# UNIVERSITY OF NOVA GORICA GRADUATE SCHOOL

# TOXIC EFFECTS OF SELECTED NEONICOTINOIDS THROUGH DIFFERENT ORGANISATIONAL LEVELS: IN VITRO AND IN VIVO STUDIES

**DISSERTATION** 

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"Science... never solves a problem without creating ten more."

G. B. Shaw

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#### **ABSTRACT**

Neonicotinoids are widely applied pesticides due to their higher affinity for insect nicotinic acetylcholine receptors. These compounds are extensively applied to control pest insects in different agricultural crops; however they can also affect non-target organisms (humans or biota). Still a limited number of studies are referring to neonicotinoids in terms of potential hazard for the additive/cumulative effects on human health and to toxic effects of their transformation products on aquatic non-target organisms.

In the scientific literature, data on the effects of IMI, its agricultural products and its transformation products are not sufficient. Due to this fact during this research we wanted to test the toxic effect of imidacloprid-IMI (as pure compound or commercial formulation Confidor 200SL) and its four transformation products (6-chloronicotinic acid-6CNA, desnitro-IMI, olefin-IMI and 5-hydroxy-IMI) applying a battery of acute *in vivo* and *in vitro* bioassays. *In vivo* assays included as model organisms: luminescent bacteria *Vibrio fischeri*, green microalgae *Desmodesmus subspicatus* and amphipod crustacean *Gammarus fossarum*, while *in vitro* assays involved bacteria *Salmonella typhimurium* and sensory neuronal F11 cell line.

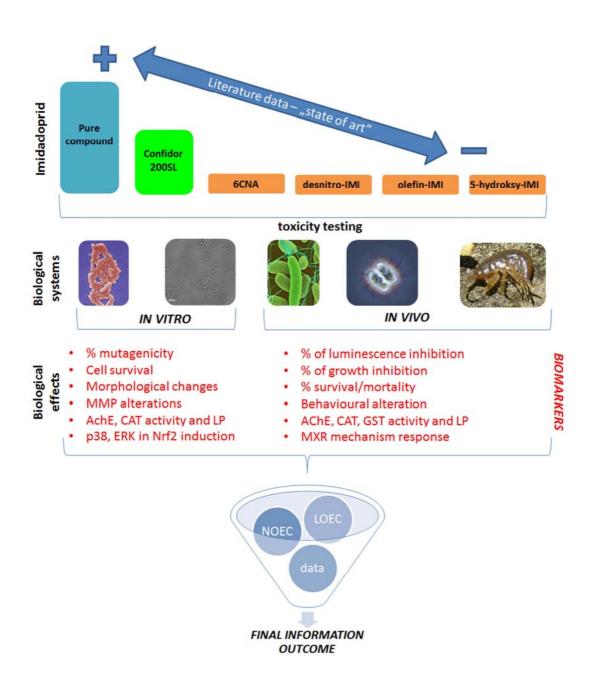
In this work we studied the effects of IMI (1–4000  $\mu$ M; 0.25–1022.4 mg L<sup>-1</sup>) on F11 cell line applying a battery of measured parameters. IMI showed significant cytotoxic effects in F11 cells only at concentrations higher than 1000  $\mu$ M (255.6 mg L<sup>-1</sup>), confirming its overall low toxicity. More evident reduction of cell survival was in presence of Confidor 200SL, desnitro-IMI, 5-hydroxy-IMI and olefin-IMI compared to pure compound alone. F11 cells exposed for 24 h to IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) increased the presence of picnotic nuclei, loosed plasma membrane integrity and showed strong degradation of cytoskeleton elements. After 48 h, IMI induced a drastic re-organisation of the cytoskeleton leading to nuclear condensation. F11 cells exposed to IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) showed early activation of p38-and ERK-mediated intracellular pathways after 30 min, significantly increased lipid peroxidation after 24 and 48 h exposure and slight loss of mitochondrial activity after 15 min at same concentration. Inhibitors of p38 were only moderately sufficient to block IMI cytotoxicity, while anti-oxidant compounds such as vitamin C and E were more protective. These data indicate that oxidative stress induction has a potential

crucial role by which IMI exerts its toxic effects in neuronal cells and suggest caution in potential long-term sensitisation of peripheral sensory neurons even at sub-threshold doses. Results relative to possible mutagenic impact of IMI, Confidor 200SL and 6CNA tested with *S. typhimurium* reverse mutation test did not cause any mutations.

In case of *in vivo* tested models, oxidative stress responses and behavioural changes after 24 h in the crustacean amphipod *G. fossarum* were investigated as well as the growth rate in freshwater algae *D. subspicatus* after 96 h exposure to IMI, its commercial formulation Confidor 200SL and its transformation product 6CNA. Algal growth has shown significant sensitivity to Confidor 200SL and 6CNA acid when compared to IMI. In the case of amphipods, low doses of IMI (102.2 μg L<sup>-1</sup>) were sufficient to induce lipid peroxidation, while Confidor 200SL induced increased catalase activity (511.3 μg L<sup>-1</sup>) and peroxidation (255.6 μg L<sup>-1</sup>). 6CNA altered significantly only antioxidant mechanisms (catalase activity) without changing peroxidation levels. Furthermore, toxicity testing of aqueous solutions for three IMI transformation products (5-hydroxy-IMI, desnitro-IMI and olefin-IMI) was performed with the luminescent bacteria. Different toxic levels of luminescent inhibition were demonstrated also for the IMI products tested on *V. fischeri*.

Given the very different responses of organisms to tested compounds, it would be reasonable to extend the research not only to detect other effects, but also to implement safer use and more efficient removal. These study measurements of different biological effects and responses are helpful to understand the mechanism of IMI or its transformation products-induced oxidative stress. Obtained data demonstrate potential harmful effects of neonicotinoid-based pesticides on non-target organisms. In general, toxicity testing of pesticides should be performed on models at different levels of biological organisation integrated with a multi-biomarker approach providing a complete functional 'picture' and better comparison of obtained data.

#### **GRAPHICAL ABSTRACT**



#### **POVZETEK**

Neonikotinoidi se pogosto uporabljajo kot pesticidi zaradi njihove visoke afinitete vezave na nikotinske holinergične (ali acetilholinske) receptorje žuželk. Neonikotinoidi se pretežno uporabljajo za zatiranje insektov na različnih kmetijskih pridelkih, žal pa lahko vplivajo tudi na netarčne organizme (ljudi, druga živa bitja). Omejeno število študij sicer obravnava neonikotinoide kot potencialno nevarne v smislu aditivnih/kumulativnih učinkov na zdravje ljudi in strupenih učinkov njihovih produktov pretvorbe (npr. intermediatov, vmesnih produktov) na netarčne vodne organizme.

V znanstveni literaturi podatkov o učinkih IMI, njegovih kmetijskih pripravkov ter produktih pretvorbe ni dovolj. Zato smo v sklopu naših raziskav preučevali učinke imidakloprida (IMI), komercialnega pripravka Confidor 200SL ter štirih produktov pretvorbe (6-kloronikotinska kislina-6CNA, deznitro-IMI, olefin-IMI in 5-hidroksi-IMI) z uporabo različnih akutnih *in vivo* kot tudi in *vitro* bioloških preskusov. *In vivo* preskuse smo izvedli z naslednjimi modelnimi organizmi: luminescenčnimi bakterijami *Vibrio fischeri*, zelenimi algami *Desmodesmus subspicatus* ter raki postranicami *Gammarus fossarum*, medtem ko so *in vitro* biološki preskusi vključevali bakterije *Salmonella typhimurium* ter senzorske nevronske celice F11.

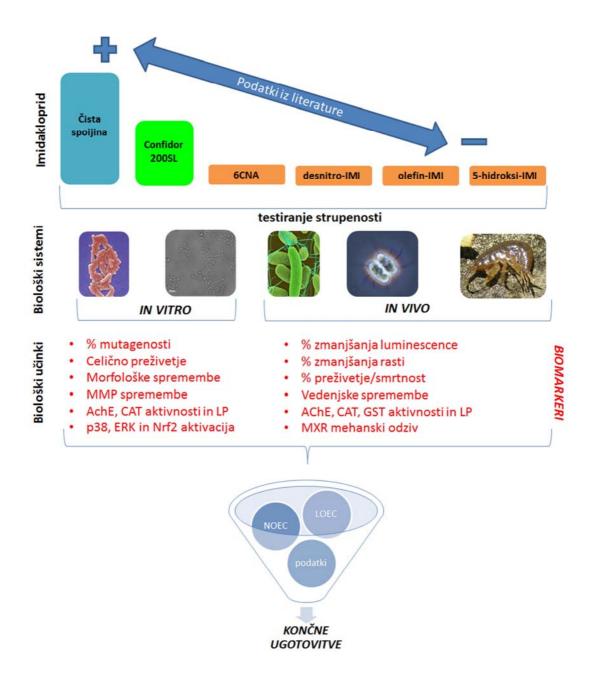
Za ugotavljanje učinkov IMI-ja (1–4000 μM; 0.25–1022.4 mg L<sup>-1</sup>) smo uporabili F11 celično linijo, pri kateri smo merili serijo različnih parametrov. Rezultati so pokazali citotoksične učinke le pri koncentracijah IMI, višjih od 1000 μM (255.6 mg L<sup>-1</sup>), s čimer smo potrdili njegovo poznano nizko toksičnost. Opaznejše zmanjšanje preživetja celic smo zabeležili v prisotnosti Confidorja 200SL, deznitro-IMI-ja, 5-hidroksi-IMI-ja ter olefin-IMI-ja v primerjavi z imidaklopridom. Ko smo F11 celice izpostavili IMI-ju za 24 ur (4000 μM; 1022.4 mg L<sup>-1</sup>) je prišlo do porasta piknotičnih jeder, izgube integritete plazemske membrane ter znatnega propada citoskeletnih elementov. Po 48 h je IMI induciral intenzivno reorganizacijo citoskeleta v smeri kondenzacije jeder. F11 celice, izpostavljene IMI-ju (4000 μM; 1022.4 mg L<sup>-1</sup>) so po 30 min izkazovale zgodnjo aktivacijo p38- in ERK intracelularnih poti, po 24 in 48 h izpostavitvi pa značilno povišano stopnjo lipidne peroksidacije ter rahlo znižanje mitohondrijske aktivnosti po 15 min. Inhibitorji p38

so le delno blokirali IMI-jevo citotoksičnost, medtem ko so druge antioksidativne spojine, kot sta vitamina C in E, delovale bolj zaščitno. Ti rezultati nakazujejo, da ima indukcija oksidativenga stresa pomembno vlogo, s čimer IMI strupeno deluje na nevronske celice. Zato je potrebna previdnost v primeru dolgotrajne preobčutljivosti perifernih senzornih nevronov že pri koncentracijah, ki so pod pragom strupenosti. Rezultati testiranj z uporabo *S. typhimurium*, vezanih na možne mutagene učinke IMI-ja, Confidorja 200SL ter 6CNA, niso pokazali nikakršnih mutacij.

V primeru *in vivo* bioloških testov, smo v rakih postranicah *G. fossarum* preučevali odzive, vezane na oksidativni stres ter spremembe obnašanja po 24 h izpostavitvi, v sladkovodnih algah *D. subspicatus* pa hitrost rasti po 96 h izpostavitvi IMI-ju, komercialnemu pripravku Confidorju 200SL ter pretvorbenemu produktu 6CNA. Rast alg je bila manjša pri izpostavitvi Confidorju 200SL in 6CNA v primerjavi z IMI. V primeru rakov so le nizke koncentracije IMI-ja (102.2 μg L<sup>-1</sup>) inducirale lipidno peroksidacijo, medtem ko je Confidor 200SL induciral tako katalazno aktivnost (511.3 μg L<sup>-1</sup>) kot peroksidacijo (255.6 μg L<sup>-1</sup>). 6CNA je vplivala na antioksidativni mehanizem (katalazna aktivnost) medtem ko na peroksidacijo ni vplivala. Nadaljnji poskusi, pri katerih smo luminescenčne bakterije izpostavili vodnim raztopinam treh produktov pretvorbe IMI-ja (5-hidroksi-IMI, deznitro-IMI in olefin-IMI) so pokazali različno strupenost v primerjavi z IMI-om.

Glede na zelo različne odzive organizmov na testirane spojine, bi bilo smisleno raziskave razširiti ne le na ugotavljanje drugih učinkov, pač pa tudi na varnejšo uporabo ter učinkovitejše odstranjevanje. Omenjena raziskava z meritvami različnih učinkov in odzivov v bioloških preskusih omogoča razumevanje mehanizma oksidativnega stresa, povzročenega zaradi delovanja IMI-a ter njegovih produktov pretvorbe. Dobljeni rezultati namreč kažejo na morebitne škodljive učinke neonikotinoidnih pesticidov na netarčne organizme. Na splošno pa lahko trdimo, da moramo strupenostne teste s pesticidi izvesti na modelnih sistemih na različnih nivojih biološke organizacije z uporabo večjega števila biomarkerjev. Na ta način zagotovimo učinkovitejšo primerjavo podatkov ter popolnejšo sliko delovanja preiskovanega pesticida in razumevanje pri katerih koncentracijah ima substanca kvarne učinke.

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- **Figure 32:** A. Bright field representative images of F11 cells in control and after 24 and 48 h IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) exposure. Scale bar: 50  $\mu$ m. B. Values of MTT assay in F11 cells after exposure to IMI for 24 and 48 h; data presented as percentage of control shown as dashed line (n=9). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 33:** Representative microscopy images of F11 cells morphological characterisation in basal conditions and after IMI exposure (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) for 48 h. Cells were stained with different fluorescent markers, as indicated for each image. To note: stronger nuclear DAPI staining signal after treatment and resilient degradation of cellular proteins (actin staining). Scale bar 100  $\mu$ m.

- **Figure 34:** Morphology of F11 cells in control and after IMI exposure (4000 μM; 1022.4 mg L<sup>-1</sup>) for 48 h in standard growing conditions (FBS) or in absence of serum (SF). A. Nuclei are counterstained with DAPI (blue), β-Tubulin III immunofluorescence staining is shown in green. To note: protein aggregates and nuclear condensation in F11 cells after IMI exposure (yellow rounded rectangles). B. Nuclei are counterstained with DAPI (blue), β-Tubulin III immunofluorescence staining is shown in red. To note: IMI stimulated processes elongation (arrows). Scale bar: 100 μm.
- Figure 35: Values of MTT assay in F11 cells after exposure to IMI, Confidor 200SL and 6CNA for 24 h (A) and 48 h (B); data presented as percentage of control shown as dashed line (n=6). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).
- **Figure 36:** Values of MTT assay in F11 cells after exposure to IMI, desnitro-IMI, 5-hydroxy-IMI and olefin-IMI for 24 h (A) and 48 h (B); data presented as percentage of control shown as dashed line (n=6). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 37:** A, B. Representative fluorescence microscopy images of F11 cells in control conditions and in the presence of IMI 400 μM (102.2 mg  $L^{-1}$ ) and 4000 μM (1022.4 mg  $L^{-1}$ ) immunostained with antibodies anti-active Erk (A) or active p38 MAPK (B). Scale bar: 50 μm. Histograms quantify the grey values of immunopositive signals after background value subtraction. Data have been normalised on total cell number (counterstained with DAPI) and expressed as arbitrary units (AU). To note: different activation profile. C. Graphs quantify p38 signal intensity during the time after application of IMI at 4000 μM to F11 cells (left) and nuclear translocation of activated phosphorylated p38 during time after IMI application (right). (n= 3). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 38:** Representative microscopy images of translocated phosphorylated p38 MAPK in nucleus of F11 cells exposed to IMI (4000 μM; 1022.4 mg L<sup>-1</sup>) for 10 min. Scale bar 10 μm.
- **Figure 39:** A. Representative example of F11 cells microscopy images analysed in control and after IMI incubation at 4000 μM (1022.4 mg  $L^{-1}$ ) for 30 min. Cells are immunolabelled with anti-Nrf2 antibodies. Scale bar: 100 μm B. The histogram describes the time course (10, 30, 60, 120 and 240 min) of the relative average quantification of Nrf2 immunofluorescence reactivity in control and in IMI-exposed cells (4000 μM; 1022.4 mg  $L^{-1}$ ). Histograms quantify the grey values of immunopositive signals after background value subtraction. Data have been normalised on total cell number (counterstained with DAPI) and expressed as arbitrary units (AU). (n= 3), p < 0.05 (\*).

- **Figure 40:** A. *In vivo* imaging of live cells exposed to IMI 4000 μM (1022.4 mg L<sup>-1</sup>) 15 and 60 min. Representative microscopy images show JC-1 fluorescence of control (red) and IMI-treated F11 cells (green). Scale bar 100 μm. Histogram represent JC-1 green-red fluorescence signal ratio already significant after 15 min from IMI application. Fluorescence values were expressed as the ratio of fluorescence intensity of J-aggregates (red) to fluorescence intensity of monomers (green); ratio results are presented as fold increase/decrease of the control values (JC-1/control in arbitrary units AUs) (n=3). B. Catalase activity in F11 cells tested 24 or 48 h after IMI application 1000 μM (255.6 mg L<sup>-1</sup>) and 4000 μM (1022.4 mg L<sup>-1</sup>) (n =3). C. Quantification of TBARS products released in cell growth medium by cell exposed to IMI 1000 μM and 4000 μM for 24 and 48 h (n=3). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 41:** Values of MTT assay in F11 cells after 24 and 48 h exposure to 1000 μM (255.6 mg L<sup>-1</sup>) (A) and 4000 μM (1022.4 mg L<sup>-1</sup>) IMI (B) in presence of p38 MAPK inhibitor SB203580, *N*-acetyl-cysteine (NAC) or mixed tocopherols (vitamin E); data presented as percentage of control shown as dashed line (n=4). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*) compared to IMI-exposed cells.
- **Figure 42:** Dose response curve for aqueous solution of olefin-IMI (A), desnitro-IMI (B) and 5-hydroxy-IMI (C) for *V. fischeri* luminescent bacteria within 30 minutes of exposure.
- Figure 42a: Gamma values ploted against their corresponding chemical concentration (A) olefin-IMI, (B) desnitro-IMI, (C) 5-hydroxy-IMI.  $EC_{20}$  value is given by the point of intersection with the x-axis at  $\Gamma = 0.25$ .
- **Figure 43:** *D. subspicatus* % of algal growth compared to control after exposure to IMI (A) Confidor 200SL (B) and 6CNA (C) at 24, 48, 72 and 96 h. The inside graph represents exposure to negative control (known co-formulants only). Data are reported as mean  $\pm$  standard error (n=3). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).
- **Figure 44:** Mortality rate of *G. fossarum* after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). (n=50). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 45:** Whole-body CAT activity ( $\mu$ mol/min/mg protein) of *G. fossarum* measured after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75% of all readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $^{\bullet}$ ). (n=10). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 46:** Whole-body GST activity (nmol/min/mg protein) of *G. fossarum* measured after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75% of all

readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $\bullet$ ). (n = 10). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).

- **Figure 47:** Whole-body LP of *G. fossarum* (expressed in absorbance units of TBARS products) measured after 24 h of exposure IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75% of all readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $^{\bullet}$ ). (n=10). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).
- **Figure 48:** Inhibitory potential of IMI, Confidor 200SL and 6CNA (0.7 or 7.6 mg L<sup>-1</sup>) and copper (3 and 10 μg L<sup>-1</sup>) on the accumulation of fluorescent Rh B dye in *G. fossarum*. Results are expressed in fluorescence units (f.u.) normalised on total animal wet body weight as an average of three separated experiments. (n=3). p < 0.05 (\*).
- **Figure 49:** Schematic diagram describing the possible cytotoxic model of IMI and its activated molecular pathways in human neuroblastoma F11 cells. (YNM, 2010, with slight modifications).

#### ABBREVIATIONS AND SYMBOLS

a. i. active ingredient

ACh acetylcholine

AChE acetylcholinesterase

nAChe nicotinic acetylcholine receptor

ATP adenosine triphosphate

AU absorbance unit/ arbitrary unit

BDNF brain derived neurotrophic factor

BSA bovine serum albumin

BTH butylatedhydroxytoulene

CAT catalase

CCD colony collapse disorder

CDNB 1-chloro-2,4-dinitrobenzene

CGRP calcitonin gene-related peptide

DMSO dimethyl sulfoxide

DTNB 5,5 dithiobis-2-nitrobenzoic acid

DMEM Dulbecco's modified Eagle medium

 $EC_{50(20)}$  median (50 %; 20%) effective concentration

ECPA European Crop Protection Association

ECVAM European Centre for Validation of Alternative Methods

EFSA European Food Safety Authority

em. Emission

ERA Environmental Risk Assessment

ETS electron transport system

ex. Excitation

FAO Food and Agricultural Organisation

FBS foetal bovine serum

GDP gross domestic product

 $GP_x(GSH-P_x)$  glutathione peroxidase

GSH Glutathione

GST glutathione-S-transferase

G6PD glucose-6-phosphate dehydrogenase

HPLC-DAD high performance liquid chromatography – diode array

detector

HQ hazard quotient
IMI Imidacloprid

IC<sub>50</sub> median (50 %) inhibition concentration

IF Immunofluorescence

IPM Integrated Pest Management

LC<sub>50</sub> median (50 %) lethal concentration

LC-MS liquid chromatography-mass spectrometry

 $LD_{50}$  median (50 %) lethal dose LGIC ligand-gated ion channel

LO(A)EC(L) lowest observed (adverse) effect concentration (level)

LP lipid peroxidation

MAPK mitogen activated kinase

MDA Malondialdehyde

MMS methyl methanesulphonate

MRL maximum residue level

MRP multidrug resistance protein

MXR multixenobiotic resistance

NAC *N*-acetyl-*L*-cystein

NADP β-nicotinamide adenine dinucleotide phosphate sodium salt

hydrate

NGF neuronal growth factor

NMP N-methylpirrolidone

NO(A)EC(L) no observed (adverse) effect concentration (level)

Nrf2 nuclear factor (erythroid-derived 2)-like 2

NRC National Research Council
PBS phosphate buffered saline

PEC predicted environmental concentration

PLL poly-*L*-lysine hydrochloride

PKC protein kinase C

ppb part per billion

ppm part per million

PPP plant protective products
ROS reactive oxygen species

RNS reactive nitrogen species

RQ risk quotient

RT room temperature

SOD superoxide dismutase

TBA thiobarbituric acid

TBARS thiobarbituric acid reactive substances

TCA trichloroacetic acid

TN total nitrogen

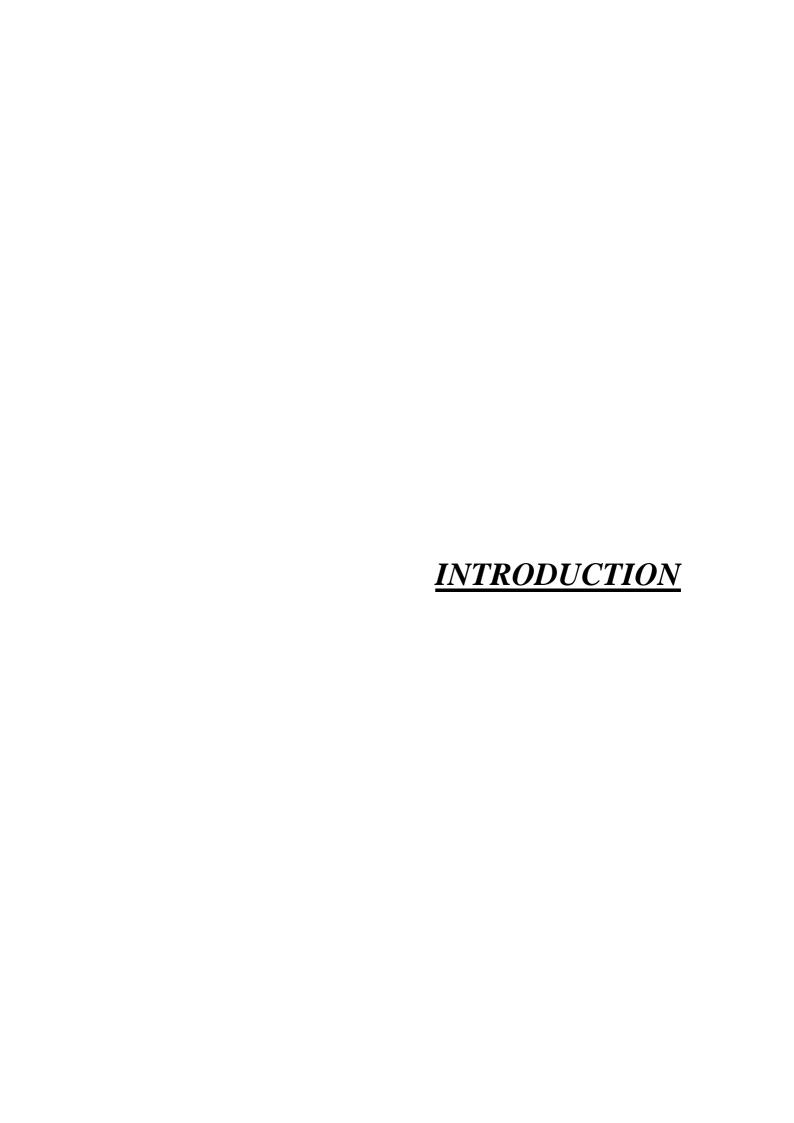
TOC total organic carbon

TP transformation product

US EPA US Environmental Protection Agency

WHO World Health Organisation

6CNA 6-chloronicotinic acid



#### 1. INTRODUCTION

Pesticides are the pollutants increasingly present in the surrounding environment, which are often persistent and can be bioaccumulated through the biological chains such as soil-plant-food or water-aquatic organism-food (Preston, 2002). The usage of high amount of pesticides in environment represents a possible risk for biota and human health due to their potential toxic action. Pesticide substances are **biologically active** and must be tested to ensure that their use will not give rise to any unacceptable risks to **non-target organisms** (i.e. humans, animals and plants) or to the environment. As part of these testing, pesticides are fully investigated for potential acute, sub-chronic and chronic toxic effects with the use of different **biological systems** and with the measurement of selected **biomarkers** or conventional **endpoints**.

## 1.1. Pesticides in general

As the Food and Agriculture Organisation of the United Nations (FAO) defined, pesticide is any substance or mixture of substances intended for preventing, destroying, repelling, or mitigating any pest, including vectors of human or animal disease or weed which can cause harm during or otherwise interfering with the production, processing, storage, transport or marketing of food, agricultural commodities, wood and wood products or animal feedstuffs. Pesticides can be classified according to their **target**, their **mode or period of action**, or their **chemistry**. They may be chemical substances, biological agents (such as viruses or bacteria), antimicrobials, disinfectants or devices used against any pest (Saravi and Shokrzadeh, 2011). Mainly, agricultural pesticides are divided into five categories based on the targeted pest and include: insecticides, herbicides, fungicides, rodenticides, and fumigants (Saravi and Shokrzadeh, 2011). Recently, the European Food Safety Authority (EFSA) tried to replace the expression 'pesticide' due to its negative implications, with the new term-'plant protective product' (PPP).

Pesticides are a highly diverse group of compounds and present one of the most important groups of chemical stressors in the environment (Hanazato, 2001; Liess and Ohe, 2005). As an alternative to first generation pesticides, modern pesticides have been designed to avoid some of the deleterious effects of these pesticides that had a long residence time in the environment and accumulated in the food chain (Meleiro Porto *et al.*, 2011). Newer classes of highly selective, systemic and single mode activity pesticides were introduced in the early 1990s and promised to address pest more specifically. Over the past twenty years, a class of systemic insecticides called **neonicotinoids** gained increasing interest in the agricultural sector. The term 'neonicotinoid pesticides' comprises a group of several different insecticides, but it is usually used to indicate the four that are most widely applied: imidacloprid, thiacloprid, clothianidin and thiamethoxam. One of these representatives, imidacloprid (IMI), was the main focus of this research.

In the last decades, the world has known for a continuous growth of pesticide usage, both in number of chemicals and quantities sprayed over the agricultural fields (Carvalho, 2006). According to the data supplied by European Crop Protection Association (ECPA), the total amount of PPP used in the European Union (reported as tonnes of active ingredient (a. i.) for the main PPP categories) increased steadily in the 1990s, stabilised in the late '90s and these data have not altered significantly between 1992 and 2003 (Fig. 1A). Especially, is important to notice that the use of insecticides more than doubled between 1992 and 2000, with only a slight decrease from 2000 to 2003 (Fig. 1B) (EUROSTAT, 2007).

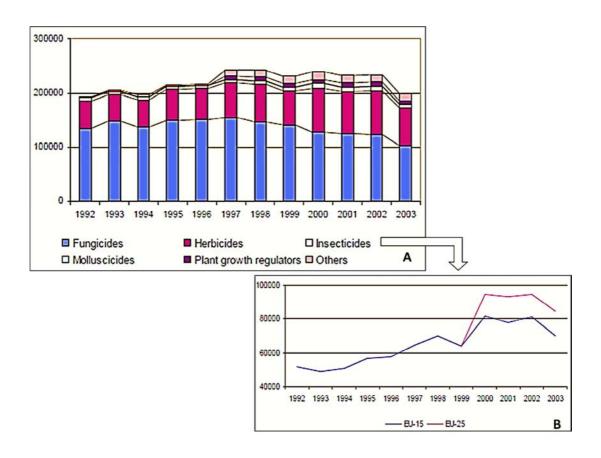


Figure 1: (A) Use and composition of PPP in 15 Member States of the European Union (EU-15), 1992-2003 (y-axis; in tonnes of a. i.) (B) Use of PPP-insecticides in all Member States of European Union (EU-25) and in EU-15, 1992-2003 (y-axis; in tonnes of a. i.). (EUROSTAT, 2007).

The Community regulatory framework concerning pesticides focuses particularly on the placing on the market and the end of the life cycle of such products. The most relevant legislative measures concerning PPPs are:

- 1) Directive 91/414/EEC on the placing of PPPs on the market.
- 2) Regulation (EC) No. 396/2005 on maximum residue levels of pesticides in food and feed.
- 3) Water Framework Directive (WFD) 2000/60/EC.

Lately, the EC Directive 91/414/EEC has been replaced by EC Directive 1107/2009. The new legislation published in December 2009 which ispart of the European Union Thematic Strategy on Pesticides and is composed of four elements:

- 1) Plant Protection Products Regulation 1107/2009
- 2) Sustainable Use Directive (SUD) 2009/128/EC
- 3) Machinery Directive 2009/127/EC which sets out standards for new equipment
- 4) Statistics Regulation 1185/2009 the key elements of this regulation are the provision of annual sales data and the provision of data every five years on usage on crops and the pesticides applied.

These above-mentioned legislations are also applied in Slovenia that is one of the Member States of the EU. In comparison to other Central and Eastern European countries, the macroeconomic importance of agriculture in Slovenia is relatively low. Share of agriculture, hunting and forestry in its gross domestic product (GDP) has been decreasing in the last decade (Slabe, 2001). The PPP market in Slovenia is dominated by herbicides and fungicides in comparable proportions (Fig. 2). Over the last years, total pesticide products use has been increasing. Average use of pesticide products in 2000 was estimated at 3.1 kg ha<sup>-1</sup> of agriculture land (GIS, 2003).

The use of **neonicotinoids** has been approved in the EU and they are currently in use. IMI, thiamethoxam, acetamiprid and thiacloprid are registered also on the Slovenian market. However, since May 2008, the insecticides thiamethoxam, clothianidin and IMI were banned for treatment of corn seeds (Zea mays), sugar beet seeds (Beta vulgaris) and oilseed rape seeds (Brassica napus) (The Official Gazette of the Republic of Slovenia No. 50/2008). Despite all the precautions taken in order to avoid the poisoning of bees, a large number of bee's colonies died in the eastern part of Slovenia during April 2011 (Pomor čebel v Pomurju dobiva nove razseznosti, 2011). Due to these repetitive negative events, the Phytosanitary Administration of Slovenia banned completely clothianidin, IMI and thiamethoxam for any seed treatment, also other EU member states such as France, Germany and Italy imposed similar bans for IMI and clothianidin. Moreover, clothianidin is completely banned in Slovenia for any agricultural use (The Official Gazette of the Republic of Slovenia No. 31/2011). It was estimated that in Slovenia approximately 2,400 kg of neonicotinoids was sold in the year 2004 (Phytosanitary Administration of the Republic of Slovenia, personal communication, 2005; Žabar, 2012).

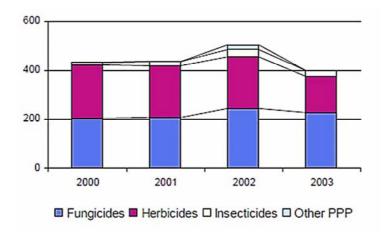


Figure 2: Use and composition of PPP in Slovenia, 2000-2003 (y-axis; in tonnes of a.i.) (EUROSTAT, 2007).

### 1.2. Neonicotinoids – Imidacloprid

**Neonicotinoids** are a group of insecticides derived from nicotine isolated from the tobacco plant (*Nicotiana tabaccum*) which presents insecticidal activity and has been used extensively as natural insecticide. The developing road from nicotine to neonicotinoids was long and complicated. Neonicotinoids effective control of insect pests and helminthic parasites has been achieved by targeting invertebrate (insect) nAChRs (Matsuda *et al.*, 2005; Tomizawa and Casida, 2003; 2005). The major commercial insecticides targeting insect nAChRs were not derived from natural products but rather from the discovery of synthetic nitromethylene heterocyles (Kagabu, 1997). Introduction of the 6-chloro-3-pyridylmethyl and nitroimine moieties led to the development of the first type of nicotinic insecticide called imidacloprid (Kagabu, 1997).

**Imidacloprid** (**IMI**) is one of the major representatives of the new generation of neonicotinoid insecticides. It was patented for the first time in 1985 by Bayer and was placed on the market in 1991. Today it has been made commercially available by Bayer AG and Nihon Tokushu Noyaku Seizo KK. It is a nicotine derived compound (neonicotinoid) with a large potential distribution due to its agonistic action on insect nAChRs and its selective toxicity to insects over vertebrates (Tomizawa and Casida, 2003). IMI [1-[(6-chloro-3 pyridynil) methyl]-*N*-nitro-2-imidazolidinimine] has the molecular formula C<sub>9</sub>H<sub>10</sub>ClN<sub>5</sub>O<sub>2</sub> (Fig. 3), with a molecular weight of 255.7 g mol<sup>-1</sup>. In appearance, it consists of colourless crystals.

Figure 3: Chemical structure of imidacloprid.

This component is a relatively new pest control substance, which is having the fastest growing sales worldwide (Tomizawa and Casida, 2005) and is generating increasing concern on its possible impacts on natural ecosystems (Matsuda *et al.*, 2001; Jemec *et al.*, 2007). Elbert *et al.* (2008) pointed out remarkable data revealing the turnover toward insecticidal seed treatment. By the year 2005 seed coating developed into a €535 million market, with a 77 % share for neonicotinoid insecticides.

IMI's physicochemical properties render it useful for a wide range of application techniques, including foliar, seed treatment, soil drench, and stem application (Millar and Denholm, 2007) (Table 1). It is a versatile, broad-spectrum, systemic insecticide with activity against sucking insects (e.g. aphids, leafhoppers, whiteflies, and termites), several species of *Coleoptera*, *Diptera*, *Lepidoptera* and parasites on a different types of crops (Tomlin, 1994; Tomizawa and Casida, 2005). In addition, it is also applied as veterinary medicine against parasites and fleas in dogs and cats. Other important neonicotinoids are acetamiprid, clothianidin, dinotefuran, thiacloprid and thiamethoxam.

Commercial formulations of IMI used for the control of different pest insects are available in a very large number (e.g. Admire, Merit, Confidor 200SL, Provado and Gaucho) (PMRA, 2005). Formulations of IMI are present as: a slurry or flowable concentrate for seed treatments, granule, wettable powder, soluble concentrate, and suspension concentrate, water dispersible granules, and dustable powder (Tomlin, 2004). To make a pure pesticide molecule applicable and highly functional, a. i. are combined with solvents or surfactants to accomplish an effective action of these formulations in agricultural applications. These supplementary substances often represent the highest proportion in pesticide marketed mixtures and include co-formulants (e.g. dimethyl sulfoxide (DMSO), N-methylpyrrolidone (NMP), propylene carbonate and other solvents) that could modify IMI's toxicity and bioavailability. Formulations of IMI include also other chemicals such as crystalline quartz silica and naphthalene chemicals which have also associated toxicological characteristics (Cox, 2001). Even a minor concern regarding the toxicity of these compounds and their possible synergistic effect with IMI or other ingredients should be continuously considered (Tobiassen et al., 2003; Surgan, 2005). It is important to

notice that the commercial formulations are the ones applied in the environment with relevant soil (50 g ha<sup>-1</sup> to 320 g ha<sup>-1</sup>) and foliar concentrations (73 mg L<sup>-1</sup> to 150 mg L<sup>-1</sup>). Recent evaluation of the data relative to different formulations noted high levels of IMI in leaves and in blossoms of treated plants, and increases in residue levels over time (CDPR, 2011). Data indicated that the use of these IMI formulations on annual basis may be at the end cumulative. Due to recent findings certain commercial products within the class of neonicotinoids (containing a. i. IMI) were placed under re-evaluation and need further studies (CDPR, 2011).

Even if IMI has been in use for a relatively short period of time compared to other common pesticides, it is considered to being used in the largest quantity worldwide of all insecticides (Cox, 2001; Ware and Whitacre, 2004). Comparative toxicity studies indicate IMI's adverse effects to some aquatic invertebrates and have noticed high species-specific response to IMI, which suggests that IMI toxicity data may not be generalised (Jemec *et al.*, 2007). In addition, IMI can be applied at very low rates and during whole year.

In the present research, IMI - as **pure compound** and as **commercial formulation Confidor 200SL** was selected for further studies and investigated for potential toxicity to different non-target organisms (Fig. 4).

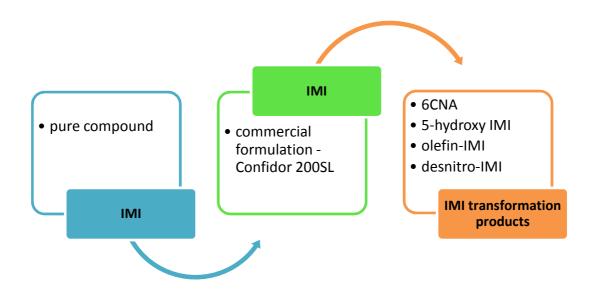


Figure 4: Schematic diagram of selected compounds which were tested on different biological systems during this study.

Table 1: Selected properties of IMI.

## **Imidacloprid**

		Ref. No.
Physico-chemical properties		2103. 110.
CAS number	138261-41-3	(1)
Water solubility at 20°C (mg L <sup>-1</sup> )	(a) 510 (b) 610	(a) (2) (b) (1, 3)
$K_{oc}$ (soil organic carbon–water partitioning coefficient)	(a) 210 (20 °C) (b) 249-268 (c) 109-411 (20°C)	(a) (4) (b) (5) (c) (3)
$Log K_{ow}$ (octanol-water partition coefficient)	(a) 0.57 (b) 0.92	(a) (1) (b) (4)
Average application rates: soil (g ha <sup>-1</sup> ) and foliar (mg L <sup>-1</sup> ) applications	(a) 50-320 (b) 73-150	(a) (6) (b) (7)
Environmental fate		Ref. No.
Detected aquatic concentrations (µg L <sup>-1</sup> )	<ul> <li>(a) 1 (surface water, Florida, USA)</li> <li>(b) 6.7 (ground water, Florida, USA)</li> <li>(c) 14 (Lake Wales Ridge, USA)</li> <li>(d) 300 (surface water, Denmark)</li> </ul>	(a) (8) (b) (9) (c) (10) (d) (11)
Estimated aquatic concentrations (μg L <sup>-1</sup> )	<ul> <li>(a) 36.04 (acute surface water exposure); 17.24</li> <li>(chronic surface water exposure)</li> <li>(b) 22 (accidental direct spray in a pond or stream);</li> <li>1.8-7.3 mg L<sup>-1</sup> (accidental spill in a small pond)</li> </ul>	(a) (9) (b) (12)
Aqueous photolysis DT <sub>50</sub>	(a) 3 h (simulated sunlight, 30°C) (b) >1 h (simulated sunlight, 24 °C)	(a) (13) (b) (9)
Hydrolysis DT <sub>50</sub> (days)	(a) > 30 (b) 355 (at pH 9); stable at pH 5 and 7 (c) 36.3 at pH 4; 41.6 at pH 7	(a) (9) (b) (14) (c) (15)
Soil photolysis DT <sub>50</sub> (days)	38.9	(9)
Soil anaerobic DT <sub>50</sub> (days)	27.1	(9)
Soil aerobic DT <sub>50</sub> (days)	(a) 156 (b) 997	(a) (3) (b) (9)
Field dissipation DT <sub>50</sub> (days)	(a) 26.5 – 229 (b) 83 - more than 365 (c) 7 (corn); 12 (sandy loam); turf grass (61-107) (d) 96 (bare soil)	(a) (9) (b) (16) (c) (17) (d) (3)

**References:** (1) Tomlin, 2004; (2) Yen and Wendt, 1993; (3) Krohn and Hellpointer, 2002; (4) Nemeth-Konda et al., 2002; (5) Oi, 1999; (6) PMRA, 2005; (7) www.bayercropscience.com; (8) Jemec et al., 2007; (9) Fossen, 2006; (10) US Geological Survey, 2003; (11) Tennekes, 2010; (12) SERA, 2005; (13) Kagabu and Medej, 1995; (14) US EPA, 2005; (15) Sarkar et al., 1999; (16) Sabbagh et al., 2002; (17) CDPR, 2006.

### 1.2.1. Imidacloprid mode of action

IMI mode of action is based on the interference of the neurotransmission in the nicotinic cholinergic nervous system. IMI binds to nicotinic acetylcholine receptors (nAChRs) at the neuronal and neuromuscular junctions in insects and vertebrates. The nAChRs are members of the cys-loop ligand-gated ion channel (LGIC) superfamily consisting of five membrane proteins arranged around a central cation permeable pore (Sine and Engel, 2006) (Fig. 5). These ion channels rapidly transduce the actions of the excitatory chemical neurotransmitter acetylcholine (ACh) to membrane depolarisation at the level of synapses. The receptor is normally present in a closed state; however, after ACh binding, the complex channel opens a pore and becomes permeable for cations.

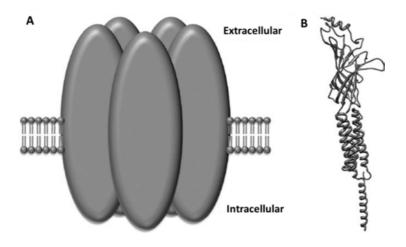


Figure 5: Nicotinic acetylcholine receptors (nAChR) and subunit structure. (A) Schematic representation of nAChR, presenting the pentameric arrangement of subunits around a central cation-permeable pore. (B) Three dimensional structure of an individual nAChR subunit showing the structure of the polypeptide backbone (Millar, 2009).

Similarly to the naturally occurring signal transmitting ACh, IMI stimulates certain nerve cells by acting on a receptor protein. IMI fits to the receptor that normally receives the molecule of ACh and irreversibly blocks postsynaptic nAChRs (Tennekes, 2010) (Fig. 6). In contrast to ACh, which is quickly degraded by enzyme acethylcholinesterase (AChE), IMI is inactivated either very slowly or not at all

(Cox, 2001). Prolonged activation of the nAChR by IMI causes desensitisation and blocking of the receptor and leads to paralysis and death.

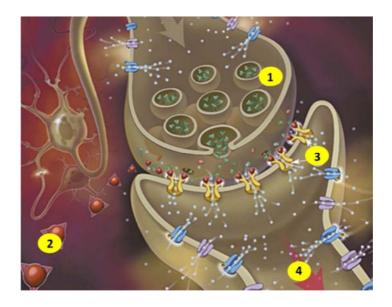


Figure 6: Site and mode of action of imidacloprid on insect neuronal cells. (1) Vesicles of acethylcholine, (2) imidacloprid, (3) nicotinic receptors blocked open and (4) constant neuromuscolar stimulation. (Bayer HealthCare  $AG^{\otimes}$ ).

To understand the activity of IMI on insect and mammalian nAChRs some structural properties are important. The IMI molecule consists of a 6-chloro-3-pyridyl group and a 2-nitroimino-imidazolidine group, bridged by a methylene group (Fig. 3). Toxicity, binding and receptor activation are affected by all three of these components and also by the overall hydrophobicity of the molecules.

Insect nAChRs - high affinity IMI binding sites in insect nAChRs were detected in a broad range of insects, including green peach aphid, glassy-winged sharpshooter, whitefly, cockroach, migratory locust and fruit fly (Zhang *et al.*, 2000; Tomizawa and Casida, 2003). The 2-nitroimino-imidazolidine group plays a key role in the selectivity of insect nAChRs. Neonicotinoids are not protonated and instead of an easily protonated nitrogen they have an electronegative nitro or cyano pharmacophore. This electronegative pharmacophore is proposed to associate with a cationic subsite in the insect nAChRs (Tomizawa and Casida, 2005). In general, the potency of IMI for insect brain nAChRs is considerably higher than for mammalian channels. For example, the binding affinity of IMI to nAChRs of *Drosophila* sp. is

over 550-fold greater than the affinity to the mammalian receptors (Tomizawa, 2001).

**Mammalian nAChRs** - structural studies suggested that the electron deficient nitrogen atom of the imidazolidine group in IMI molecule corresponds to the protonated form of nicotine and interacts with the mammalian nicotinic receptors (Matsuda *et al.*, 2000). The binding affinity and agonist potency of IMI have been reported for several vertebrate species. IMI showed a partial agonist activity with the recombinant chicken  $\alpha 4\beta 2$  and  $\alpha 7$  receptors (Matsuda *et al.*, 1998; 2000). IMI presented also agonist action on nAChRs in BC3H1 muscle cells and mouse N1E-115 neuroblastoma (Zwart *et al.*, 1994).

It is important to describe several additional effects that occur at the receptor and cellular level after exposure to IMI. It was demonstrated that IMI (after 3 days of exposure) up-regulates the α4β2 nAChR subtype in mouse M10 cell line (Tomizawa and Casida, 2000). This up-regulation is often associated with the receptor desensitisation (Pauly et al., 1996). At the cellular level, the immediate effect of nAChRs activation is an increase in the level of intracellular calcium and consequent membrane depolarisation which then activates a complex downstream signalling pathway (Berg and Conroy, 2002). One of the key components involved in the nAChR downstream signalling pathway is the extracellular signal-regulated kinase (ERK), also known as mitogen-activated protein kinase (MAPK). In the study of Tomizawa and Casida (2003) IMI activated the ERK cascade in mouse neuroblastoma N1E-115 cells after 30 min incubation. The IMI stimulation of  $\alpha 4\beta 2$ receptor was joined with the phosphorylation of ERK in a Ca<sup>2+</sup> and protein kinase C (PKC)-dependent manner (Tomizawa and Casida, 2003). ERK pathway is a necessary intermediate in the signalling from the nAChR to expression of specific genes (Chang and Berg, 2001). Several human neuropathologies have been linked to genetic alterations of nAChRs genes (Lindstrom, 2002). These receptors are also involved at different levels in several neurodegenerative diseases such as Parkinson and Alzheimer's. There is a growing number of facts (even if studies are not entirely consistent) which correlates long-term low-dose pesticide exposure along with the specific agricultural working conditions to a variety of disease conditions including cancers, reproductive health problems and a range of neurological disorders (Parrón et al., 2011).

#### 1.2.2. Environmental fate and behaviour of imidacloprid

After the application of pesticides used for pest control in agriculture a lot of different transportation and transformation processes may occur. The fate of a pesticide comprises its behaviour in all three major compartment of the environment: air, soil and water. In general, the principal routes of dissipation for IMI in the environment are aqueous photolysis, microbial degradation and uptake by plants.

**Air:** Some formulations of IMI are applied as sprays, allowing for possible off-site movements through drift. IMI's low vapour pressure (2 x 10<sup>-7</sup> Pa at 20° C) and low soil adsorption coefficient indicates that its volatilisation from soil and leaf surfaces may not be the main route of dissipation. Combination of these characteristics with low Henry's law constant of 6.5 x 10<sup>-11</sup> atm m<sup>3</sup> mol<sup>-1</sup> makes it unlikely that IMI will be present in the air in measurable amounts after application by any method. For example, an air monitoring study of IMI residues has been performed by the California Department of Pesticide Regulation (CDPR) in cropped field treated with IMI for control of glassy-winged sharpshooter (*Homalodisca coagulata*). Air samples collected immediately after spray applications in this area did not detect any residues of IMI (Walters *et al.*, 2001; Segawa *et al.*, 2004).

**Soil:** The relevant water solubility and low K<sub>oc</sub> for IMI indicate a low predisposition for adsorption to soil particles (Table 1). Field studies have produced a wide range of half-life values (DT<sub>50</sub>). DT<sub>50</sub> is the time required for 50% of the field-applied pesticide to dissipate and in case of IMI can range anywhere from approximately 80 days to 2 years (Table 1). Some authors consider IMI as relatively immobile in soil and do not expect its significant leaching behaviour (Krohn and Hellpointner, 2002), while some studies indicate the opposite (Gupta *et al.*, 2002). IMI persistence in soil depends on a wide range of factors such as: soil type, pH, and presence or absence of ground cover. Very important factors, which also influence the mobility, are the chemical nature of active substances and the formulation characteristics (Gupta *et al.*, 2002). Furthermore, degradation of IMI in soil is decreased if organic materials used to improve soil quality (organic fertilisers) are added (Rouchaud *et al.*, 1996).

Water: Based on solubility of IMI and its persistence, IMI is considered to have high leaching potential. The solubility of IMI in water is relatively high, and its octanol-water partitioning coefficient is quite low (Table 1). Generally, IMI is persistent in water and not easily biodegradable (Tišler *et al.*, 2009). The influences of pH and applied formulation on the persistence of IMI in water have also been studied, and it was found that a higher pH, meaning alkaline conditions, increases half-life time and consequently its persistence. At environmentally relevant pH values, hydrolysis of this insecticide is greater than 30 days, but it can be rapidly degraded through photolysis with a half-life of 1-3 hours (Tišler *et al.*, 2009). The formulation of the pesticide also presents a significant effect on IMI persistence. For example, the powder formulation Gaucho 70WS was more persistent in water than the liquid formulation Confidor 200SL (Sarkar *et al.*, 1999).

# 1.3. Transformation products of imidacloprid

Pesticides can be transformed in the environment, crops or organisms into a large number of different products, called **transformation products** (**TPs**). It is widely known that TPs may have different properties which allow them to occur in areas which usually cannot be reached by a pesticide itself. Due to their higher polarity and mobility in the soil-water environment, TPs may reach surface- and ground-water more easily than the parent compound (Hernández *et al.*, 2008). These products can be more toxic (Bavcon Kralj *et al.*, 2007) and more persistent than the parent compound (Kolpin *et al.*, 2009).

IMI's chemistry is founded on nitromethylene derivatives, the activity or stability of which is modified mainly by three processes: (1) change of the heterocyclic ring to an imidazolidine ring; (2) modification of the nitromethylene moiety to nitroimino; (3) and introduction of a pyridyl moiety (Iwaya and Kagabu, 1998). All transformation processes are related to these three possible pathways.

The TPs of IMI in soil are typically IMI urea, 6-chloronicotinic acid (6CNA) and 6-hydroxynicotinic acid (Scholz and Spiteller, 1992; Rouchaud et al., 1996) (Fig. 7). It is important to notice that IMI degradates express mainly higher water solubility and consequently higher leaching or contamination potential of aquatic compartments. 6CNA is one of the final transformation products of IMI and due to its high water solubility (2 g L<sup>-1</sup>) it may leach from soil into the aquatic environment. IMI's major photolysis breakdown product in water are 6-chloronicotinic aldehyde, 6-chloro-N-methylnicotinacidamide, 6-chloro-3-pyrydylmethylethylendiamine, IMIurea, 6-hydroxynicotinic acid, and a minor breakdown product is IMI-guanidine (Bacey, 2001). The major TPs resulting from incubation of aqueous samples under the environmentally relevant non-sterile conditions and light exposure were desnitro-IMI, IMI-urea and 6CNA (Mulye, 1995) (Table 2). Under dark, non-sterile, anaerobic conditions, desnitro-IMI was the major TP of IMI and has been found to be more persistent than its parent compound (Mulye, 1995). IMI taken up by the crops is metabolised to several major products including desnitro-IMI, 5-hydroxy-IMI and 6CNA (Table 2).

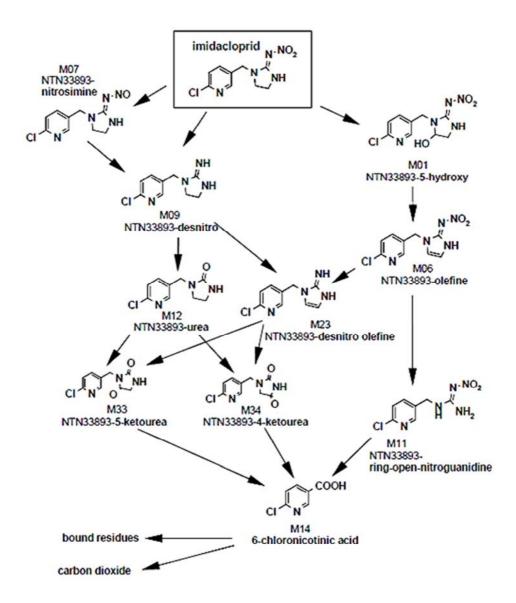


Figure 7: Probable degradation pathway in soil (main TPs) (IMIDACLOPRID (206) by U. Banasiak).

The TPs produced through *in vivo* metabolism of neonicotinoid compounds in animals (i.e. mouse and rat model organisms) are numerous and well known metabolites. The metabolism of IMI is complex and has been studied in goats, rodents and laying hens (Tomizawa and Casida, 2005). Two major routes are present for IMI metabolism in the rat as shown in Fig. 8 (Thyssen and Machmer, 1999). In the first route, IMI undergoes oxidative cleavage to imidazolidine and hydrolysis to 6CNA. The second route involves hydroxylation in the imidazolidine ring (formation of 4- and 5-hydroxy-IMI), followed by elimination of water and formation of olefin-IMI (JMPR, 2001; CDPR, 2006).

Table 2: Chemical structure, IUPAC name and selected properties of four major IMI's TPs: 6CNA, desnitro-IMI, olefin-IMI and 5-hydroxy-IMI.

IMI's TP	CAS number and chemical structure	IUPAC name	Molecular weight (g mol <sup>-1</sup> )	Solubility $(g L^{-1})$
6CNA	5326-23-8 COOH	6-chloropyridine- 3-carboxylic acid	157.6	2
desnitro-IMI	115970-17-7  **NH2 NH	1-(6- chloropyridin-3- ylmethyl) imidazolidin-2- ylideneamine hydrochloride	247.13	180-230
olefin-IMI	1115248-04-8 N_NO <sub>2</sub>	1-(6-chloro-3- pyridylmethyl)- <i>N</i> - nitro-1,3-dihydro- 2 <i>H</i> -imidazol-2- ylideneamine	253.65	> 0.51
5-hydroxy- IMI	1115248-02-6 CI N N H OH	(E)-3-(6-chloro-3- pyridylmethyl)-2- (nitroimino) imidazolidin-4-ol	271.66	> 0.51

Different studies performed with IMI and its TPs indicated that the desnitroor guanidine-TPs may be activators of toxicity in mammals and detoxification
products in insects, while the opposite is true for the olefin- and nitrosoimine-TPs
(Schulz-Jander and Casida 2002; Schulz-Jander *et al.*, 2002). One of these TPs,
desnitro-IMI (Table 2), is of particular interest, because it presents preference for
mammalian versus insect nAChRs and could be related with possible toxic effect
also in humans (Tomizawa and Casida, 1999; Tomizawa and Casida, 2000).
Important enzymes in IMI biotransformation are cytochrome 450 (liver microsomal
CYP450) and cytosolic aldehyde oxidase (AOX) (Honda *et al.*, 2006). CYP3A4

oxidises the imidazolidine moiety and reduces the nitroguanidine substituent (Schulz-Jander and Casida 2002; Schulz-Jander *et al.*, 2002) (Fig. 7), while AOX reduces the nitroguanidine moiety of IMI (Schulz-Jander *et al.*, 2002; Dick *et al.*, 2005). Metabolism of the neonicotinoid can decrease or increase its toxicity/potency, based on the compound and specificity of the nAChR.

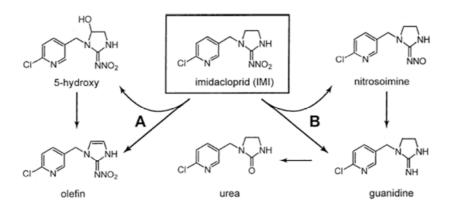


Figure 8: IMI metabolism by human CYP450 (A) hydroxylation or desaturation of imidazolidine moiety with formation of hydroxyl- and olefin-IMI; (B) cleavage and reduction of nitroimine substituent to form urea, nitrosoimine and guanidine metabolites. (Schulz-Jander and Casida, 2002).

The safe and effective use of pesticides requires a clear understanding of the parent compound, but also of its TPs. All legislations related to the placement of pesticides on market focus only on parent compounds rather than marketed formulations or TPs. No particular control or monitoring over the presence of IMI TPs is performed and there is a great necessity to implement this part of legislation by establishing the MRLs levels for several TPs present in the environment. Although IMI is continuously studied, fewer investigations have been conducted on its TPs. There is a single study relative to the toxicity of 6CNA on aquatic invertebrates performed on midge *Chironomus tentans* with LC<sub>50</sub> (96 h) higher than 1 mg L<sup>-1</sup> (Bowers and Lam, 1988). There is a great need to improve our knowledge relative to IMI's TPs by understanding their possible toxic effects to non-target organisms. In this study in addition to the selected parent compound **IMI** and its **commercial formulation Confidor 200SL**, four major TPs of IMI: **6CNA**, **olefin-IMI**, **desnitro-IMI** and **5-hydroxy-IMI** were also included (Fig. 4).

# 1.4. Pesticide effects on different levels of biological organisation

Pesticides released into the environment may have several adverse ecological effects ranging from long- to short-term effects in the ecosystem. Pesticides hold a distinctive position among environmental contaminants due to their acute and chronic toxicity. After the application of the pesticides (with spraying through sprinkling or by wetting the seeds or crops); the pesticides can influence different soil or aquatic organisms (Fig. 9). The indirect influence of pesticides to this group of organisms called **non-target organisms** should not be neglected. Compounds currently applied in agriculture are biologically very active and their specificity could not be improved to such extent that effects on non-target species can be avoided (Schulz *et al.*, 2002).

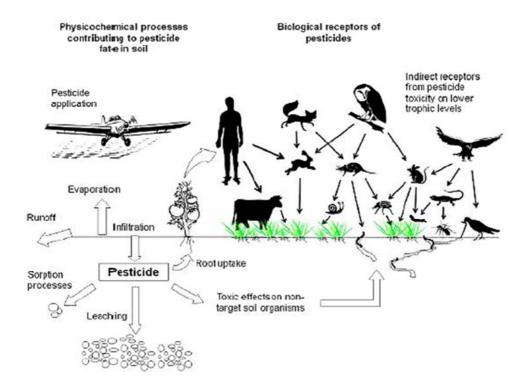


Figure 9: The general scheme of pesticides environmental fate and influence of different organisms in the environment (Sanchez-Hernandez, 2011).

The toxicity of a pesticide is determined by quantifying **biomarkers** (or **conventional endpoints**) of experimental organisms to a series of increasing chemical's doses in different **toxicological bioassays**. This relationship between administered dose and organism response is graphically represented as the doseresponse curve (Fig. 10).

A common measure used to define toxicity at a certain dose is the  $LD_{50}$  ( $LC_{50}$ ) – median lethal dose or concentration at which more than 50 % of organisms die. The  $EC_x$  is the effective concentration of test sample at which x % (e.g. 10, 20 or 50 %) of its maximum effect is reached.

The specific point on the dose-response curve where the more susceptible animals are first affected by a pesticide dose is termed the **threshold level**. The threshold level is the beginning of the linear response region of the curve and is the separation between the "No Observed (Adverse) Effect Level/Concentration" (NO(A)EL/NO(A)EC) and the "Lowest Observed (Adverse) Effect Level/Concentration" (LO(A)EL/LO(A)EC). The NO(A)EC value is the highest tested concentration that does not yet cause a statistically significant effect compared to the control, while the LO(A)EC value is the lowest tested concentration that elicits a statistically significant effect compared to the control.

In addition, the effects of a pesticide vary not only with the concentration/dose but also with the duration of exposure that can be:

- acute (short-term exposure with high concentrations)
- sub-chronic (intermediate-term exposure)
- chronic (long-term exposure with low concentrations)

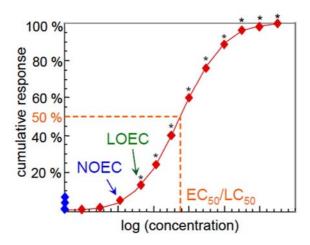


Figure 10: General scheme of a dose response curve with the toxicity parameter NOEC, LOEC and  $EC_{50}$ .  $\Diamond$  Control,  $\Diamond$  treatment, \* significant difference compared to control (Kienle et al., 2011).

The use of NOEC and LOEC data in regulatory aspects of ecotoxicology has been severely criticised since a series of published data in the 1990s (Hoekstra and Van Ewijk, 1992; Noppert et al., 1994; Chapman et al., 1996; OECD, 1998). Although an OECD (1998) workshop recommended that NOECs should be discarded out of international standards, guidance and toxicity testing, NOEC and LOEC estimations still persist in more recent OECD statistical publications (2006) because "...the NOEC is still required in many regulatory standards from many countries and in some cases where a detailed determination of an ECx is not relevant and the alteration of the study design is too costly to fullfil the requirements for regression models". In general, despite these criticisms, NOECs and LOECs are continuously generated and reported regularly in the literature since no standardised alternative is completely accepted.

Environmental risk assessment (ERA) of environmental chemicals (e.g. pesticides) is predominantly based on the results of laboratory toxicological bioassaya and studies as previously explained. This assessment is typically based on a comparison of effect estimations and exposure concentrations. ERA is completed in four steps: hazard identification, dose-response assessment, exposure assessment, and risk characterisation (EC, 2003). Besides ERA, there is another interesting term in this field known as: Integrated Pest Management (IPM) which was formalised by

the US Academy of Sciences in 1969. IPM is largely a combination of the integrated control, which combines biological and chemical control of pests with pest management. In addition, the European Commission is promoting low pesticide-input farming in Member States with the application of IPM. IPM relies on minimising pesticide use through the implementation of alternative methods to control pests, diseases and weeds. Community-wide standards for IPM are being developed and will be mandatory across the EU from 2014 (Hillocks, 2012).

In general, toxicity testing is conducted to determine the potential human health and environmental hazards of chemicals (e. g. pesticides) and their products. Human toxicology, by definition, is concerned with the effects of chemicals on a single species: man. In this case the research and testing are undertaken with one major objective, protection of the health of the individual. Pesticides are considered as a significant source of diverse pollutants that can cause health implications in humans (Hellweg and Geisler, 2003). Contrary to most other chemicals, they are deliberately emitted to the environment to control undesirable organisms, but can also reach non-target life forms such as humans. We are exposed to pesticides in the food, water and air we use and in these cases the duration of exposure, the role of pollutant mixtures, the mode of action should be evaluated and pesticide use in agriculture must be subjected to continuous monitoring.

In contrast to the human toxicology is ecotoxicology, which not only deals with the impact of chemicals on individuals of different species, but also deals with the influence of chemicals on supra-organismal levels. Ecotoxicology has evolved mainly from three different disciplines: toxicology, applied ecology and environmental chemistry. It as an interdisciplinary environmental science that deals with the interactions between environmental chemicals and biota, thereby focusing on adverse effects at **different levels of biological organisation** from the molecular, cellular, and organism level, up to populations and ecosystem (Fig. 11). Ecosystems include a large numbers of species and pesticides effect may be identified at different biological levels, i.e. individuals of a species, populations of species and groups of species (communities) with trophic links. Pesticide effect can be simple/direct or indirect due to complex food web interactions at higher levels of biological integration (Preston and Snell, 2001; Preston, 2002). Direct effects may depend on

the mode of action of the compound (Barata and Baird, 2000), the concentrations (Naddy *et al.*, 2000), and duration of the exposure (Reynaldi and Liess, 2005). These straight-forward effects of pesticide exposure on susceptible species include lowering of survival rate (Barry *et al.*, 1995), or effects on different sub-lethal endpoints such as food consumption, growth (Hooper *et al.*, 2005), and brood production (Hosmer *et al.*, 1998). Pesticide exposure affects swimming behaviour, filtration rates as well as predator-prey relationships (Dodson *et al.*, 1995). After pesticide application was noted also reduction in diversity, density, and biomass of many invertebrate species in aquatic communities (Berenzen *et al.*, 2005; Schafer *et al.*, 2007).

Currently there are present two main approaches that reflect the trends in ecotoxicological researches. One way is to work in real ecosystems which are represented by simplified but still realistic model systems like mesocosms. In these models are simulated and investigated complex interactions and feedback mechanisms which occur in ecosystems. An alternative approach is to develop a mode of action based effect evaluation (Escher et al., 1997) by investigating responsible molecular mechanisms of toxic action. These modes of action can be solved using whole-organism (in vivo) or in vitro model systems in order to develop and measure specific biomarkers that can be applied as early warning signals. In eco-toxicological research, cellular and biochemical effect studies including mechanisms of toxic action are equally important as studies in cell culture and whole organism models because the primary interaction between chemicals and biota occurs at the surface of/or in cells. In this way, cellular toxicology provides an essential concept in understanding eco-toxicological processes, as it plays a key role in identifying toxicological effects at higher biological levels. Taken together, these two described approaches integrate each other and will finally improve the understanding of the effects of pollutants (e.g. IMI) on living systems (biota and humans).

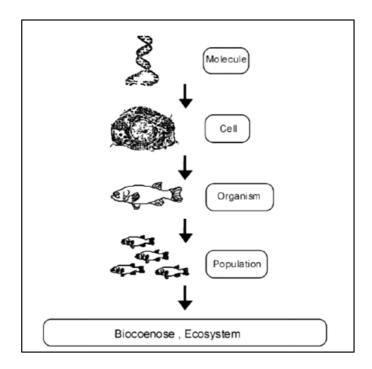


Figure 11: Ecotoxicology, an environmental science focusing on different levels of biological organisation (Fent, 2001).

### 1.4.1. Imidacloprid - toxicity studies on non-target aquatic organisms

IMI can contaminate surface- and ground-water by runoff or leach from agricultural areas and lead to pulse or localised pesticide contaminations (Fossen, 2006; Gupta *et al.*, 2002). Detected aquatic concentrations indicate measured levels of IMI going from 14  $\mu$ g L<sup>-1</sup> up to 0.3 mg L<sup>-1</sup> for surface waters, while the estimated concentrations for accidental spills reach high values going from 1.8 mg L<sup>-1</sup> up to 7.3 mg L<sup>-1</sup> (Table 1).

IMI based on its mode of action is in specific toxic to insects and aquatic insects; more than to other aquatic invertebrates (Overmyer *et al.*, 2005), crustaceans and fish (Tišler *et al.*, 2009). Freshwater fish do not appear to be particularly sensitive to IMI, with toxic effects occurring at relatively high concentrations over 80,000 μg L<sup>-1</sup> (CCME, 2007). In specific, only the early life stages of fish exhibit higher sensitivity (Cox, 2001). In case of freshwater plants there is evidence that IMI may be harmful to plants and algae at higher concentrations (Table 3). The acute toxicity of IMI was tested on different species of **green algae** (Heimbach 1989; Gagliano and Bowers 1991). These studies gave a NOAEC and EC<sub>50</sub> values for biomass and growth higher than 119 mg L<sup>-1</sup> a. i. for 5 day test with *Selanastrum capricornutum* and 10 mg L<sup>-1</sup> a.i. for *Scenedesmus subspicatus*. In comparison, a 4 day NOEC of 6.69 mg L<sup>-1</sup> and 4 day LOAEC of 9.98 mg L<sup>-1</sup> was determined for the diatom (*Navicula pelliculosa*) after exposure to IMI formulation with 21.6 % of a. i. (Hall, 1996). In general, it appears that algae are at least three orders of magnitude less sensitive to IMI than many insect and ostracod species.

Furthermore, the major part of studies referring to the persistence and toxicity of IMI in the aquatic environment is focused to different **non-target aquatic invertebrates** (Table 4). IMI induces toxic effects in several aquatic invertebrates at different concentrations having a species-specific action and effects. Using survival and growth endpoints a 96 h and 10 day NOEC values have been determined at 1.24 μg L<sup>-1</sup> a. i. and 0.67 μg L<sup>-1</sup> a. i., respectively for *Chironomus tentans* (Gagliano, 1991). Effects at such low concentrations of IMI were reported also by Stoughton (2008) for the same species, with a LC<sub>50</sub> for 96 h of 5.75 μg L<sup>-1</sup> a. i. The sensitivity of crustaceans to IMI appears to vary. A recent study noted that ostracod species seem to be more sensitive than cladocerans e.g. water flea *Daphnia magna* (Sánchez-

Bayo and Goka, 2006). Amphipods appear to have intermediate sensitivity in comparison with other crustaceans. Stoughton (2008) reported in case of the amphipod *Hyallela azteca* a LC<sub>50</sub> for 96 h of 65.4 μg L<sup>-1</sup> of a. i. IMI. For the amphipod *Gammarus pulex*, a 28 day LOEC for mortality endpoint was determinate at 256 μg L<sup>-1</sup> of a. i. IMI (Hendel, 2001). In a 19 week microcosm study IMI was applied to the surface of tanks containing a variety of phytoplankton, zooplankton, and macro-invertebrates at two week intervals with concentrations ranging from 0 to 0.180 mg L<sup>-1</sup> a. i. Statistically significant decreases in populations of total macro-invertebrates as well as individual taxa (mayflies, midges, beetles and amphipods) were most frequently observed at IMI's concentrations ranging from 0.02 to 0.180 mg L<sup>-1</sup> a. i.

All investigations span from single species toxicity tests in laboratory (Jemec et al., 2007; Chen et al., 2010; Lukančič et al., 2010; Azevedo-Pereira et al., 2011) to complete indoor/outdoor stream mesocosms studies under constant exposure (Pestana et al., 2009) and short-pulse exposure conditions (Stoughton et al., 2008; Mohr et al., 2012). The authors observed usually modifications in survival, behaviour and population growth rate; while some of them evaluated biochemical alterations at molecular level and compared the toxic effects of pure compound IMI and its formulated version (Jemec et al., 2007; Chen et al., 2010).

Table 3: Selection of some representative IMI toxicity studies on freshwater algae.

Test organism	Exposure/measured endpoint	Toxicity data	References
Green algae Scendesmus subspicatus	<ul> <li>sub-chronic toxicity</li> <li>concentrations of 0,</li> <li>0.1, 1, and 10 mg L<sup>-1</sup></li> <li>biomass and growth</li> </ul>	72 and 96 h EC <sub>50</sub> : $> 10 \text{ mg L}^{-1}$ 72 and 96 h NOAEC: 10 mg L <sup>-1</sup>	Heimbach, 1989
Green algae Selanastrum capricornutum	<ul> <li>chronic toxicity</li> <li>nominal (measured)         concentrations of 0, 15.6         (14.1), 25.9 (24.1), 43.2         (41.1), 72 (69.5), and 120         (119) mg L<sup>-1</sup></li> <li>biomass and growth</li> </ul>	EC <sub>50</sub> (5 d): >119 mg L <sup>-1</sup> NOAEC (5 d): >119 mg L <sup>-1</sup>	Gagliano and Bowers 1991
Green algae Pseudokirchneriella subcapitata	<ul><li>chronic toxicity</li><li>growth inhibition</li></ul>	EC <sub>50</sub> (5 d) and NOEC: < 119 mg L <sup>-1</sup>	Gagliano and Bowers, 1991
Diatom Navicula pelliculosa	<ul> <li>sub-chronic toxicity</li> <li>concentrations: 0, 0.16, 0.42, 1.05, 2.64, 6.69, and 17.0 mg L<sup>-1</sup></li> <li>growth inhibition</li> </ul>	4 d NOAEC: 6.69 mg L <sup>-1</sup> 4 d LOAEC: 9.88 mg L <sup>-1</sup> 4 d EC <sub>50</sub> : 12.37 mg L <sup>-1</sup>	Hall, 1996
Blue-Green algae Anabaena flosaquae	<ul> <li>sub-chronic toxicity</li> <li>concentrations of 0, 24.9, 40.5, 68.2, 121.3, and 193.3 mg L<sup>-1</sup></li> </ul>	EC <sub>50</sub> (4 d): 32.8 (30.4-34.6) mg L <sup>-1</sup> NOEC (4 d): 24.9 mg L <sup>-1</sup>	Bowers, 1996; Mulye, 1997

Table 4: Selection of some representative IMI toxicity studies on freshwater invertebrates.

Test organism	Exposure/measured endpoint	Toxicity data	References
ACUTE/SUB-CHRONIC TOXICITY			
Water Flea Daphnia magna	<ul> <li>static 48 h acute toxicity test</li> <li>3 trials, 4 replicates per concentration, 10 animals</li> </ul>	Water Flea: LC <sub>50</sub> (48 h): 10.44 mg L <sup>-1</sup>	Song et al., 1997
Mosquito Aedes aegypti	<ul><li>each species per replicate</li><li>mortality</li></ul>	Mosquito: LC <sub>50</sub> (48 h): 0.044 mg L <sup>-1</sup>	Song and Brown, 1998
Amphipod crustacean  Hyalella azteca	<ul> <li>static sub-chronic toxicity test</li> <li>2 replicates per concentration, 10 per replicate</li> <li>mortality, immobilisation, abnormal effects (e.g. lethargy or surfacing)</li> </ul>	(a) $LC_{50}$ (96 h): 0.526 mg $L^{-1}$ $EC_{50}$ (96 h) for immobilisation : 0.055 mg $L^{-1}$ NOAEC (96 h) for immobilisation and abnormal effects: 0.00035 mg $L^{-1}$ (b) $LC_{50}$ (96 h) for juveniles: 65.43 $\mu$ g $L^{-1}$ (IMI) $LC_{50}$ (96 h) for juveniles: 17.44 $\mu$ g $L^{-1}$ (Admire) NOEC (96 h) similar for the two products; 54.24 $\mu$ g $L^{-1}$ (IMI); 48.75 $\mu$ g $L^{-1}$ (Admire).	(a) England and Bucksath, 1991  (b) Stoughton et al., 2008
Midge Chironomus tentans	<ul> <li>static sub-chronic toxicity test</li> <li>2 replicates per concentration, 10 chironomids per replicate</li> <li>mortality</li> </ul>	LC <sub>50</sub> (96 h): 0.0105 mg L <sup>-1</sup> NOAEC (96 h) for survival: 0.00124 mg L <sup>-1</sup>	Gagliano, 1991
Water louse Asellus aquaticus L.  Stream scud Gammarus fossarum Koch.	<ul> <li>static acute 48 h toxicity test</li> <li>mortality</li> <li>electron transport system (ETS) activity and respiration level</li> </ul>	Water louse (a.i. in Confidor 200SL): $LC_{50}$ (48 h): 8.5 mg $L^{-1}$ $EC_{50}$ (24 h): 0.8 mg $L^{-1}$ Stream scud (a.i. in Confidor 200SL; 20 % of a. i. IMI). $LC_{50}$ (48 h): 0.8 mg $L^{-1}$ $EC_{50}$ (24 h): 0.07 mg $L^{-1}$	Lukančič et al., 2010

Table 4: Selection of some representative IMI toxicity studies on freshwater invertebrates (continued).

Test organism	Exposure/measured endpoint	Toxicity data	References
ACUTE/SUB-CHRONIC TOXICITY  Midge Chironomus tentans	<ul> <li>constant sub-chronic toxicity test with larvae</li> <li>mortality</li> </ul>	LC <sub>50</sub> (96 h): 5.75 μg L <sup>-1</sup> (IMI) LC <sub>50</sub> (96 h): 5.4 NOEC (96 h): 1.03 μg L <sup>-1</sup> (IMI) and 5.11 μg L <sup>-1</sup> (Admire) The difference between the effects of IMI and Admire (commercial formulation) was not significant	Stoughton et al., 2008
Microcrustacean Ceriodaphnia dubia Richard	<ul><li>acute toxicity test</li><li>survival</li><li>population size</li></ul>	LC <sub>50</sub> (48 h): 2.07 μg L <sup>-1</sup> (Admire Pro; 42.8% a. i. IMI) 8 days' exposure to a mixture of the nonylphenol polyethoxylate, R-11 and IMI resulted in a population size 3 times smaller than with R-11 alone, and 13 times smaller than with IMI only	Chen et al., 2009
Planktonic cladoceran Chydorus sphaericus	<ul><li>acute toxicity</li><li>mortality</li></ul>	LC <sub>50</sub> (24 h): 161.95mg L <sup>-1</sup> LC <sub>50</sub> (48 h): 132.67 mg L <sup>-1</sup> EC <sub>50</sub> was also determined, and was 2-13 times lower under dark than under normal (16 h light : 8 h dark) laboratory conditions	Sánchez-Bayo and Goka, 2006
Water flea Daphnia magna	<ul> <li>chronic 21 d toxicity test</li> <li>4 replicate per concentration, 6 daphnids per replicate</li> <li>mortality, immobilisation</li> </ul>	(a) EC <sub>50</sub> (21 d) for immobilisation: >7.3mg L <sup>-1</sup> LOAEC: 3.6 mg L <sup>-1</sup> 3.6 and 7.3 mg L <sup>-1</sup> : Significantly reduced adult daphnid length compared with control 7.3 mg L <sup>-1</sup> : significantly reduced survival; significantly reduced reproduction compared with control	(a) Young and Blakemore, 1990

Table 4: Selection of some representative IMI toxicity studies on freshwater invertebrates (continued).

Test organism	Exposure/measured endpoint	Toxicity data	References
CHRONIC TOXICITY		(b) NOEC (21 d): 1.25 mg L <sup>-1</sup>	(b) Tišler <i>et al.</i> , 2009
		The toxicity of commercial formulation Confidor SL 200 was intensified in comparison to the a.i. (c) $LC_{50}$ (10 d): 9 mg $L^{-1}$ $LC_{50}$ (28 d): 0.91 µg $L^{-1}$ (a.i. in Admire)	(c) Sánchez-Bayo and Goka, 2006
Midge Chironomus tentans	<ul><li>chronic 28 d toxicity test</li><li>constant exposure</li></ul>	NOEC (28 d): 1.14 μg L <sup>-1</sup> (Admire) NOEC (96 h pulse exposure): 3.47 μg L <sup>-1</sup> after 10 d	Stoughton et al., 2008
Amphipod crustacean Hyalella azteca	<ul><li>chronic 28 d toxicity test</li><li>constant exposure</li></ul>	LC <sub>50</sub> (28 d): 7.08 μg L <sup>-1</sup> NOEC (28 d constant exposure): 3.44 μg L <sup>-1</sup> (Admire) NOEC (96 h pulse exposure): 11.93 μg L <sup>-1</sup> after 10 d	Stoughton et al., 2008
mesocosmos  phytoplankton, zooplankton, macroinvertebra tes (Hyalella azteca)	<ul> <li>19 week microcosm study</li> <li>4 surface applications at 2-week intervals</li> </ul>	Amphipods were the most sensitive species, significant decrease in populations of total macroinvertebrates and individual macroinvertebrate taxa. Study recommended 0.006 mg L <sup>-1</sup> as NOEC for regulatory action.	Morning et al., 1992
Macroinvertebrates	<ul> <li>20 d mesocosmos experiment</li> <li>three pulses of IMI (duration of 24 h)</li> </ul>	IMI pulses reduced invertebrate abundance and community diversity compared to control.  Reduction of oxygen consumption in stoneflies.	Pestana et al., 2009
Macrozoobenthos community	<ul> <li>3-week mesocosmos experiment</li> <li>3 IMI pulses of 12 h (1 week apart)</li> </ul>	Caddisfly <i>Neureclipsis</i> sp. reacted after first pulse and most sensitively.  Insect larvae (ephemerids and dipteran) negatively affected after repeated pulses.  Effects on insects larvae more pronounced in summer, due to increased temperature.  Emergence is the most sensitive endpoint.	Mohr et al., 2012

### 1.4.2. Imidacloprid - toxicity studies on non-target mammalian models

Toxicity testing on mammalian experimental models is necessary to completely elucidate possible IMI adverse effects, specific mechanisms of actions and molecular pathways. **Humans** are also a **non-target group** which could be affected by IMI action through common routes of exposure. Diet is the dominant exposure pathway for pesticides (McKone *et al.*, 2007; Lu *et al.*, 2008). In general, food ingestion, air inhalation and ingestion of drinking water are possible ways of insecticide intake by general public. In addition, agricultural workers are one of the particularly exposed groups to acute (high) and chronic (low) concentration of pesticides after continuous applications, spraying and accidental spillage. Exposure to pesticides has detrimental effects on human health and is considered as the major contributor to most diseases of great public significance.

According to World Health Organisation (WHO) IMI is considered as moderately toxic and its classified by US EPA system as toxicity class II and class III agents because it blocks specific neuron pathway which are more abundant in insect than warm blooded animals. However, several reports have shown high clinical toxicity of IMI in human suicidal attempts (Wu et al., 2001; Proenca et al., 2005; David et al., 2007; Shadnia and Moghaddam, 2008). The oral LD<sub>50</sub> of IMI is of 450 mg/kg body weight (b.w.) in rats and 131 mg/kg b. w. in mice (Najafi et al., 2010). Experimental animals presented signs of toxicity at doses lower than those causing mortality, regardless of the species, formulation or TP administered. In most studies, clinical signs of toxicity, including staggering, sedation, apathy, tremors and convulsions were observed shortly after dosing; these signs were typically resolved in all animals before the end of a study (within 14 days). Temporary decrease in body weight was also a common symptom of IMI-treated animals. In fact, the principal toxicological finding in sub-chronic and chronic oral testing was the reduction in body weight. Monitoring of body weight provides information on the general health status of animals which can be also important explanation of reproductive effects (Aly et al., 2009). Moreover, the liver was the principal target organ as demonstrated by the hepatic necrosis or hypertrophy in rats and dogs, elevated activities of serum enzymes, hepatic mixed-function oxidase and alteration of clinical chemistry parameters such as triglycerides, cholesterol and the blood clotting time (CDPR, 2006). There are some reports showing IMI being negative in a battery of genotoxicity tests such as test for DNA damage and repair capacity, or *in vitro* gene mutation tests and *in vivo* chromosomal aberration tests (JMPR, 2001). On contrary, in some mammalian *in vitro* tests IMI caused sister chromatid exchange and chromosomal aberrations at doses which confirmed to have also cytotoxic effect (JMPR, 2001). Regarding reproductive and developmental toxicity for IMI were reported disproportionally high number of male foetus and lower foetal body weight in rats (CDPR, 2006). The offspring of treated rat mothers exhibited also significant sensory-motor impairments after 30 post-natal days and this effect was associated with increased AChE activity in brain and in plasma (Abu-Donia *et al.*, 2008). An acute oral neurotoxicity study performed by Sheets (1994) suggested that neurobehavioral and pathological effects are seen only after high dose exposure to IMI, while all noted effects at lower doses were not on the nervous system (SERA, 2005).

Throughout the scientific literature there are present different studies with mammalian models relative to IMI toxicity and induced morphological and pathological alterations. These studies report LD<sub>50</sub> as well as relative NOEC and LOEC values for different measured endpoints. Only recently the research in this area is also focusing on IMI's mediated toxicity which involves excessive production of reactive oxygen species (ROS), induced oxidative stress and different activated molecular pathways. These current studies include experimental organisms and different mammalian or other cellular models (Table 5).

Table 5: Selection of some representative IMI toxicity studies on mammalian organisms and different neuronal cell models.

Test organism	Exposure/measured endpoint	Toxicity data	References
Central nervous system and liver of rats	• injected intravenously; 2 h exposure at 3/4 of LD <sub>50</sub> (424-475 mg/kg/b.w.)	Induction of nitric oxides in liver. Significant LP. Different responses of CAT, SOD, GPx. Upregulation of inflammatory cytokines. Results	Duzguner and Erdogan, 2010
rats	<ul> <li>antioxidant enzymes</li> <li>lipid peroxidation (LP)</li> <li>nitric oxide induction</li> </ul>	suggested that IMI cause oxidative stress and inflammation in central nervous system and liver.	Lidogan, 2010
Male mice	<ul> <li>oral administration; 24 h exposure to 1/10 of LD<sub>50</sub></li> <li>antioxidant enzymes</li> <li>lipid peroxidation (LP)</li> <li>survival rate (LD<sub>50</sub>)</li> </ul>	Increase in LP level and the activities of antioxidant enzymes including CAT, SOD, GPx and GST. G6PD activity remained unchanged, while the level of GSH content was decreased. Vitamin C might ameliorate IMI-induced oxidative. Induction of oxidative stress is perhaps the central mechanism by which this pesticide exert its toxic effect.	El-Gendy <i>et al.</i> , 2010
Female rats (ovaries)	<ul> <li>daily oral administration; 90 days of exposure to 5, 10 and 20 mg/kg/day)</li> <li>morphology</li> <li>hormones</li> <li>antioxidant enzymes</li> </ul>	Significant alterations of SOD, CAT, GPx, GSH and LP (at 20 mg/kg b.w.). Significant toxicological impact on ovary of female rats as evident by pathomorphological changes, hormonal imbalance and generating oxidative stress.	Kapoor <i>et al.</i> , 2011
Male rats	<ul> <li>daily oral administration; 60 days of exposure to 225 and 112 mg/kg b.w.</li> <li>behaviour</li> <li>hormone</li> <li>tissue morphological analyses</li> </ul>	Decreased movement, staggering, trembling, diarrhoea and spasms. Increased thickness of <i>tunica albuginea</i> , edema in connective tissue, cytoplasmic granulation. Noted decrease in serum testosterone levels.	Najafi <i>et al.</i> , 2010

Table 5: Selection of some representative IMI toxicity studies on mammalian organisms and different neuronal cell models (continued).

Test organism	Exposure/measured endpoint	Toxicity data	References
ORGANISM MODEL	<ul> <li>daily oral administration; 90</li> </ul>	IMI at 5 and 10 mg/kg/day has not produced any	
Female rats	days of exposure to 5, 10 (NOEL) and 20 mg/kg/day)  SOD, CAT, GPx,GSH and LP in liver, brain and kidney	changes in SOD, CAT, GPx, GSH and LP. Changes of these parameters were significant at 20 mg/kg/day in liver. Also, alterations of SOD, CAT, GPx in brain and LP in kidney. 10 mg/kg/day considered as NOEL through antioxidant enzymes and LP.	Kapoor et al., 2010
Female rats	<ul> <li>daily oral administration; 90 days of exposure to 5, 10 (NOEL) and 20 mg/kg/day)</li> <li>survival, body weight and biochemical parameters</li> </ul>	Decrease in the body weight at 20 mg/kg/day and at necropsy body weights of liver, kidney and adrenal were increased. Elevation of serum enzymes, glucose and decreased the activity of AChE in serum and brain.	Bhardwaj et al., 2010
MAMMALIAN NEURONAL CELLS  Mouse neuroblastoma N1E-115 cell line	<ul> <li>30 min (0-180 min) incubation (0.0001-1000 μM)</li> <li>ERK kinase cascade</li> <li>intracellular calcium (Ca<sup>+2</sup>) mobilisation</li> </ul>	IMI, desnitro-IMI and nicotine induced ERK activation in a dose-dependent manner triggered by primary action at the α4β2 nAChR with an involvement of intracellular Ca <sup>+2</sup> mobilisations. Intracellular Ca <sup>+2</sup> activate a sequential pathway from protein kinase C (PKC) to ERK.	Tomizawa and Casida, 2002
Neuronal SH-SY5Y and pulmonary A549 cell lines	<ul> <li>3 day incubation (0.2-2.7 mM)</li> <li>growth rate (LOEC, IC50)</li> <li>cell count, total protein content</li> <li>heat shock proteins (HSP)</li> <li>glucose regulated proteins</li> </ul>	Neuronal cells more sensitive than pulmonary. For SH-SY5Y: LOEC – 1.2 mM (IMI) and 0.3 mM (Confidor 200SL); IC <sub>50</sub> – 1.6 mM (IMI) and 0.8 mM (Confidor 200SL). For A549: LOEC – 1.2 mM (IMI) and 0.3 mM (Confidor 200SL); IC <sub>50</sub> – 1.8 mM (IMI) and 1.3 mM (Confidor 200SL). Commercial formulation (Confidor 200SL) indicates an additional additive toxic effect. Insecticide concentrations higher than IC <sub>50</sub> were found to induce an under-expression of all cytosolic HSP probably resulting from a general inhibition of protein synthesis	Skandrani <i>et al.</i> , 2006

*Abbreviations:* IC<sub>50</sub> (median inhibition concentration).

Table 5: Selection of some representative IMI toxicity studies on mammalian organisms and different neuronal cell models (continued).

Test organism	Exposure/measured endpoint	Toxicity data	References
MAMMALIAN NEURONAL CELLS	3 day incubation	IMI up-regulate α4β2 AchRs by five- to eightfold	
Mouse fibroblast M10 cells	<ul> <li>up-regulation of α4β2 AchRs and mechanisms of up-regulation</li> </ul>	with EC <sub>50s</sub> of 70,000 and 19,000 nM. On contrary, desnitro-IMI TP gave up-regulation by eightfold and EC <sub>50s</sub> of 870 and 500 nM, respectively. Determinated IC <sub>50</sub> for IMI was of 2600 nM and for densitro-IMI of 8.2 nM.	Tomizawa and Casida, 2000a
Primary cultures of cerebellar neurons from neonatal rats	<ul> <li>excitatory Ca<sup>2+</sup> influx assay</li> <li>applications of doses between 1, 10 and 100 μM</li> </ul>	Significant excitatory Ca <sup>2+</sup> influxes were evoked at concentrations greater than 1 μM  The firing patterns, proportion of excited neurons, and peak excitatory Ca <sup>2+</sup> influxes showed differences from those induced by nicotine.	Kimura-Koroda, 2012
Cholinergic synapses of the stellate cells of the mouse cochlear nucleus	<ul> <li>determine to what extent IMI affects the nAChRs</li> <li>exposure doses based on EC<sub>50</sub> against mammalian nAChRs of 70 μM</li> <li>whole-cell patch-clamp recording</li> </ul>	Puff application of 1 µM had no significant effect on the membrane properties of the neurons, while a concentration of 10 µM caused a significant depolarising shift in the membrane potential. Concentrations >50 µM caused a significant depolarising shift in the membrane potential. Exposure to IMI at concentrations >10 µM for <1 min can change the membrane properties of neurons that have nAChRs and, as a consequence, their function.	Bal et al., 2010

Table 5: Selection of some representative IMI toxicity studies on mammalian organisms and different neuronal cell models (continued).

Test organism	Exposure/measured endpoint	Toxicity data	References
OTHER NEURONAL CELL MODELS  Cultured GFP tagged cholinergic neurons from the third instar larvae of the genetic model organism Drosophila melanogaster	<ul> <li>5 s applications of 100 μM dosed at 5 min intervals</li> <li>fura-2-based calcium imaging</li> </ul>	Dose-dependent increase in intracellular Ca <sup>+2</sup> . Increase in Ca <sup>+2</sup> involve also voltage gated calcium channels.	Jepson et al., 2006
Neural Kenyon cell of cockroach Periplaneta americana	<ul> <li>incubation for 1, 3, 5, 8 and 24 h at 1, 5, 10, 50 and 100 μM</li> <li>cell viability (Trypan blue)</li> <li>adenylate kinase activity (AK)</li> </ul>	Significant decrease of cell viability after IMI exposure. Increase of AK activity. Higher concentrations of IMI needed for intracellular mechanisms of IMI toxicity.	Benzidane et al., 2011
Neurons isolated from the three thoracic ganglia of the cockroach, <i>Periplaneta americana</i>	<ul> <li>application of 100 μM for 20 s</li> <li>agonist actions of on nAchRs</li> <li>single electrode voltage clamp recording</li> </ul>	Strong excitation symptoms followed by prostration and death. Relatively weak partial agonists causing only 20–25 % of the maximum ACh current. Agonist efficacy, but not affinity, was positively correlated with insecticidal activity.	Tan et al., 2007
Central nervous system neurons of the stick insect (Carausius morosus)	<ul> <li>Brief U-tube application of 10<sup>4</sup> M doses for 1 s</li> <li>patch-clamp experiments</li> <li>effects on nAchRs</li> </ul>	Generally depressive symptoms, characterised by stillness and weakness, while also variably inducing postural changes such as persistent ovipositor opening, leg flexion or extension and abdomen bending that could indicate excitation of certain neural circuits.  Selective desensitisation of certain nAChR subtypes can account for the insecticidal actions of IMI.	Oliveira et al., 2011

# 1.4.3. Imidacloprid's transformation products - toxicity studies on non-target organisms

No particular monitoring over the presence of **IMI TPs** is performed. Although IMI is continuously investigated, fewer studies have been conducted on its TPs (Table 6). Major part of studies on IMI's TPs is performed mainly on common honey bee Apis mellifera (Nauen et al., 2001; Suchail et al., 2001) (Table 6). Bees, including honey bees, bumblebees and solitary bees are the prominent and economically most important group of pollinators worldwide. There is a great concern regarding the decline of Apis mellifera across Europe and world. This event has been termed as colony collapse disorder (CCD) (Fig. 12). Although the supposed causes are still currently analysed, the extensive use of chemical pesticides, such as neonicotinoids (e. g. IMI) may have contributed to loss of pollinators. Laboratory studies demonstrate that IMI is acutely toxic to bees with LD<sub>50</sub> (48 h) values ranging from 3.7 to 230.3 ng bee<sup>-1</sup> (SERA, 2005). For chronic laboratory tests with IMI was demonstrated a high worker bee loss when honey bees consumed contaminated pollen at 40 µg kg<sup>-1</sup> and sugar water at 0.1, 1 and 10 µg L<sup>-1</sup> (Decourtye et al., 2001; 2003). Sub-lethal effects have been noted in bumblebees (members of the bee genus Bombus) at concentration that could be considered harmless and included altered foraging behaviour, lower reproduction and decreased pollination (Mommaerts et al., 2010). Additional sub-lethal effects consist of impairment of olfactory memory, influenced mobility, reduced learning, and a decrease in 'dancing', which leads to reduced foraging activity (Desneux et al., 2007). Sub-lethal dose as low as 0.1 ng of IMI per honeybee seemed also sufficient to disturb navigation in honeybees, loss of foragers and consequently loss of food supply to the colony which then continuous weakening (CST, 2003). All these facts seriously imply that concentrations of several parts per billion (ppb) in the diet of social insects can already harm them on colony level. A recently published review of Blacquière et al. (2012) summarises the data relative to the neonicotinoids in bees and stresses the fact that TPs of IMI also contribute to the toxicity to beneficial arthropods. At the moment, most studies were conducted on TPs of IMI in bees and showed that olefin-, dihydroxy- and hydroxy-IMI were more toxic based on oral LD<sub>50</sub> compared to urea-IMI and 6CNA (Nauen et al., 2001).



Figure 12: (A) Frames taken from colonies suffering from CCD with few bees and a large numbers of developing larval and pupae bees. (B) Frame taken from a healthy colony full of bees. (Oldroy, 2007).

Based on structural considerations, the following TPs of IMI may be of toxicological significance: 6CNA, 4- and 5-hydroxy IMI, olefin-IMI, desnitro-IMI and the nitrosoimine compound (a contaminant of IMI preparation and its metabolite). All of these compounds contain the 6-chloropyridinyl moiety and are included in the tolerances established for the IMI residues, although they should be considered separately.

The toxicity of the major TPs of IMI (6CNA, 4 - or 5-hydroxy-IMI and olefin-IMI) has not been fully tested in mammals (CDPR, 2006). Several TPs were tested for acute toxicity by oral administration to rats and for their ability to induce point mutations (mostly by Ames test with bacteria *Salmonella typhimurium*). The TPs showed moderate acute toxicity after oral administrations and clinical signs similar to those described after IMI exposure. Overall, they were found to be less toxic than the parent compound, and all tests for genotoxicity gave negative results (SERA, 2005). However, one of the TPs, desnitro-IMI (Table 2), is important, because its mode of action prefers mammalian *versus* insect nAChRs (Tomizawa and Casida, 2000), as already mentioned in section 1.3 of this manuscript. From a toxicological point of view, the formation of the TP desnitro-IMI is of particular interest, as this product presents a clearly higher toxicity to mammals than IMI and is produced as the major degradation product or metabolite in environment and human liver microsomes, respectively (CDPR, 2006).

The predominant TPs associated with toxicity in insects are olefin-, dihydroxy- and hydroxy-IMI. There is present a general IMI toxicity pattern of exposure which involves an immediate onset of neurotoxicity, followed by a delayed mortality, usually 4 h after exposure. It seems based on the data that unmodified IMI may be responsible for the primary neurotoxic effect, while olefin-, hydroxy- and dihydroxy-IMI which appear at approximately 4 h after exposure may be responsible for mortality (SERA, 2005).

Studies in invertebrates showed that the olefin- and hydroxy- compounds had similar acute toxicity as the parent compound, while 6CNA did not act as a nicotinic agonist (Nauen, *et al.*, 2001). None of the assessed IMI TPs (urea-IMI, desnitro-IMI and 6CNA) were as acutely toxic as a.i. IMI alone in tests with the midge (*C. tentas*) or amphipod (*H. azteca*) (Bowers 1996; Bowers and Lam 1998; Rooney and Bowers 1996; Dobbs and Frank 1996).

In general, there are a limited number of studies relative to the toxicity of IMI TPs to aquatic invertebrates (Table 6) which warrants expanding our knowledge in this field. To this end, it is necessary to investigate their effects on freshwater biota, especially on non-target aquatic organisms. In particular, these investigations should be performed as a part of the analysis of pesticide impact on vulnerable aquatic ecosystems.

Table 6: Selection of some representative IMI's TPs toxicity studies on non-target organisms.

Test organism/tested TP	Exposure/measured endpoint	Toxicity data	References
MAMMALIAN ORGANISMS  Rat (olefin-IMI)	<ul><li>survival</li><li>TP purity of 98 %</li></ul>	Acute toxicity LD <sub>50</sub> for male of 3500 and for female rats of 1100 mg/kg b.w.	JMPR, 2001
Rat (nitroso TP)	<ul><li>survival</li><li>TP purity of 98.1 %</li></ul>	Acute toxicity LD <sub>50</sub> for male of 1980 and for female rats of 3560 mg/kg b.w.	JMPR, 2001
Wistar rat (nitroso TP)	<ul> <li>15 males and 15 females</li> <li>0, 100, 300 and 1000 ppm for 12 week (~ 13, 35 and 110 mg/kg b.w.)</li> </ul>	Water intake was decreased in the groups at 1000 ppm. At 300 ppm, higher lymphocyte counts and lower numbers of polymorpho-nuclear cells were observed. The NOAEL was 100 ppm (part per million), equal to 13 mg/kg b.w. per day	Krötlinger, 1992
Mouse (nitroso TP)	<ul> <li>Oral application (100 mg/kg b.w.)</li> <li>Animals sacrificed after 24, 48 and 72 h post-dosing</li> <li>Micronucleus test</li> <li>TP purity 98.9 %</li> </ul>	The 48 h sacrifice group showed a statistically significant increase over controls in micronucleated polychromatics. There may be a weak effect of the tested TP on micronucleus formation under these conditions.	Herbold, 1989
Rat (nitroso TP)	<ul> <li>3 or 4 males/dose, 2 or 3 females/dose, 7 weeks old</li> <li>Oral administration at doses of 300, 1000, 1400, 1800 and 2500 mg/kg b.w. (males); and 1400 and 2500 mg/kg b.w. (females)</li> <li>Drawback: no control used</li> </ul>	$LD_{50} > 2500$ mg/kg. No mortality. Non-specified poisoning symptoms reported at all doses tested. Authors report "the poisoning symptoms were rather different from those seen in the study on a.i. IMI alone.	Nakazato, 1991
Honey bee ( <i>Apis mellifera</i> ) (4-,5-dihydroxy-, desnitro-,urea-IMI and 6CNA)	<ul> <li>3 cages of 20 bees, 3 replicates</li> <li>acute oral toxicity</li> </ul>	The LD $_{50}$ values for the TPs were each $>$ 1000 ng/bee $(>10,\!000~\mu\text{g/kg}).$	Suchail et al., 2001

Table 6: Selection of some representative IMI's TPs toxicity studies on non-target organisms. (continued)

Test organism/tested TP	Exposure/measured endpoint	Toxicity data	References
Honey bee ( <i>Apis mellifera</i> ) head membrane preparations (6CNA, urea-IMI, 4-hydroxy-, 4,5-dihydroxy-IMI)	• binding studies with IMI TPs to determine displacement of <sup>3</sup> H IMI	No effective displacing of IMI from its binding site by 6CNA and urea-IMI even at high concentrations (0.1 mM). The affinity of the other TPs for the IMI binding site decreased in the following order: olefin-IMI > 4-hydroxy-IMI >> 4,5-dihydroxy-IMI. 6CNA and urea-IMI are not biologically active via the IMI receptor in the honey bee. Results were confirmed by electrophysiological studies.	Nauen et al., 2001
Honey bee (Apis mellifera) (5-hydroxy-IMI)	<ul> <li>20 late summer worker bees of unknown age per treatment, 3 replicates</li> <li>oral administration</li> </ul>	48 h LD <sub>50</sub> : 153.5 ng/bee (125.9 - 196.9)	Decourtye et al., 2003
Honey Bee ( <i>Apis mellifera</i> ) (5-hydroxy-IMI)	<ul> <li>newly emerged worker bees, 60-163 bees per treatment</li> <li>nominal concentrations of 0, 7.5, 15, 30, 60, 120, and 140 µg/kg sucrose solution</li> <li>chronic mortality(11 d exposure)</li> </ul>	NOAEC (mortality): 120 μg/kg LOAEC (mortality); 240 μg/kg	Decourtye et al., 2003
Honey Bee ( <i>Apis mellifera</i> ) (5-hydroxy-IMI, desnitro-IMI, olefin-IMI and urea-IMI)	<ul> <li>3 cages of 30 bees per experiment, 3 replicates</li> <li>10-day chronic mortality study</li> <li>tested concentrations of 0, 0.1, 1, and 10 ug L<sup>-1</sup> (0.010, 0.1 and 1 ng/bee/day)</li> </ul>	All TPs caused mortality within 72 h after the onset of intoxication (trembling, tumbling, coordination problems). 50% mortality was reached by day 8 for all tested TPs. Except 0.1 ug L <sup>-1</sup> 5-hydroxy-IMI (reached 40% mortality by end of study). All TPs had similar timing of mortality. 5-hydroxy-IMI showed evidence of dose-response.	Suchail <i>et al.</i> , 2001

Table 6: Selection of some representative IMI's TPs toxicity studies on non-target organisms. (continued)

Test organism/tested TP	Exposure/measured endpoint	Toxicity data	References
AQUATIC INVERTEBRATES  Amphipod crustacean (Hyalella azteca) (desnitro-IMI)	<ul> <li>14-21 d old, 10 organisms per concentration, 2 replicates</li> <li>96 h static acute toxicity of</li> <li>measured concentrations of 0, 5.6, 11.0, 22.1, 43.8 and 86.8 mg L<sup>-1</sup></li> </ul>	96 h LC <sub>50</sub> : 51.8 mg a.i L <sup>-1</sup> .  96 h EC <sub>50</sub> (immobilisation): 29.0 mg a.i. L <sup>-1</sup> .  96 h NOAEC (mortality): 22.1 mg a.i. L <sup>-1</sup>	Rooney and Bowers, 1996
Amphipod crustacean ( <i>Hyalella azteca</i> ) (urea-IMI)	<ul> <li>7-21 d old, 10 organisms per concentration, 2 replicates</li> <li>96 h static acute toxicity of</li> <li>nominal (measured) concentrations of 0, 6.2 (5.81), 12.5 (11.80), 25 (23.46), 50 (46.80), and 100 (94.83) mg a.i. L<sup>-1</sup>.</li> </ul>	96 h LC <sub>50</sub> : > 94.83 mg a.i L <sup>-1</sup> , 96 h EC <sub>50</sub> (immobilisation): > 94.83 mg a.i L <sup>-1</sup> 96 h NOAEC: 94.83 mg a.i. L <sup>-1</sup>	Dobbs and Frank, 1996a
Midge (Chironomus tentans) (desnitro-IMI)	<ul> <li>2 replicates, 10 chironomids per replicate</li> <li>96-hour static acute toxicity</li> <li>nominal (measured) concentrations of 0, 0.1, (0.12), 1.0 (0.87), 10.0 (8.19) and 100 (82.8) mg a.i L<sup>-1</sup></li> </ul>	96 h LC <sub>50</sub> : >82.8 mg a.i L <sup>-1</sup> 96 h EC <sub>50</sub> (sub-lethal effects): 17.0 mg a.i.L <sup>-1</sup> 96 h NOAEC (mortality and sub-lethal effects): 8.19 mg a.i. L <sup>-1</sup> . Sub-lethal effects included spotted coloration and erratic behaviour.	Bowers, 1996
Midge (Chironomus tentans) (urea-IMI, 6CNA)	<ul> <li>16 d old, 10 chironomids per concentration, 2 replicates</li> <li>96 h static acute toxicity</li> <li>urea-IMI and 6NCA at concentrations of 0, 0.1 (0.10), 1 (1.0), 10 (10.04) and 100 (99.80) mg a.i. L<sup>-1</sup></li> </ul>	Urea-IMI: 96 h LC <sub>50</sub> : > 99.80 mg a.i. $L^{-1}$ , 96 h EC <sub>50</sub> (sub-lethal effects): >99.80 mg a.i $L^{-1}$ , 96 h NOAEC: 99.80 mg a.i. $L^{-1}$ 6CNA: 96 h LC <sub>50</sub> : > 1 mg a.i. $L^{-1}$ NOAEC=1 mg a.i. $L^{-1}$	Dobbs and Frank, 1996b Bower and Lam, 1988

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Table 6. Selection of some representative IMI's TPs toxicity studies on non-target organisms. (continued)

Test organism/tested TP	Exposure/measured endpoint	Toxicity data	References
CELL MODELS	• doses: -/+ S9: 0.1, 0.3, and 1	Cytotoxicity of nitroso TP at the mid and high dose	
Hamster V79 Cells (nitroso TP)	mg mL <sup>-1</sup> cultures harvested at 7 (high dose only), 18 and 28 h (high dose only) after start of the exposure  chromosomal aberration test	indicated by a decline in mitotic index and at the high dose by a decline in plating efficiency (-S9 only). No increase in chromosome aberrations. Nitroso-IMI is not clastogenic in this system under these conditions.	Heidemann, 1989
CHO-K1 Cells (nitroso TP)	<ul> <li>doses, -/+ S9: 0, 0.25, 0.5 and 1 mg mL<sup>-1</sup></li> <li>exposure time: -S9, 24 and 48 h +S9, 4 h</li> <li>chromosome aberrations</li> </ul>	Possible slight increase in % cells with chromosome aberrations under -S9 condition.	Usami <i>et al.</i> , 1988
Primary Hepatocytes of Male Rats (nitroso IMI)	<ul> <li>0.04, 0.13, 0.44, 1.33, 4.44,</li> <li>13.33, 44.44, 133.33, 444.44,</li> <li>and 1333.33 mg mL<sup>-1</sup></li> <li>18 h exposure; 3 replicates</li> </ul>	Severe cytotoxicity observed only above 133.33 mg mL <sup>-1</sup> .	Fautz, 1989
Mouse fibroblast M10 cells (desnitro-IMI)	<ul> <li>3 day incubation</li> <li>up-regulation of α4β2 AChRs and mechanisms of up- regulation</li> </ul>	Desnitro-IMI TP gave up-regulation by eightfold and EC <sub>50s</sub> of 870 and 500 nM, respectively. Determinated IC <sub>50</sub> for densitro-IMI (8.2 nM).	Tomizawa and Casida, 2000
Mouse neuroblastoma N1E-115 cell line (desnitro-IMI)	<ul> <li>30 min (0-180 min) incubation (0.0001-1000 μM)</li> <li>ERK kinase cascade</li> <li>intracellular calcium (Ca<sup>+2</sup>) moblisation</li> </ul>	Desnitro-IMI induced ERK activation in a dose- dependent manner triggered by primary action at the $\alpha 4\beta 2$ nAChR with an involvement of intracellular $Ca^{+2}$ mobilisation.	Tomizawa and Casida, 2002

To note: Studies performed by Usami et al., 1988 and Fautz, 1989 were evaluated as "unacceptable, but possibly upgradeable" in CDPR report (2006).

## 1.5. Biological systems: in vitro and in vivo models

Adequate and reliable data regarding adverse effects of exposure to chemical agents are crucial in order to protect human health and environment. The principal source of this information involves often the use of laboratory experimental animal and associated tests. For ethical, scientific and economic reasons, over thirty years there has been present an intensive debate and research how to **reduce**, **refine** and **replace** these animal tests (**3Rs**), without compromising the high level of human health and environmental protection demanded by the European Community. For these purposes in October of 1991 was created the European Centre for Validation of Alternative Methods (ECVAM), pointing to a requirement in the Directive 86/609/EEC on protection of experimental animals used for scientific purposes. The Directive 86/609/EEC was subsequently revised and replaced by the currently active Directive 2010/63/EU which has further enforced the role of ECVAM. In addition, the implementation of the new EU chemicals Regulation REACH (EC, 2006) and the 7<sup>th</sup> Amendment of the Cosmetics Directive (EC, 2003) motivated the community for development of easier and scientifically more reliable alternative toxicity tests.

In order to understand the mode of action and interactions of pesticides in humans or animal organisms we have to rely on *in vitro* techniques which are relevant for the development of sensitive eco-toxicological endpoints. *In vitro* bioassays as simplified model systems can, to a certain extent, replace toxicity tests on whole organisms and can be used to test the toxic potential of chemicals or contaminated environmental samples. *In vitro* bioassays based on specific cellular mechanisms measure cellular toxic effects with the use of cell cultures or transgenic bacteria or yeast. Cell cultures provide the finest system for studying toxic molecular mechanism, by permitting the cells to be studied in a monitored and isolated environment, which regulates their activities. In general, the major advantage of cell line usage is the reduction of the complexity of study systems, replacement of animal tests (3Rs) and offering benefits in terms of ethical consideration. Mammalian cells belong to tissues, organs and specialised cell types (e.g. neurons) and therefore have a substantial level of internal spatial complexity or cellular activity which results in definite molecular mechanisms. Primary or stable cell cultures maintain their

intrinsic cell-tissue molecular characteristics and therefore can provide tissue-specific and potentially useful informations on pesticide exposure toxicology. Often is impossible to transfer obtained mouse/rat facts to effects on humans, so the use of human-derived cell cultures may offer additional data. However, a major problem of *in vitro* systems is that they typically cannot take in consideration species-specific differences in sensibility and make difficult to extrapolate obtained results to whole organism models. It is often mentioned in literature that the use of *in vitro* models is 'a compromise between convenience and relevance' (Brandon *et al.*, 2003, Pelkonen *et al.*, 2005; Abbas, 2010).

The understanding of toxicant uptake and responses at whole organisms, not only cell, level has also a high ecological importance and is almost impossible to completely eliminate the need of organism models in ecotoxicology studies. For a better evaluation of integrative effects on whole organisms, validated and standardised *in vivo* assays with test species from different trophic levels (bacteria, algae and invertebrates) are applied. The use of standardised and well-established test (e.g. ISO Standard tests) in laboratory experiments is desirable as it allows achieving reproducible and comparable data. These bioassays measure effects on parameters such as growth, mortality, reproduction and feeding activity (conventional endpoints), as well as effects based on more specific biochemical / molecular biomarkers.

In addition, the monitoring of health and environment risks should be integrated with chemical analyses, screening of pollutants and possible TPs production. For example, the accessibility of LC-MS techniques in modern research has revolutionised analytical methods used for the analysis of organic molecules including pharmaceutically active compounds, veterinary drugs and pesticides (Kuster *et al.*, 2006). The up-to-date LC-MS instrumentation has excellent sensitivity and allowed the development of methods that permit the simultaneous analysis of at least 50 pesticides in different samples with high efficiency (Alder *et al.*, 2006).

Throughout this research a battery of eco-toxicological assays was applied. These assays utilised test organisms and cell models from different taxonomical and organizational level providing a possible comprehensive view of the IMI toxicity by reflecting different species susceptibility to IMI (as pure compound or as commercial formulation) and to its TPs. In this study, the following assays adopting a neuronal

cell line F11 and a bacterial strain *S. typhimurium* were referred to as *in vitro* bioassays related to specific measured endpoints. The term *in vivo* bioassays was used for assays with luminescent bacteria (*Vibrio fischeri*), algae (*Desmodesmus subspicatus*) and aquatic crustacean (*Gammarus fossarum*).

#### 1.5.1. *In vitro* models applied in this research

**Neuronal cell line F11** - in the field of *in vitro* toxicology, *in vitro* techniques based on cellular neuronal line models can be used to analyse the neurotoxic effects of chemical substances such as pesticides and their mechanisms of neurotoxicity. In this way, the cellular, morphological or functional changes analysed by the use of this in vitro models can be associated with mechanisms linked to toxic effects. The objective of this in vitro test was to use immortalised neuronal F11 cell line derived from a fusion between mouse embryonic neuroblastoma N18TG-2 and rat dorsal root ganglion (DRG) neurons (Platika et al., 1985) to better understand IMI toxic mode of action (Fig. 13). In addition, as little toxicity found for IMI in different published studies, we decide to investigate if IMI could sensitise sensory neurons that innervate the skin and that are on of major players of dermal exposure route (likely to occur in occupational toxicity to pesticides). As their origin, DRG are a complex neuronal population constituted by peptidergic and non-peptidergic neurons of different size, expressing different sub-population of neurons generating different molecules. Peptidergic neurons are characterised by a strong exocytosis of many different neuromodulators and inflammatory peptides such as: neuronal growth factor (NGF), brain-derived neurotrophic factor (BDNF), calcitonin gene-related peptide (CGRP), substance P, atrial natriuretic peptide, adenosine triphosphate (ATP) and glutamate. Upon depolarising stimuli, sensory neurons are transmitting sensory environmental stimuli to the brain, and may undergo to long-lasting activation that may result in diseases characterised by hypersensitisation of peripheral fibres. Since IMI is a nAChR agonist, it is presumed that in F11 cells can operate acting on α3 subunit of nAChRs, which is highly expressed in this cell line. A detailed characterisation of F11 cell model was performed to evaluate IMI effects on cell morphology/survival, signalling pathways, and to identify additional mechanisms of IMI's action (e. g. IMI-induced oxidative stress). On the F11 cell model were tested the cytotoxic

effects of IMI (as pure compound or as commercial Confidor 200SL) and also all four IMI TPs (6CNA, desnitro-IMI, 5-hydroxy-IMI, olefin-IMI).

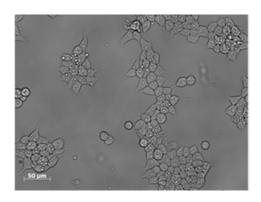


Figure 13: Image of F11 cell line in culture taken by bright field microscopy (20x).

In vitro bacterial reverse mutation (Ames) test - the bacterial reverse mutation test uses amino-acid requiring strains of S. typhimurium (Fig. 14) to detect point mutations, which involve substitution, addition or deletion of one or a few DNA base pairs. Point mutations are the cause of many human genetic diseases and there is substantial evidence that point mutations in oncogenes and tumour suppressor genes of somatic cells are involved in tumour formation in experimental animals and humans. The particular bacterial S. typhimurium strain (LT2) with a deep rough mutation (rfa) is relatively harmless and is useful in the determination of the possible genotoxic potential of several samples. This strain is used in Ames test, an ISO/DIS Standard 16240 (2005) test and is able to measure mutagenicity of chemicals that induce point or frame-shift mutations. However, it is important to note that the bacterial reverse mutation test utilises prokaryotic cells, which are different from mammalian cells in their uptake, metabolism, chromosome structure and DNA repair processes. Therefore, the in vitro test such as this one generally requires the use of an external source of metabolic activation to produce in vitro derivatives by liver detoxification enzymes. In vitro metabolic activation systems cannot mimic entirely the mammalian in vivo conditions and often does not provide direct information on the mutagenic and carcinogenic potency of tested substance in mammals. Generally, the bacterial reverse mutation test is rapid, inexpensive, and is employed as an initial screen for genotoxic and point mutation-inducing activity.

This method was included in our experimental strategies to explore mutagenic or carcinogenic potential of tested IMI, its commercial formulation and TP (6CNA).

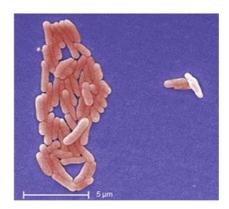


Figure 14: Coloured scanning photomicrograph under a magnification of 5000x shows numbers of clustered Gram-negative Salmonella typhimurium bacteria (AP Photo/CDC, Janice Haney Carr).

### 1.5.2. In vivo models applied in this research

The test organisms incorporated in these *in vivo* assays included species of different taxonomic group and different trophic level as microorganisms, plants and invertebrates.

Bioluminescent marine bacteria Vibrio fischeri – bacteria play an important role in the environmental fate of applied pesticides and other chemical compounds. For over 25 years the luminescent bacteria test with marine bacterium V. fischeri is used as a basic bioassay for eco-toxicological testing of chemicals, waste water and eluents or leachates from soil and sediment. V. fischeri is a Gram-negative bacterium with a unique characteristic, bioluminescence, which is controlled by a small set of genes known as the lux operon (Fig. 15). The lux operon is a 9 kb fragment that consists of genes that code for the subunits of luciferase (luxAB) and for enzymes (luxCDE) that convert organic compounds to oxidised substrates.

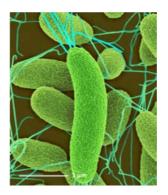


Figure 15: Coloured scanning photomicrograph under a magnification of 8000x of Vibrio fischeri bacteria a flagellated, bioluminescent, curved rod-shaped marine prokaryote (Dr. Dennis Kunkel Microscopy, Inc./Visuals Unlimited, Inc.; with slight modification).

Luminescence is a chemical reaction in which the enzyme luciferase oxidises organic compounds, such as long chain aldehyde and reduces flavin mononucleotide, so in this way releases free energy in the form of blue-green light at 490 nm. The reaction that takes place in *V. fischeri* is described schematically in Fig. 16.

The bioluminescence produced by the marine bacterium V. fischeri is the basis for ISO standard toxicity bioassay that has been used to assess the toxicity of aqueous solution of known or unknown samples. Differences in the amount of light produced (inhibition of bioluminescence) can therefore be correlated to the organism's metabolism and is proportional to the toxicity of the tested sample. The luminescence is the test criterion which is measured after a contact time of 30 min (between bacteria and tested sample) taking into account a correction factor (Cf) that is derived from the intensity changes of control samples during the exposure time. The inhibitory effect of the sample on the light emission of V. fischeri can be determined as a 30 min EC<sub>50</sub> value (in case that 50 % of inhibition is not reached is possible to determine also the EC<sub>20</sub> value - concentration that causes 20% inhibition relative to control). The V. fischeri light emission inhibition is measured according to the ISO-Guideline No. 11348-Part 1-3 (2007) as defined in a Standard Operation Procedure (SOP). Measurements can be carried out using freshly prepared bacteria (Part 1), as well as liquid-dried (used in this research; Part 2) or freeze-dried (Part 3) bacterial preparations. The ISO guideline 11348 (2007) recommends testing at least one of the three reference substances (3,5-dichlorophenol, potassium dichromate and zinc sulphate heptahydrate) parallel to each testing to ensure the evaluation of the toxicity test.

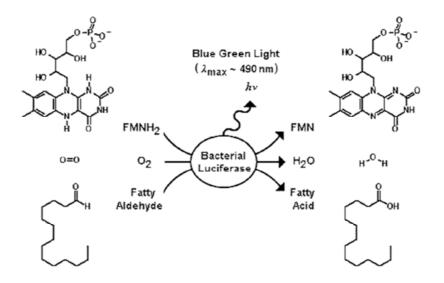


Figure 16: The net chemical equation of the bacterial luciferase catalysed reaction (http://www.photobiology.info/Lin.html).

The effect of IMI on V. fischeri was already assessed by Tišler et al. (2009) and showed a 30 min IC<sub>50</sub> of 61.9 mg L<sup>-1</sup> for IMI and of 56 mg L<sup>-1</sup> (0.028 % v/v) for Confidor 200SL, while Kungolos et al. (2009) determined an IC<sub>50</sub> of 226 mg L<sup>-1</sup> for IMI. An additional study was performed by Dell'Arciprete et al. (2009) and reported the inhibition of V. fischeri luminescence of  $32 \pm 7$  % at concentration 80 mg L<sup>-1</sup> for IMI. Data relative to TPs include a 30 min EC<sub>50</sub> of 15.1 mg L<sup>-1</sup> for 6CNA (Žabar, 2012). The luminescent bacteria was applied as a model during this research to test three more TPs of IMI (desnitro-IMI, olefin-IMI and 5-hydroxy-IMI) which at the moment were not tested for their possible toxic effects towards aquatic organism.

Freshwater green unicellular algae *Desmodesmus subspicatus* - among the potential test organisms, unicellular green algae are commonly used for toxicity tests. It is very important to expand the knowledge about harmful effects of toxic substances in aquatic environment on growth of single-cell algae, since they are an important component of freshwater systems, being the primary producer of organics and oxygen in the trophic food chain (Berden-Zrimec *et al.*, 2008). Any adverse impact on algae is likely to affect organisms at higher trophic levels and may have important consequences for the health status of the whole aquatic ecosystem (Franklin *et al.*, 2000). One of the useful ISO/DIS Standard 8692 (2004) test is the freshwater algal growth inhibition test with single cell algae. This test determines the

concentration of contaminant which reduces the growth of microalgae by the means of chlorophyll fluorescence measurements. Test organisms used in this test are planktonic freshwater algae, *D. subspicatus* (Chodat) Hegewald *et* Schmidt (formerly *Scenesdesmus subspicatus* (Fig. 17). This algae species are green algae belonging to the order of *Chlorococcales* (*Chlorophyta*, *Chlorophyceae*) and are usually unicellular in culture. The recommended strains are available in uni-algal, non-axenic culture from SAG (Collection of Algal Cultures, Germany), ATCC (American Type Culture Collection, USA) or CCAP (Culture Centre of Algae and Protozoa) collections.

Freshwater algae have been used in several toxicity studies (Riva *et al.*, 2002; Bengston *et al.*, 2005). Primarily, the measurement of algae growth inhibition of tested samples is used to evaluate the impact of chemicals in aquatic environments. In this respect, unicellular algae *D. subspicatus* were also included in our research to investigate the sub-chronic toxicity of IMI, Confidor 200SL and 6CNA

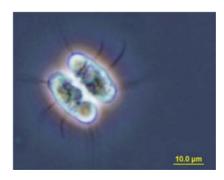


Figure 17: Image of Desmodesmus subspicatus algae taken by bright field microscopy (20x).

Freshwater stream scud Gammarus fossarum — Invertebrates having simple molecular machinery are often used as good model organisms to explore basic molecular mechanisms occurring in altered physiological states. Particularly, crustaceans are suitable for biomonitoring in aquatic toxicity tests due to their prolific breeding, high abundance in nature and sensitivity to anthropogenic toxic compounds in water bodies which they inhabit (Fernandez-Alba *et al.*, 2002). In addition, they are widely distributed, easy to collect and less migratory than fishes.

Amphipoda is one of the most diverse and widespread crustacean's orders. They play a key role in the detritus breakdown process and are commonly used in freshwater risk assessment (Rinderhagen *et al.*, 2000; MacNeil *et al.*, 2000).

Moreover, the use of these species is logistically interesting because they can be sampled throughout the year and easily identified, manipulated and maintained/used in the laboratory for ecotoxicological bioassays.

Amphipods (i. e. gammarids) are flattened from side to side and lack of the carapace covering the thorax. They have seven thoracic and six abdominal plated segments, which support and provide shelter for the gills and other soft parts of the animal. The head carries two pairs of antennae, the stalk-less eyes, and the mouthparts (Schram 1989, Barnard and Karaman, 1991). The name 'amphipod' comes, from having seven pairs of walking legs of which the first four reach backwards and the fifth to seventh reach forwards. Typical body length for an adult amphipod is from a few millimetres to a few centimetres, depending on the species. The life cycle of gammaridean amphipods typically consists of five different periods. Since the sexes of amphipods are separate, eggs develop in the brood pouch of the females. Unlike most crustaceans, the amphipods lack a free-living larval stage and juveniles look very much like the adults. Once the juveniles are big enough to start living on their own, they are released. The growth of amphipods is connected to the change of the rigid exoskeleton. After a varying time period and a certain number of moults, juveniles achieve characteristics typical to their sex.

G. fossarum Koch. 1835 is one of most abundant species in freshwater ecosystems in Slovenia (Fig. 18). They are translucent slightly brown benthic animals with alternated periods of swimming and longer periods of burrowing or crawling. Gammarids are often found in high densities in headstreams where they are an important reserve of food for vertebrates, i.e., fish, bird and amphibian species (Friberg et al., 1994; MacNeil et al., 2000). They are an important component of freshwater ecosystems since they play a key role in leaf litter breakdown process and are important for material transfer in the food web (MacNeil et al., 2000; Maltby et al., 2002).



Figure 18: Image of stream scud Gammarus fossarum (Peter Balej; http://www.biolib.cz).

All these previously described facts explain why amphipods are often used in eco-toxicological studies and why this organism models was included in this study. Furthermore, crustaceans as well as insects belong to arthropods and due to this crustacean-insect relationship they could present a potentially suitable non-target group for IMI testing (Strausfeld, 1998). In general, aquatic biota is extremely variable going from uni- to multi-cellular organisms. Thus for aquatic eco-toxicity testing is essential to use a battery of bioassays representing differences in sensitivity of different test organisms from various trophic levels to cover this variability (Fig. 19).

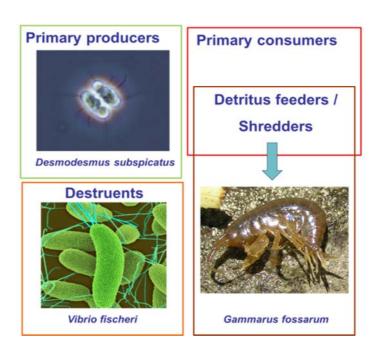


Figure 19: The use of a battery of bioassays involving different model organisms from different trophic levels as an essential tool for predicting adverse effects to the aquatic ecosystem. (Kienle et al., 2011; with slight modifications)

## 1.6. Biological effects and responses: multi-biomarker approach

The use of molecular biology and biochemical methods to understand toxicity mechanisms provides ecotoxicologist useful tools which permit them to investigate possible biological effects or responses due to the impact of all pollutants released from both natural and anthropogenic sources in aquatic ecosytems (Carajaville et al., 2000; Sarkar et al., 2006). To analyse such toxicological, biochemical and physiological process is useful to measure specific biomarkers. Biomarkers are defined as quantitative measures of changes in the biological system that respond to either (or both) exposure to, and/or doses of xenobiotic substances that lead to biological effects. They show the interaction between biological system and potentially harmful chemical, biological and physical action (WHO, 1993). According to the NRC (1987) and WHO (1993), biomarkers can be subdivided into three classes: biomarkers of exposure, of effect and of susceptibility. The use of term biomarker is often restricted to cellular, biochemical, molecular, or physiological changes that are measured in cells, body fluids, tissues, or organs within an organism and are indicative of xenobiotic exposure and/or effect changes that occur at the organism, population and more complex levels. One of the functions of biomarkers is to provide early warning signals of biological effects, and that it is generally believed that sub-organismic (molecular, biochemical and physiological) responses tend to precede those that occur at organism or higher levels (Fig. 20).

Different biomarkers were used in order to test potential adverse effects of IMI and its related compounds on model systems described in sections 1.5.1 and 1.5.2 of this manuscript. The selected biomarkers included in this study are described as follows:

#### IN VITRO MODELS

## • Bacteria S. typhimurium

- mutagenicity potential

### • Neuronal F11 cell line

- survival/cytotoxic effects
- structural/morphological parameters
- molecular/sub-cellular parameters

### IN VIVO MODELS

## • Marine bacteria V. fischeri

- bacterial luminescence inhibition

## • Unicellular green algae D. subspicatus

- rate of algae growth inhibition

## • Amphipod G. fossarum

- mortality rate
- oxidative stress parameters (e.g. enzyme activity)
- multixenobiotic resistance (MXR) mechanism
- behavioural endpoints

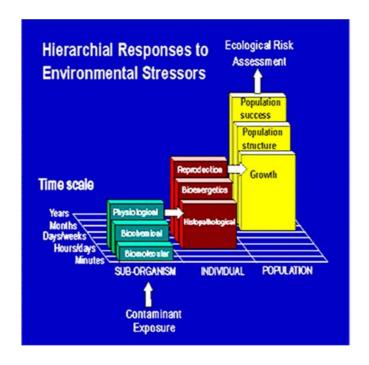


Figure 20: Hierarchical responses to environmental stressors (Adams, 1990).

### Biochemical biomarkers (stress enzymes) as indicator of stressful effects

Different classes of **pesticides** may be related to enhanced production of reactive oxygen species (ROS) or reactive nitrogen species (RNS) which could contribute to the toxicity of these compounds (Banerjee *et al.*, 2001). Basic cellular metabolism in aerobic organisms involves the production of oxygen free radicals and non-radical ROS (Halliwell and Guttridge, 2007). The imbalance between the generation and the neutralisation of ROS by antioxidant mechanisms within an organism generates the **oxidative stress** (Livingston, 2001; Valvadinis *et al.*, 2006) (Fig. 21). Pesticides are known to induce oxidative stress by:

- induction of ROS production as by-products of detoxifying mechanisms,
- alteration of normal mitochondrial respiration processes or
- by their own reduction/oxidation properties.

Pesticide induced oxidative stress can than provoke oxidant-mediated effects (such as increased activities of antioxidant enzymes) and oxidant-mediated toxicities (such as oxidation of lipids, mutations or cell death) (Zama *et al.*, 2007; Tebourbi *et al.*, 2011). Enhanced ROS are also implicated in pesticide-induced neurotoxicity, based

on their role in biochemical changes that give rise to neuronal cell damage and death (Jia and Misra, 2007). Moreover, oxidative stress induced by pesticides could be one of the sources for neurodegenerative disorders such as Alzheimer's or Parkinson's disease (Bogdanov *et al.*, 2001; Huang *et al.*, 2004).

The primary line of defence is offered by general antioxidant systems, which have been demonstrated to scavenge ROS and free radicals. There are essentially three systems which include: (i) water soluble reductants (e.g. glutathione, ascorbate), (ii) lipophilic reductants (e.g. α-tocopherol, or β-carotene) and, (iii) enzymes (e.g. catalase, peroxidase or superoxide dismutase). From the point of view of biomarkers, a condition of oxidative stress can be shown through evaluation of specific biomarkers of oxidative damage to macromolecules such as lipids, proteins and nucleic acids (Woo *et al.*, 2006). Generation of ROS and their action affects antioxidant enzymatic activities, lipid peroxidation and mitochondrial function. In addition, oxidative stress by environmental pollutants activates several signalling molecular pathways such as: MAPK pathways or extracellular signal-regulated kinase (ERK) which is involved in cellular apoptosis – cell death (Ki *et al.*, 2012).

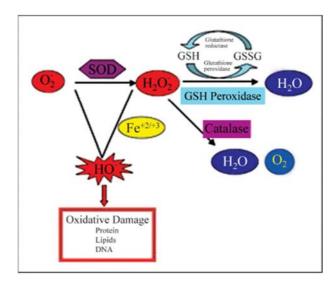


Figure 21: An unbalance between the production of pro-oxidants and antioxidants in cell might lead to strengthened production of ROS which could lead to serious cellular damage (Wakamatsu et al., 2008).

In fact, even if pesticide toxicity is often restricted to their mode of action, as in case of IMI to insect neuronal transmitter receptors, it is important to expand our knowledge regarding other provoked mechanisms (e.g. ROS production) which could contribute to the toxicity of this insecticide. In this research, in order to have a better understanding of the toxic action of IMI and its TPs the involvement of induced ROS production was investigated by measurement of oxidative stress biomarkers such as antioxidant enzyme catalase (CAT), detoxifying enzyme glutathione-S-transferase (GST) and the levels of lipid peroxidation (LP) damage in non-target aquatic organism (G. fossarum) (Malev et al., 2012) and in neuronal F11 cell line.

CAT is one of the antioxidant enzymes and an essential defence against the potential toxicity of superoxide anions like hydroxyl free radical. CAT extensively decomposes hydrogen peroxide formed during oxidative stress (Winston and Di Giulio, 1991). A change in CAT activity is often an indicator of a cellular damage after exposure to chemicals, and therefore it is considered as an early environmental stress biomarker in different cases. Most studies about CAT activity focus on organisms after exposure to pesticides, hydrocarbon (Brown *et al.*, 2004) and heavy metals (Laszczyca *et al.*, 2004).

GST belongs to the class of phase II detoxifying enzymes that catalyses the conjugation of glutathione with xenobiotics and cytotoxic aldehydes produced during lipid peroxidation (Banerjee et al., 2001; Bebe and Panemangalore, 2005). GSTs enzymes catalyse the reaction of xenobiotic compounds with the -SH group of reduced glutathione (GSH), thereby neutralising their electrophilic sites and rendering the products more water-soluble (Edwards *et al.*, 2000). In addition to catalysing GSH conjugation, GSTs also exhibit glutathione peroxidase (GSH-P<sub>X</sub>) activity, which suggests a role in protection against oxidative stress. Consequently, GST activity it is a potential biomarker of a detoxification process induced by very large number of pollutants capable of generating oxidative stress. For example, in insects, GST has been induced by exposure to a number of chemicals, including pesticides, such as lindane (Lagadic *et al.*, 1993), paraquat (Parkes *et al.*, 1993) and oxadiazolone (Hunaiti *et al.*, 1995).

LP is considered an oxidation-related biomarker and it represents the negative effects caused on the lipid membranes by ROS. LP occurs when the natural antioxidant defences are overcome and fatty acid hydro-peroxides formed with the consequence of membrane and membrane-bound enzymes destabilisation (Porter et al., 1995; Linden et al., 2008). One of the consequences of oxidative stress is the peroxidation of lipids. Lipid peroxides, derived from polyunsaturated fatty acids, are unstable and decompose to form a complex series of compounds. These include different reactive carbonyl compounds of which the most abundant is called malondialdehyde (MDA) and is used as a marker of LP induced by xenobiotic stressors (Del Rio et al., 2005). The most widely used assay for lipid peroxidation is the thiobarbituric acid reactive substances (TBARS) method (Draper et al., 1993). MDA as a secondary lipid peroxidation product in biological samples reacts with thiobarbituric acid (TBA) under strong acidic condition and heating. This reaction is actually the formation of a pink-colour product (TBA-MDA) which can consequently be measured by the colorimetric method (Lykkesfeldt, 2007). The detailed reaction is shown in Figure 22.

Figure 22: Schematic illustration of MDA-TBA reaction and formation of a colour product (Janero, 1990).

AChE is often used as a biochemical biomarker that indicates stress effects due to pesticide exposure. AChE plays a key role in regulation of cholinergic nervous transmission. It is responsible for the hydrolytic degradation of ACh, which is the primary neurotransmitter in sensory and neuromuscular systems in most organisms. The active site of AChE contains a serine hydroxyl group that binds to anticholinesterase insecticides (with structure similar to Ach), which inhibit the AChE, generating a phosphorylated unreactive enzyme and causing serious damage to neuronal structures. AChE inhibition leads to overstimulation of the nervous system, which results in neurotoxicity and cellular death. Monitoring studies based

on the inhibition effects are majorly related to the presence of organophosphorus insecticides and carbamates (Fulton and Key, 2001). However, evidence of the modulation of esterase activities by other organic chemicals (e.g. surfactants), metals and indirect alteration by neonicotinoids (e.g. IMI) has been described (Jifa *et al.*, 2005; Dondero *et al.*, 2010). In our study we tested possible indirect inhibitory effects on freshwater amphipods exposed to IMI (Malev *et al.*, 2012). This measurement was also performed on gills of neonicotinoid exposed mussels and showed an interesting outcome and change of AChE activity (Dondero *et al.*, 2010). In this study was observed a significant inhibition at the lowest and at the intermediate tested IMI concentration.

### Morphological, sub-cellular and molecular changes as indicator of stressful effects

In the test that involved IMI testing on neuronal F11 cell model was evaluated not only the final endpoint of cell survival (cytotoxicity), but also subcellular and molecular factors which can be evoked upon cell exposure to stressful stimuli. Immunofluorescence (IF) appears as a powerful in vitro technique to assess susceptibility of different cell to xenobiotics. Using the IF method is possible for example to detect endogenous proteins induced by different contaminants (Marsili et al., 2008) as well as alterations of cellular ultrastructure (Liu et al., 1996; Segner and Braunbeck, 1998). In our case was possible to detect the changes in the cell cytoskeleton (i.e. actin, tubulin, intermediate filaments) and other ultra-structural alterations. One specifically studied example was mitochondria that are not only an important source of cellular energy but they also preserve intracellular  $\text{Ca}^{2+}$  levels and regulate calcium dependent cytotoxicity (Foster et al., 2006). Various alterations in the number, size shape and mitochondria membrane potential can occur in different stressful conditions and can be detected not only ultra-microscopy analysis with (transmission electron microscopy; TEM), but also with the usage of specific molecular dyes (e.g. JC-1) detectable with optical imaging as performed in this research.

Standard high priority targets other than the above mentioned cytoskeleton proteins were also P-glycoprotein. Organisms protect cellular targets via efflux transporters, such as P-glycoprotein (P-gp) and multidrug resistance protein (MRP)

present in cell membranes (Cole and Deeley, 1998, Endicott, 1989). P-gp are ATP dependent transport protein which limit the permeability of xenobiotics into the cell, and recognise or eliminate potentially cyto- and genotoxic metabolites produced during metabolism/detoxification of these compounds (Deeley *et al.*, 2006). This system has been termed **multixenobiotic resistance** (**MXR**) **mechanism** in aquatic organisms and may be induced or inhibited by a variety of chemicals (Bard, 2000; Kurelec, 1992). MXR mechanism, mediated by the activity of membrane efflux transporters, represents a first line of defence in aquatic organisms exposed to environmental contaminants.

Immunochemical analysis with mammalian monoclonal antibody to cross react with P-gp counterparts has been used to detect the presence of P-gp in aquatic organisms (Sauerborn Klobučar, 2010; Bard *et al.* 2002; Cooper *et al.*, 1999). Expression of MXR proteins is often reflective of xenobiotic levels to which an organism is exposed (resistance to environmental toxins is likely conferred by the overexpression of these proteins).

Another common method to detect MXR activity is through **dye exclusion assays**. This assay applies specific fluorescent dyes (rhodamine B, rhodamine 123 and calcein AM) to test the ability of cells to extrude a known MXR substrate. Under normal conditions with fully functional transporters the applied dye is readily expelled from the cells giving low fluorescence values. On contrary, when MXR efflux function is compromised by the use of an MXR inhibitor the fluorescent dye accumulates within the cell (Fig. 23). Dye exclusion assays are relevant for evaluating chemicals, such as environmental contaminants, for their ability to inhibit MXR. This methodology was applied also in this research to investigate the MXR mechanism in aquatic amphipods stressed by neonicotinoid insecticide IMI and its TP 6CNA. In addition was test also a possible synergistic effect of heavy metal (copper) and IMI or 6CNA.

Copper-based compounds are intentionally introduced into the environment as killing agents for fungi and algae. One of the copper based compounds mostly present is copper sulphate (CuSO<sub>4</sub>) which can be found in most running waters. CuSO<sub>4</sub> is in large amounts used as fungicide in agriculture, especially in viticulture as an agent to protect grapes against bacteria and can appear concurrently with IMI in the environment. Copper (as copper oxychloride or copper hydroxide or copper

sulphate tribasic) as foliar fungicide with protective action and IMI as systemic insecticide with protective and curative action are all used in potatoes, tomatoes and numerous vegetable (Tomlin, 2000). It is also used as an algaecide and herbicide in irrigation and municipal wastewater treatment. Trade names of its commercial formulations are Agritox<sup>®</sup>, Basicap<sup>®</sup>, and Bluestone<sup>®</sup>. Copper may be released and become free to catalyse the formation of highly reactive hydroxyl radicals and initiate oxidative damage (Gaetke and Chow 2003; Bopp *et al.*, 2008). CuSO<sub>4</sub> is water soluble and it doesn't have additional solvent or metabolite that can be toxic to animals. It presents seasonal applications, so very high instant concentrations can be locally observed in freshwaters (de Oliveira-Filho *et al.*, 2004). Graves *et al.* (2004) found copper concentrations up to 77.4 mg L<sup>-1</sup>, which were attributed to the wide use of several forms of Cu as fungicides for tomatoes and direct application of copper sulphate as an algaecide or herbicide in golf courses, lakes and ponds.

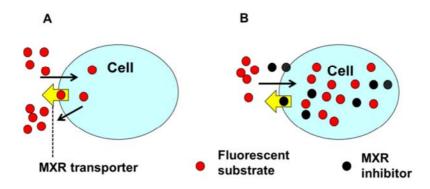


Figure 23: (A) Cell with normal MXR function where transporters recognise the fluorescent dye and expel it from cell which has a low fluorescence value. (B) Cell with disrupted MXR function exposed to an MXR inhibitor or chemosensitizer accumulate the fluorescent dye leading to higher cell fluorescence. (Epel et al., 2008 with slight modifications).

MXR inhibition, can occur when there is an abundance of substrates present, leading to saturation of the efflux system. In this case the pumps become overwhelmed and chemicals are accumulated within the cell (Epel *et al.*, 2008). Another way is through chemicals known as "chemosensitizers" (inhibitors, modulators) which inhibit MXR function and thus represent specific environmental contaminants. A wide range of pesticides have been shown to act as MXR inhibitors. Presence of chemosensitizers in the environment may increase the absorption of

xenobiotics and consequently their cellular toxic influence in organisms causing harmful physiological/pathological damage (Žaja *et al.*, 2006).

Cells possess adaptive mechanisms to counteract environmental stresses and stimuli that can continuously occur. Usually, these systems may be divided in different categories which include some of the previously mentioned mechanisms such as nucleophilic trapping processes (catalysed by GSTs) or efflux transporters that export toxic compounds or metabolites (MXR). In general, induction of these protective responses requires different components on of which is nuclear factor (erythroid-derived 2)-like 2 (Nrf2), a transcription factor responsible for the induction of downstream genes involved in cell protection from deleterious effects of oxidative stress. Nrf2 target genes encode phase II detoxifying enzymes, antioxidant proteins, ABC transporters, scavenge receptors, chaperone proteins, and so forth (Kobayashi and Yamamoto, 2005). Under normal conditions Nrf2 is sequestered in cytoplasm by a cytoskeletal protein, Keap1. Situation is modified under stressful conditions, when electrophiles and/or ROSs cause dissociation of Nrf2 from Keap1. As a consequence, Nrf2 is translocated to the nucleus where leads to the activation of cytoprotective genes involved in electrophile conjugation, excretion of xenobiotics, ROS scavenging and stabilisation of cellular redox potential. The activation of the Nrf2 – antioxidant response element signalling pathway is a major mechanism in the cellular defence against oxidative or electrophilic stress. Interestingly, mutations in Nrf2 gene, and lack of Nrf2 function in rodents and in humans, has been associated to many chronic diseases (Hayes and McMahon, 2009). Role of Nrf2 transcription factor in cellular response to oxidative stress after IMI exposure was used in this research as an 'indirect' marker of IMI-induce oxidative stress.

Molecular changes after the influence of various stressors can induce multiple signalling pathways represented by the one stimulated by **mitogen-activated protein kinases (MAPK)** - ERK, JNK and p38 family members. MAPK are members of a family of serine/threonine protein kinases activated by dual phosphorylation at threonine and tyrosine positions (188 and 190, respectively). These MAPKs are implicated in cellular processes such as gene regulation, metabolic reactions, cell proliferation/differentiation, cell mobility and survival or death (Roux and Blenis,

2004). A thoroughly and comprehensive review the functions of MAPKs is presented by Gerits *et al.* (2007). In general, these kinases are important to elicit signal transduction and nuclear factor activation responsible for controlling the expression of individual genes. To test mechanisms of IMI acute toxicity was evaluated the induction of downstream pathways like p38 MAPK or ERK in F11 cell model.

### Mutagenicity parameters as indicator of stressful effects

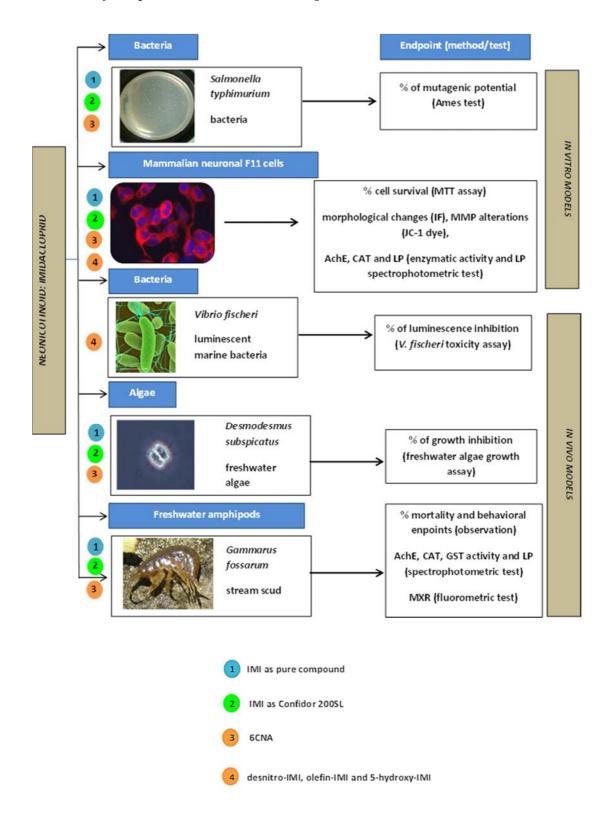
Many environmental chemical pollutants in the aquatic ecosystems have the capability to harm and change genetic material of natural biota. At molecular levels DNA is one of principal site of xenobiotics action (Frenzilli et al., 1999), so the assessment of DNA damage is of primary concern when determining the pollutionrelated stress in living organisms (Klobučar et al., 2003; Malev et al., 2010; Klobučar et al., 2012). Maintenance of DNA integrity is extremely important for all living organisms, which have developed useful and different mechanisms for protection of genetic material. Genotoxic substances can cause effects such as carcinogenesis, teratogenesis, embryotoxicity as well as a suite of health disorders referred to as genotoxic disease syndrome which is manifested as impaired enzyme function, enhanced protein turnover, impairment immune responses, production of initiators of cytotoxic injuries, inhibition of growth, decreased fecundity and faster ageing that can have adverse effects on stability of ecosystems (Kurelec, 1992). Mutagens are the most dangerous compounds present in ecosystems, because at low concentrations their effect can have consequences at higher levels of organisation that extend for several generations (Bolognesi and Degan, 2001). Furthermore, most chemicals exert their effects via both genotoxic and metabolically toxic mechanisms operating simultaneously. Therefore, there is an increasing need for sensitive assays to monitor the genotoxicity and mutagenicity of the environment. One of these standardised assays called Ames test was also applied in this study to assess the mutagenic potential of the tested neonicotinoid compounds.

## Behavioural endpoints

Measurements in this category include assessments of behavioural responses to exposure such as feeding, mating, parental care, and other behavioural modification. Behaviour is considered as a useful tool in ecotoxicology since is one of the early warning indicators of toxicant stress (Pestana *et al.*, 2009). During experimental exposure analysed individual biochemical biomarkers should be linked to behavioural responses whenever this is possible (Xuereb *et al.*, 2009; Hellou, 2011). A potential advantage of tested behavioural endpoints is in the detection of novel chemicals or chemicals with neurological effects (e.g. IMI), which may provoke more pronounced behavioural changes before effects are detectable at other levels of biological organisation.

# 1.7. Experimental research design

Figure 24: Schematic diagram of the experimental design with tested models and measured biomarkers after exposure to IMI and its TPs during this research.



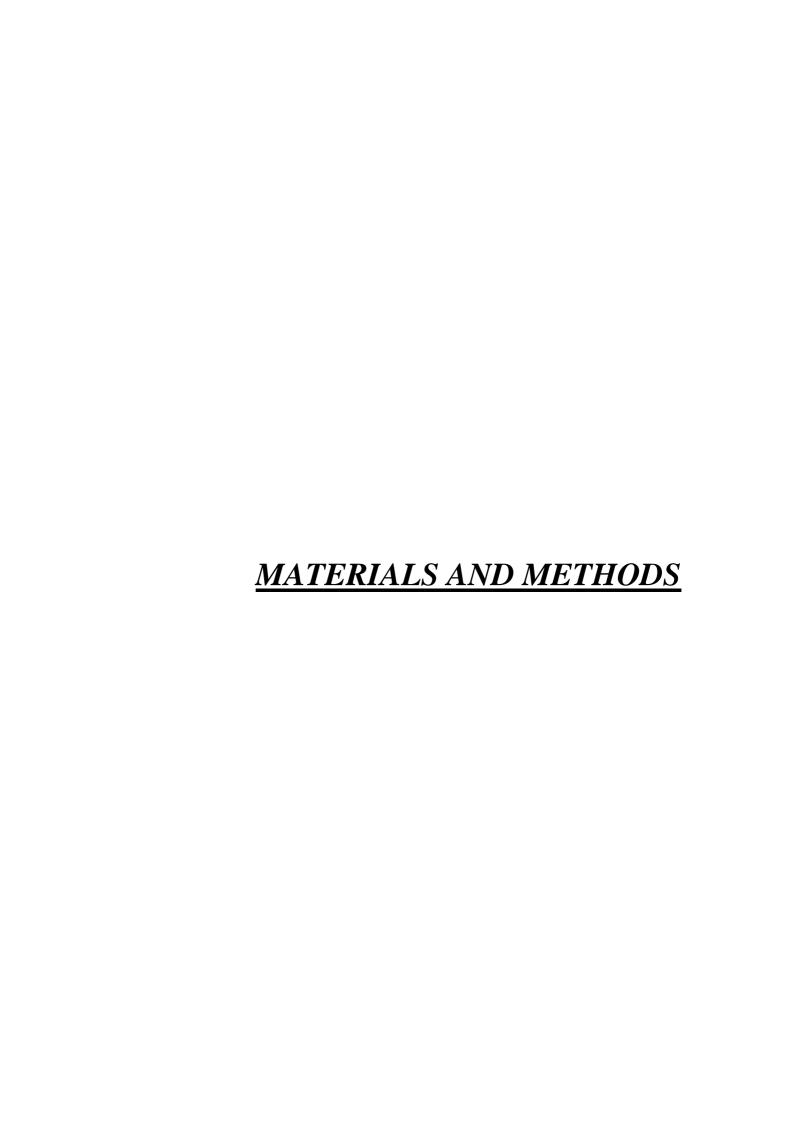
## 1.8. Experimental research objectives

This research project intends to provide a full picture of IMI mode of action and its possible harmful effects for the environment and non-target groups of organisms (biota or human). During this study was proposed the application of a battery of eco-toxicological assays which included bacteria, algae, invertebrates and mammalian cell culture to test the toxicity potential of IMI and it TPs. In addition, these assays were integrated with the measurement of several biomarkers such as biochemical, molecular and physiological responses.

### Main objectives of the research project were:

- testing of acute and sub-chronic IMI effects on biochemical, survival and growth parameters in non-target organisms: amphipod *G. fossarum*, unicellular green algae *D. subspicatus* and bacteria *V. fischeri* (a battery of test organisms from different taxonomic and trophic level was selected);
- comparison of IMI effects with effects of its commercial formulation Confidor 200SL and its TPs (6CNA, desnitro-IMI, olefin-IMI and 5-hydroxy-IMI);
- better understanding of the involvement and impact of IMI-induced ROS production and consequent oxidative stress in amphipod *G. fossarum* and in neuronal F11 cells:
- screening of the IMI, Confidor 200SL and 6CNA mutagenic potential;
- investigation of toxic effects of IMI, Confidor 200SL and IMI's four TPs on survival of neuronal F11 cells and its molecular responses discovering the mechanisms with which cells protect and activate stress pathways;
- determination of the utility of *in vitro* cell model systems represented in this study by mammalian neuroblastoma F11 cell line, to obtain information about the specific toxic mode of action of IMI.

The use of a multi-test approach was applied, as it affords a complete screening of the eco-toxicological values of IMI (as pure compound and commercial mixture) and its transformation products.



### 2. MATERIALS AND METHODS

## 2.1. Chemicals

*Tested compounds:* IMI was purchased as the Pestanal® grade chemical (99.8 % purity; Sigma-Aldrich, UK), and as a commercial formulation known as Confidor 200SL (200 g L<sup>-1</sup> of IMI a. i. form Bayer Crop Science Slovenia, Ljubljana, Slovenia). 6CNA was obtained as pure compound (97 %) from Fluka (Sigma-Aldrich, Switzerland). Desnitro-IMI, 5-hydroxy-IMI and olefin-IMI were a courtesy from Bayer Crop Science Germany. Molecular weight in g mol<sup>-1</sup> of all tested compounds are as follows: IMI (255.66), 6CNA (157.55), desnitro-IMI (247.13), olefin.IMI (271.66) and 5-hydroxy-IMI (253.65).

Chemicals for the HPLC-DAD and LC-MS analyses: acetonitrile CHROMASOLV® and methanol CHROMASOLV® for HPLC grade (Sigma Aldrich), double deionised water (< 18 M $\Omega$  cm) prepared through the NANOpure water system (Barnstead, USA) and acetic acid glacial 100 % p.a. from Merck.

Chemicals for total organic carbon (TOC) and total nitrogen (TN) analyses: potassium hydrogen phthalate was purchased from Alfa Aesar GmbH, ammonium sulphate from Fluka, hydrochloric acid 37 % puriss. p.a. from Sigma Aldrich Company Ltd.

Chemicals for biochemical biomarkers measurement: all obtained from Sigma Aldrich: dibasic and monobasic potassium and sodium phosphate, 5,5-dithiobis-2-nitrobenzoic acid (DTNB), acetylthiocholine iodide, 1-chloro-2,4-dinitrobenzene (CDNB), *L*-glutathione (reduced form), hydrogen peroxide (30%), bovine serum albumin (BSA), Bradford reagent, trichloroacetic acid (TCA), thiobarbituric acid (TBA), malondialdehyde tetraethylacetal (1,1,3,3-tetraethoxypropane) (MDA standard), butylhydroxytoluene (BTH), 96 % ethanol, 1-butanol, hydrochloric acid (37 %).

Chemical for MXR dye exclusion assay: all from Sigma Aldrich: rhodamine B, verapamil hydrochloride, cyclosporine A and phosphate buffered saline without  $Ca^{2+}$  and  $Mg^{2+}$  (PBS).

Chemicals for algal growth inhibition test: standard growth and nutrient medium of the ISO Standard 8692 (2004) prepared according to the protocol using

double deionised water and analytical grade chemicals, potassium dichromate puriss. p.a. (Sigma Aldrich).

Chemicals for Ames test: growth and exposure medium according to the standard protocol ISO 16240 (2005). Additional chemicals all from Sigma Aldrich: 2-aminofluorene (2-AF; Sigma Aldrich), methyl methanesulphonate (MMS 99 %; Sigma Aldrich), N-methylpirrolidone, ampcillin sodium salt, citric acid monohydrate, dimethyl sulfoxide (DMSO),  $\beta$ -nicotinamide adenine dinucleotide phosphate sodium salt hydrate (NADP) and L-histidine.

Chemicals for bacterial luminescence inhibition test: sodium hydroxide p.a. from AppliChem, sodium chloride from Carlo Erba Reagenti, hydrochloric acid 37 % puriss. p.a. from Sigma Aldrich Company Ltd..

Chemicals used with neuronal F11 cell model: supplied from PAA-Austria: Dulbecco's modified Eagle's medium (DMEM) - high glucose (4.5 g  $L^{-1}$ ) with L-glutamine (with and without Phenol red indicator), foetal bovine serum (FBS) heat inactivated, penicillin/streptomycin 100 x. Purchased from Sigma Aldrich: trypsin-EDTA solution 10 x , poly-L-lysine hydrochloride (PLL), thiazolyl blue tetrazolium bromide (MTT salt), 4',6-diamidino-2-phenylindole dihydrochloride ( $\geq$  98 %; DAPI), N-acetyl-L-cystein (NAC), mixed tocopherols (vitamin E), Nrf2 antibodies, and phosphospecific p38 (P-p38), while ERK1/2 (P-ERKs) from Cell Signalling Technology Inc. From Life Technologies were obtained: wheat germ agglutinin conjugates (WGA) Alexa Fluor® 488 conjugated lectin, Alexa Fluor® 594 phallodin, 5,5',6,6'-tetrachloro-1,1',3,3'-tetraethyl benzimidazolyl carbocyanine iodide (JC-1), SB203580 (p38 inhibitor) and Alexa Fluor® 594/488 anti-mouse or anti-rabbit IgG dyes (secondary antibodies).

All used chemicals were of the highest commercially available grade.

## 2.2. Analytical chemistry methods: HPLC and LC-MS

Residues of IMI and its TPs were extracted from analysed cell/invertebrate organism samples. Briefly, the samples were processed using a mixture of methanol/water (3:1; v/v) with previous pre-treatment that includes homogenisation or lysis. After extraction the samples were filtered (0.45  $\mu$ m pore size) and injected into the HPLC, chromatographed under isocratic reversed phase conditions and detected by MS with electrospray ionisation, in the case of LC-MS.

To ensure reliable toxicity data, the stability of IMI and 6CNA was checked. The exposure was confirmed measuring the concentrations of the specific chemicals at the beginning and end of the experimental trial, under the same condition as all toxicity tests (described in Sections 2.3.1; 2.4.2.1 and 2.4.2.2). IMI and 6CNA samples were taken in duplicates and all determinations were performed in three experiments.

IMI and 6CNA were extracted with the use of the miVac centrifugal concentrator Modular Series (Genevac). The water was evaporated (under maintained vacuum conditions at 70 °C for approximately 200 min). The dried leftover was re-dissolved in 500 μL of double deionised H<sub>2</sub>O. All prepared samples were stored in glass vials under dark at 4 °C until subjected to HPLC-DAD (UV-Vis). Previous procedure was applied for samples with lower concentration of chemicals (used for tests with amphipods), while samples with higher concentration of chemicals (used for tests with algae and F11 cell) were analysed immediately without pre-concentration step. For quantification purposes, calibration curves were prepared in the appropriate concentration range. The r<sup>2</sup> value of the regression line for IMI was 0.9999 and for 6CNA was 0.9996.

Aqueous solutions of IMI and 6CNA were analysed by HPLC-DAD (UV-Vis) consisting of an Agilent 1100 Series chromatograph, coupled with a DAD detector operating in the UV-Vis range. The separation was achieved using a Zorbax C8 column (250 mm x 4.6 mm) filled with a stationary phase Chromasil 100 (pore size 5  $\mu$ m, end-capped) produced by BIA Separations d.o.o., Slovenia. The column thermostat was maintained at 25 °C and injection volume was 75  $\mu$ L. According to Žabar *et al.* methods for IMI (2012) and 6CNA (2011) detection were applied. For IMI detection the eluents consisted of 30 % acetonitrile (A) and 70 % acetic acid

0.75 % v/v (B); isocratic elution; flow rate was 1 mL min<sup>-1</sup>. The wavelength was 270 nm and the retention time was 8.9 min. While for the 6CNA detection the eluents consisted of acetonitrile (A) and acetic acid 1.5 % v/v (B); flow rate was 1 mL min<sup>-1</sup>. The gradient elution was as follows: 0–16 min 15 % A; 16–20 min 70 % A. The wavelength was 242 nm and the retention time for 6CNA was 13.2 min.

### 2.3. *In vitro* models

#### 2.3.1. Neuronal F11 cell line

F11 cell lines were grown in DMEM (4.5 g L  $^{-1}$  glucose; with *L*-glutamine) and penicillin/streptomycin (10,000 units mL $^{-1}$  penicillin and 10 mg mL $^{-1}$  streptomycin in 0.9 % sodium chloride), supplemented with 10 % (v/v) of heatinactivated FBS in a humidified cell culture incubator at 37 °C under 5 % CO<sub>2</sub> as a buffer system.

For cell culture and experimental exposure F11 cells were plated at 1-5 x 10<sup>4</sup> cells/cm<sup>2</sup> (corresponding to 5,000-20,000 cells/well) on different types of tissue culture plates (uncoated 48 or 96-well; based on selected method) in DMEM complemented with 10 % FBS and were grown for 24 h before being used. In 6-well plates F11 cells were plated at 2 x 10<sup>5</sup> cells/cm<sup>2</sup> (70 000 cells/well). For microscope cell imaging cells were plated on plastic Petri dishes (35 mm x 10 mm) with PLL (10 µg mL<sup>-1</sup>) pre-coated glass coverslips at a concentration of 3 x 10<sup>5</sup> of F11 cells mL<sup>-1</sup> (initial density of 9000 cells per 12-mm glass coverslip) in DMEM medium complemented with 10 % FBS and were grown for 24 h before being used.

Cells were exposed to vehicle (0.004 % v/v; DMSO) alone as control group and to a.i. IMI, Confidor 200SL, 6CNA, desnitro-IMI, 5-hydroxy-IMI and olefin IMI solutions as exposure groups for 24/48 h. A stock solution of 1 M IMI and 6CNA (255.6 g L<sup>-1</sup> and 157.5 g L<sup>-1</sup>), as well of all TPs, was prepared in DMSO (solubility of IMI in DMSO is >200 g L<sup>-1</sup>) with constant mixing until complete dissolving. The test solutions were prepared by adding an appropriate volume of the stock solution in the culture medium to achieve final exposure concentrations. For all compounds was prepared an equal molar concentration range expressed in corresponding mg L<sup>-1</sup> values as presented in Table 7. The toxicity of known co-formulants incorporated in

Confidor SL 200 (as negative control – a solution consisting of 38.4 % of dimethylsulfoxide (DMSO), 37.5 % of N-methyl-2-pyrrolidone (NMP) and 24.1 % of DMEM in place of IMI) was also tested in the same range as Confidor 200SL (%; v/v). For Confidor 200SL the final concentrations were corresponding to 0.0001–0.5% (v/v) which contained 0.25–1022.4 mg L<sup>-1</sup> of a.i. IMI.

In certain experiments, cells were pre-treated with N-acetyl cysteine (NAC; 1 mM, 30 min), mixed tocopherols (vitamin E; 1 mM, 30 min) or with the p38 inhibitor SB203580 (10  $\mu$ M, 30 min), to block possible IMI effects and selected molecular pathways.

Table 7: Selected concentration range for all tested compounds during experiments with F11 cell culture.

$\mu M^*/mg L^-$	IMI	Confidor 200SL (a.i. IMI; % v/v)	6CNA	desnitro-IMI	olefin-IMI	5-hydroxy-IMI
1	0.25	0.25 (0.0001)	0.15	0.25	0.25	0.27
30	7.6	7.6 (0.003)	4.7	7.4	7.6	8.1
100	25.6	25.6 (0.012)	15.7	24.7	25.3	27.2
200	51.1	51.1 (0.025)	31.5	49.4	50.7	54.3
500	127.8	127.8 (0.06)	78.7	123.6	126.8	135.8
1000	255.6	255.6 (0.12)	157.5	247.1	253.6	271.6
4000	1022.4	1022.4 (0.5)	630	988.5	1014.4	1086.4

\*To note: in the graphical representation of F11 results all applied concentrations were shown as molar values. Equal molar concentrations were applied for better comparison of data.

### 2.3.1.1. MTT cytotoxicity assay

MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide) assay was used to measure the amount of metabolically active cells. Indirectly the assay can also determine cytotoxicity and growth rate. The MTT assay is based on the protocol described for the first time by Mossmann (1983) and is based on the conversion of yellow MTT to the purple end-product formazan by succinate-tetrazolium mitochondrial reductase, a system that belongs to the respiratory chain of metabolically active cells (Fig. 25). This assay was optimised for the F11 cell line and used in this study.

Figure 25: Scheme showing the reduction of MTT tetrazolium salt to formazan (http://en.wikipedia.org/wiki/File:Mttscheme.png).

For testing the cytotoxic effects of IMI and its TPs, assays of viability were performed in 96-well cell-culture plates. Cells were seeded at a density of 3 x  $10^4$  cells/cm<sup>2</sup> (10,000 cells/well) and cultured for 24 h before being exposed to all tested chemicals in the selected concentration range (Table 7). In addition, only in presence of IMI (1000  $\mu$ M; 255.6 mg L<sup>-1</sup> and 4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) for 24 and 48 h, cells were pre-treated with *N*-acetyl cysteine (NAC; 1 mM, 30 min), mixed tocopherols (vitamin E; 1 mM, 30 min) or with the p38 inhibitor SB203580 (10  $\mu$ M, 30 min).

Briefly, for the purpose of MTT experiments at the end of the exposure in 96-well microplates (for 24 and 48 h) cells were incubated for 2 h with 500 μL of 0.5 mg mL<sup>-1</sup> of MTT solution in DMEM cell medium without phenol red. After the incubation period, the MTT solution was carefully decanted off and formazan was extracted adding 220 μL of DMSO in each well with gentle shaking for 5 min until complete dissolution was achieved. The plates were read at 550 nm (test wavelength) and 630 nm (reference wavelength) using Tecan F200 Infinite multiplate reader (Tecan Austria GbmH, Austria). Each group was run in 2 or 3 wells and each assay was triplicated (n=9). After analyses the absorbance values and all results were presented as percentage of the control values (viability/control; % of cell survival). Percentage of cell survival and in this case toxicity of the tested compounds is expressed with the following relationship [1]:

#### % of cell survival =

[1]

### 2.3.1.2. Biochemical biomarkers assays in F11 cells

F11 cells were cultured in 6-well plates F11 at 2 x 10<sup>5</sup> cells/cm<sup>2</sup> (70 000 cells/well) in DMEM medium supplemented with 10% FBS for 24 h before addition of the various tested compounds (Table 7). All assays were run in three separated and repeated experiments (n=3), each with two or three replicates.

**LP** was estimated *in vitro* after the formation of MDA, a major by-product of LP that reacts with thiobarbituric acid (TBA). LP was measured as TBARS release into the culture medium. After pre-culturing, cells were incubated for 24 and 48 h in DMEM medium without phenol red to IMI (1000 μM; 255.6 mg L<sup>-1</sup> and 4000 μM; 1022.4 mg L<sup>-1</sup>). Afterwards, the medium was centrifuged for 10 min at 1,000 g and TBARS were measured by mixing the supernatant with 0.7 % TBA and 25 % trichloroacetic acid (TCA) and heating at 90 °C for 10 min. Butylated hydroxytoluene (BTH; 0.5 mM) was included to prevent sporadic LP. TBARS were extracted with addition of 3 mL 1-butanol and the absorbance was measured using the Perkin Elmer Lambda 25 UV/VIS spectrophotometer system at 535 nm and 600 nm, the last one to correct the non-specific turbidity. TBARS were expressed as ng MDA/mg protein using 1,1,3,3-tetramethoxypropane as MDA standard (as fold increase/decrease of control) (Zampieri *et al.*, 2009).

CAT activity in cell lysate was determined according to the method of Aebi (1974) and Jamnik *et al.* (2003) by measuring the decrease in absorbance at 240 nm for 2 min due to the decomposition of  $H_2O_2$  in a Perkin Elmer Lambda 25 UV/VIS spectrophotometer system. After exposure period of 24 and 48 h to IMI (1000  $\mu$ M; 255.6 mg L<sup>-1</sup> and 4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>), the attached cells were washed in PBS (100 mM; pH 7) and lysed in 250  $\mu$ L of appropriate lysis buffer (50 mM Tris-HCl, pH 8; 150 mM NaCl), scraped and collected in a tube. Cells were sonicated on ice two times for 10 s and centrifuged at 12,000 g for 10 min at 4 °C and the supernatant was collected and used for further analyses. The reaction mixture (2 mL) contained 0.1 mL of cell supernatant in PBS (50 mM, pH 7.0) and 1.9 mL of 10 mM  $H_2O_2$  in PBS (100 mM; pH 7.0). An extinction coefficient for  $H_2O_2$  at 240 nm of 40 L cm<sup>-1</sup> mol<sup>-1</sup> (Aebi, 1974) was used for the calculation. The specific activity of CAT was expressed as  $\mu$ moles of  $H_2O_2$  reduced per minute per mg protein (as fold increase/decrease of control).

All the data relative to biochemical biomarkers are normalised to the **total protein content**. Cell lysate used for the analysis was used also for protein determination. Protein concentration in the cell extract was determined by the Bradford method (1976) using bovine serum albumin (BSA) as a standard. This method is based on the reaction between Coomasie Brillant Blue G-250 (Bradford reagent) and the proteins present in the sample to form a complex. Coomassie Brillant Blue G-250 exists in two different colours forms, red and blue. The red form is converted to blue form upon binding of the dye to the protein. The binding of the dye to the protein is a very rapid process (approximately 5 minutes) and the protein-dye complex remains dispersed in the solution for one hour. During the procedure, 1.5 mL of Bradford reagent was added to 50  $\mu$ L of supernatant and mixed up in two replicates for each animal. The absorbance of the blue coloured complexes was measured by the spectrophotometer at 595 nm after at least 2 hours. The standards were made with BSA. As blank a mixture of 1.5 mL Bradford reagent and 50  $\mu$ L of PBS were used.

## 2.3.1.3. Immunofluorescence microscopy

For microscope imaging F11 cells were cultured in plastic Petri dishes (35 mm x 10 mm) with PLL (10  $\mu$ g mL<sup>-1</sup>) pre-coated glass coverslips at a concentration of 3 x 10<sup>5</sup> of F11 cells mL<sup>-1</sup> (initial density of 9000 cells per 12-mm glass coverslip) in DMEM medium supplemented with 10 % FBS for 24 h before addition of IMI. For analyses of cell morphology after exposure to IMI (1000  $\mu$ M; 255.6 mg L<sup>-1</sup> and 4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) the incubation lasted for 24 and 48 h in DMEM medium with or without FBS.

For p38 and ERK induction experiments, exposure to IMI was at 400  $\mu$ M (102.2 mg L<sup>-1</sup>) and 4000  $\mu$ M for 30 min and for a time-course experiment at 5, 10, 30 and 250 min incubation time (4000  $\mu$ M IMI).

For Nrf2 experiment exposure to IMI was during a time-course experiment (at 10, 30, 60, 120, 240 min incubation time) at 4000  $\mu$ M concentration. All assays were run in three separated experiments (n=3), each with ten replicates.

F11 cultures were processed for IF staining with standard protocols. F11 cells were fixed in 4 % paraformaldehyde (in PBS 100 mM; pH 7.4) for 20 min at RT.

After washing twice with ice cold PBS, the cells were incubated in blocking solution (5 % BSA, 1 % FBS and 0.1 % Triton X-100 in PBS) for 20 min at RT. After the blocking of paraformaldehyde (PFA) reactive amino residues and unspecific antigens with blocking solution cell glass-coverslips were incubated on a piece of parafilm with primary antibody in 1 % BSA/PBS in a humidified chamber for 1 h. The following specific antibodies were used: neuronal specific β-tubulin III (1:1000, Sigma Aldrich), anti-active phosphorylated p38 MAP Kinase (1:1000, Life Technologies), anti-active phosphorylated ERK 1,2 (1:300, Cell Signaling) and Nrf2 (1:200; Santa Cruz Biotechnology - to test Nrf2 expression, cells were fixed in methanol for 5 minutes at -20 °C).

After three washing steps with PBS, cells were incubated with fluorescence labelled secondary antibody in 1 % BSA for 40 min at RT in dark. For secondary immunostaining, AlexaFluor 488- or 594-conjugated antibodies (1:500; Life Technologies) were used (secondary staining alone gives no signal). All secondary antibodies are labelled with suitable fluorofores with ex. at 488 nm for green and at 594 nm for red colour. Cellular actin was visualised directly with phallodin conjugated with AlexaFluor594 (1:500, Life Technologies), while membrane integrity was evaluated with *in vivo* labeling with WGA-AlexaFluor 488 conjugated lectin (1:500, Life Technologies). After incubation with secondary antibodies cell coverslips are washed three times with PBS nuclei were counterstained with the use of DAPI, a DNA intercalator (1 µg mL<sup>-1</sup>; Sigma Aldrich) with ex. at 360 nm for blue colour. As a final step, glass coverslip with cells are mounted with a glycerol-based anti-bleaching solution VECTASHIELD® medium (Vector Labs, UK) to prevent rapid loss of fluorescence during microscopic examination.

Samples were analysed with Axioscope fluorescence microscope (Zeiss) equipped with AxioVision software (version 4.6.2.0; Zeiss). Images were acquired under controlled exposure and excitations below saturation value, and processed for background subtraction and average immunofluoresence intensity quantification with Image J software (Collins, 2007). A minimum of 200 cells were analysed per each experiment.

### 2.3.1.4. Evaluation of mitochondria through microscopy imaging

Alterations in mitochondrial membrane potential (MMP) were qualitatively assessed with the use of 5,5',6,6'-tetrachloro-1,1',3,3'-tetraethyl benzimidazolyl carbocyanine iodide (JC-1) (Hardiek *et al.*, 2001). JC-1 has advantages over other cationic dyes as it can selectively enter into mitochondria and reversibly change colour from green to red as the membrane potential increases. In healthy cells with high MMP, JC-1 spontaneously forms complexes known as J-aggregates with intense red fluorescence. On contrary, in unhealthy or apoptotic cells with low MMP, JC-1 remains in the monomeric form, which shows only green fluorescence. JC-1 dye exhibits potential-dependent accumulation in mitochondria, indicated by a fluorescence emission shift from green (~529 nm) to red (~590 nm). Consequently, mitochondrial depolarisation is indicated by a decrease in the red/green fluorescence intensity ratio (Fig. 26).

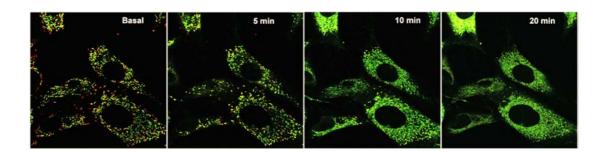


Figure 26: Representative example of JC-1 dye in action. NIH 3T3 fibroblasts stained with JC-1 showing the progressive loss of red J-aggregate fluorescence and cytoplasmic diffusion of green monomer fluorescence following exposure to hydrogen peroxide for 5, 10, and 20 min after treatment (www.invitrogen.com).

For JC-1 *in vivo* microscope imaging F11 cells were cultured in glass-bottom Petri dishes (35 mm x 10 mm) at a concentration of 3 x  $10^5$  of F11 cells mL<sup>-1</sup> in DMEM medium supplemented with 10% FBS for 24 h before addition of IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) and incubation for 15 and 60 min.

A stock solution of JC-1 (200  $\mu$ M, Life Technologies) was prepared according to manufacturer instructions. Previously prepared JC-1 solution is added (1:1000) to the IMI-treated cells in each well and cells are incubated for 15 min at 37 °C in a CO<sub>2</sub> incubator. After incubation period, cell can be analysed directly in the

culture medium since phenol red does not interfere with fluorescent staining. Healthy cells with mainly JC-1 aggregates can be detected at ex. 560 nm and em. 595 nm, while apoptotic or unhealthy cells with mainly JC-1 monomers can be detected at ex. 488 nm and em. 535 nm. Hydrogen peroxide (100 µM; 30 min), was used as a positive control for decreased MMP.

Samples were analysed with Axioscope fluorescence microscope (Zeiss) equipped with AxioVision software (version 4.6.2.0; Zeiss) with appropriate fluorescence settings. Images were acquired under controlled exposure and excitations below saturation value, and processed for background subtraction and average immunofluoresence intensity quantification with Image J software (Collins, 2007). A minimum of 200 cells were analysed per each experiment. JC-1 assay was run in three separated and repeated experiments (n=3), each with ten replicates. Fluorescence values were expressed as the ratio of fluorescence intensity of Jaggregates (red) to fluorescence intensity of monomers (green), which is often used as an indicator of cell health status. Furthermore, ratio results were presented as fold increase/decrease of the control values (JC-1/control in arbitrary units – AUs).

### 2.3.2. Ames mutagenicity test with Salmonella typhimurium

A set of histidine-requiring strains (TA 98 and strain TA 100) was used for a mutagenicity test called Ames test. Strain TA 98 contains as a marker the frame-shift mutation hisD3052, whereas TA 100 bears the base pair substitution hisG46. Strains are inoculated separately in a Bacto nutrient broth growth medium (20 mL) supplemented with 20  $\mu$ L of ampicillin solution. Bacteria are grown up overnight at 37 °C under sterile conditions in closed Erlenmeyer flasks permeable to air with caps and mixed by gentle agitation of at least 150 revolutions per minute (rpm). Earlier are also prepared minimal agar glucose plates under sterile conditions.

Test solutions of IMI, Confidor 200SL and 6CNA were prepared at 30; 100 and 1000  $\mu$ M concentration corresponding to 7.6; 25.6 and 255.6 mg L<sup>-1</sup> for IMI and Confidor 200SL and 4.7; 15.7 and 157.5 mg L<sup>-1</sup> for 6CNA. These concentrations are equivalent to: 0.053; 0.178; 1.78  $\mu$ g/plate for IMI and IMI a. i. in Confidor 200SL, 0.265; 0.89; 8.9  $\mu$ g/plate for Confidor 200SL (whole formulation) and 0.033; 0.11; 1.11  $\mu$ g/plate for 6CNA, which is the usual concentration representation in Ames

test. All test solutions are sterilised by filtration through a sterile filer with  $0.45~\mu m$  pore size. IMI and 6CNA were dissolved in DMSO, which was used at a final concentration of less than 0.004~% (v/v).

For the mutagenicity assay, the control and tested sample solution were mixed with bacterial strains (TA98 or TA100), S9 fraction and 2 mL of sterile top agar (0.6 % agar and 0.5 % NaCl containing 0.5 mM histidine and 0.5 mM biotin) and poured onto previously prepared minimal glucose agar plate [1x Vogel-Bonner salts (0.2 g L<sup>-1</sup> magnesium sulphate, 2 g L<sup>-1</sup> citric acid monohydrate, 10 g L<sup>-1</sup> dipotassium hydrogen phosphate, and 3.5 g L<sup>-1</sup> sodium ammonium phosphate), 2 % glucose, and 1.5 % agar]. By adding the S9 fraction (homogenate of rat live) directly to the Petri plates it is incorporated in the in vitro test an important aspect of the metabolism, making this test highly efficient (Ames et al., 1975). S9 treatment for enzyme induction and preparation of the S9 fraction was performed according to the guideline of ISO 16240 (2005). Arranged plates were then incubated in dark at 37 °C for 48 h. Revertant colonies appeared on a background lawn of bacteria. Spontaneous revertants in case of strain TA 98 are around 40, while for strain TA 100 near 100 colonies. Two independent experiments were conducted; each experiment consisted of three replicate plates for each treatment. All steps were performed under sterile conditions.

### 2.4. In vivo models

### 2.4.1. Toxicity test with luminescent bacteria Vibrio fischeri

The toxicity tests with luminescent bacteria were performed to determine of 30 min EC<sub>50</sub> values for aqueous solutions of three TPs of IMI. The toxicity of desnitro-IMI, olefin-IMI and 5-hydroxy-IMI (at initial concentrations of 100 mg L<sup>-1</sup>) was assessed using liquid-dried luminescent bacteria V. fischeri NRRL B-11177 with system LUMIStox, Dr. LANGE determining reduced luminescence emission after incubation with tested chemicals. Before analyses all samples were adjusted for pH at  $7 \pm 0.2$  with hydrochloric acid or sodium hydroxide solutions and added with 2 % w/v of sodium chloride salt in order to avoid possible adverse effects due to an improper pH value or unsuitable sodium chloride concentration. All luminescence values were measured with a photomultiplier LUMIStox 300 luminometer at controlled 15  $\pm$  1 °C. The toxicity test was performed in three essential steps: (1) an aliquot containing V. fischeri was added to each vial in two parallels and luminescence was measured immediately; (2) tested samples were added to vials (at various geometrical dilution levels) with bacteria and regulated to  $15 \pm 1$  °C for 30 min; (3) the luminescence of bacteria within the sample was again measured after 30 min of exposure. All dilution levels were achieved following ISO 11348-2 standard guideline (2007). The blank test was performed with 2 % w/v sodium chloride solution.

The 30 min EC<sub>50</sub> was calculated with 95 % confidence limits, according to ISO 11348-2 standard (2007) using a computer software supported model. The results are presented as luminescence inhibition in percentage and consequently, the 30 min EC<sub>50</sub> or EC<sub>20</sub> values for desnitro-IMI, olefin-IMI and 5-hydroxy-IMI were determined. This way of presenting the toxicity results is commonly adopted in the literature (Sakkas *et al.*, 2004; Dell'Arciprete *et al.*, 2009, 2010; Kitsiou *et al.*, 2009). In addition, percentage inhibitions obtained in each experiment were converted to gamma values according to first model (Jennings *et al.*, 2001), where:

Gamma (
$$\Gamma$$
) = % inhibition / (100 - % inhibition) [2]

Gamma values were plotted against their corresponding chemical concentrations, after first converting all data to natural logs (Ln), to generate Ln gamma/Ln concentration curves for each chemical. In this model, values falling within the 10-90 % inhibition range were used to fit a straight line to the Ln-transformed data by linear regression and liner regression equations have been used to calculate  $EC_{20}$  or  $EC_{50}$ . To note:  $EC_{50}$  value is given by the point of intersection with the x-axis at  $\Gamma = 1$ , while  $EC_{20}$  at  $\Gamma = 0.25$ .

## 2.4.2. Green microalgae and crustacean amphipod as test organisms

*D. subspicatus* (Chodat) Hegewald et Schmidt (formerly *S. subspicatus*) was kindly provided by the Helmholtz Centre for Environmental Research-UFZ, Leipzig, Germany. Microalgae were grown in a medium recommended by standard guidelines for freshwater algal growth inhibition test (ISO 8692, 2004).

*G. fossarum* were collected in April–July 2011 using a net (by the kick sampling method) from the stream Vogršček (Slovenia). The sampling site is in the lower Vipava Valley in Goriška region of Slovenia (45°90' N; 13°70 E). It is a small waterbed free of industrial, agricultural contamination or human activities and it can be considered unpolluted. This site has a good water quality according to ARSO data record (Agencija Republike Slovenije za okolje-Slovenian Agency for Environment; http://www.arso.gov.si/vode/poro%C4%8Dila%20in%20publikacije/povrsinske\_letn a.html) and high densities of gammarids were found.

All water samples from the sampling site and during the experimental trial were monitored for temperature, pH, conductivity, oxygen concentration and saturation with a multi-meter WTW350i (with microelectrode replacements for small volumes). In addition, TOC and TN were measured in water samples from the sampling site with a TOC Analitik Jena multi N/C 3100, calibrated with potassium hydrogen phthalate. Before being processed for the TOC and TN analyses, samples were acidified to pH 2–3 with hydrochloric acid.

Gammarids were kept during an acclimatisation period of at least 14 days in a 20 L glass aquarium supplied with thoroughly aerated original stream water. An 8/16 h light/dark natural photoperiod was maintained with the temperature at 12  $\pm$  2 °C in a temperature and humidity controlled chamber and regular water renewal every two

days. Animals were fed *ad libitum* twice a week using a pinch of dry food (e.g. TetraMina<sup>®</sup> flakes) or raw peas.

### 2.4.3. Algae toxicity test

Sub-chronic toxicity of pesticides was conducted in 96 microwell plate. The algal inoculum was taken from an exponentially growing pre-culture and added into 25 mL of growth media in order to obtain an initial cell density of 10<sup>4</sup> cells mL<sup>-1</sup>. Final volume of each well was 200 µL. Serial dilutions of tested pesticides were made in culture medium. Six replicates of controls (untreated) and three replicates of each test concentration were applied. All the plates with cover, control and treatments, were incubated for four days (96 h) at a temperature of 23 ± 1 °C and light intensity of 1100 lux. Algal growth was detected fluorometrically in intervals of 24 h over a period of 96 h in order to achieve a virtual kinetic data distribution. Analyses of chlorophyll fluorescence were performed by a Tecan Infinite® 200 PRO (Männedorf, Switzerland). Measurements were conducted using fluorescence ex. of 440 nm and by an em. of 680 nm. Before reading, tested microplates were shaken for 30 s at 100 rpm. Average of specific growth rates were calculated and subsequently used for calculation of percentage inhibition in comparison to control (Kaczala et al., 2011). IC<sub>50</sub> at 96 h (inhibition concentration that cause 50 % inhibition of algal growth) was estimated for tested compounds using linear regression analysis (Tišler et al., 2009).

Solution of 1 M IMI and 6CNA was prepared in DMSO (solubility of IMI in DMSO is >200 g L<sup>-1</sup>). Afterwards, a 10 mM (2.55 g L<sup>-1</sup> for IMI and 1.57 g L<sup>-1</sup> for 6CNA) stock solution was prepared by the addition of IMI and 6CNA (1 M) or Confidor 200SL to standard algal medium, with constant mixing until complete dissolving. The test solutions were prepared by adding an appropriate volume of the stock solution in the algal medium to achieve final concentration. For all compounds was prepared an equal molar concentration range expressed in corresponding mg L<sup>-1</sup> values as presented in Table 8. For Confidor 200SL the final concentrations were corresponding to 0.003–0.12% (v/v) which contained 7.6–255.6 mg L<sup>-1</sup> of a. i. IMI. The toxicity of known co-formulants incorporated in Confidor SL 200 (as negative control – a solution consisting of 38.4 % of DMSO, 37.5% of NMP and 24.1% of

double deionised water in place of IMI) was also tested in the same range as Confidor 200SL.

Table 8: Selected concentration range for all tested compounds during experiments with algae D. subspicatus.

μM/ mg L <sup>-1*</sup>	IMI	Confidor 200SL (a.i. IMI; % v/v)	6CNA
30	7.6	7.6 (0.003)	4.7
100	25.6	25.6 (0.012)	15.7
200	51.1	51.1 (0.025)	31.5
500	127.8	127.8 (0.06)	78.7
1000	255.6	255.6 (0.12)	157.5

\*To note: in the graphical representation of algae toxicity test all applied concentrations were shown as  $mg L^{-1}$ values.

Lower concentrations of IMI than those monitored in this experimental trial were already tested on D. subspicatus and showed no effect on algal growth up to 10 mg L<sup>-1</sup> and due to this fact were excluded (Heimbach, 1986). In addition, as an internal quality control, the bioassays were also performed on the reference chemical potassium dichromate (positive control –  $K_2Cr_2O_7$ ; 0.1–30 mg L<sup>-1</sup> (ISO 8692, 2004).

### 2.4.4. Amphipods toxicity test

Gammarids were exposed for 24 h (acute toxicity) to equal molar concentrations of IMI, Confidor 200SL and 6CNA for better comparison. A short exposure period sufficient to promote early alterations (24 h) was used also to mimic runoff-related pulse exposures to pesticides (Pick *et al.*, 1984; Werner *et al.*, 2004). The peak pesticide concentrations usually persist for about 24 h. Furthermore, *G. fossarum* from running water is greatly affected by short-term higher concentration of IMI (Lukančič *et al.*, 2010). Selected sub-lethal exposure concentrations were based on previously determined acute LC<sub>50</sub> (48 h) and EC<sub>50</sub> (24 h) values for IMI of 0.8 and 0.07 mg L<sup>-1</sup> (Lukančič *et al.*, 2010).

Solution of 1 M IMI and 6CNA was prepared in DMSO (solubility of IMI in DMSO is >200 g L<sup>-1</sup>. Afterwards, a 10 mM (2.55 g L<sup>-1</sup> for IMI and 1.57 g L<sup>-1</sup> for

6CNA) stock solution was prepared by the addition of IMI and 6CNA (1 M) or Confidor 200SL to distilled water, with constant mixing until complete dissolving. The test solutions were prepared by adding an appropriate volume of the stock solution in the original stream water to achieve final concentration. For all compounds was prepared, as mentioned, an equal molar concentration range expressed in corresponding mg L<sup>-1</sup> values as presented in Table 9. For Confidor 200SL the final concentrations were corresponding to 0.000003-0.0002% (v/v) which contained 6.3-511.3 μg L<sup>-1</sup> of a.i. IMI. The toxicity of known co-formulants incorporated in Confidor SL 200 (as negative control – a solution consisting of 38.4 % of DMSO, 37.5 % of NMP and 24.1 % of double deionised water in place of IMI) was also tested in the same range as Confidor 200SL.

The experimental trial was performed using adult male specimens. After sex determination, total body length (Becuhel and Lønne, 2002) (Fig. 27) and total wet weight was measured (animal were dried between two sheets of filter paper before being weighted). Fifty individuals per exposure concentration were used for every tested compound. Plastic Petri dishes (100 mm x 20 mm; 20 mL volume) covered in order to reduce water evaporation were used for exposure experiments. The bioassays were conducted in darkness, in a temperature and humidity-controlled chamber (12 ± 2 °C; 60 % humidity). After a 24 h exposure period, immobility or moulting and mortality were observed. Live/dead organisms were determined by gently poking and observing movement of appendages. Organisms were counted as dead if none of the appendices were moving after poking for three times. Inactive/paralysed animals were identified when only respiration movements were left (Clesceri et al., 1998). Moulted animals were counted based on the presence of the entire old exuvia in the exposure vessel (moulted amphipods were not used for biochemical parameters analyses). For each biochemical assay 10 randomly selected gammarids per concentration (from fifty individuals) were processed using wholebody homogenates due to their small body size.

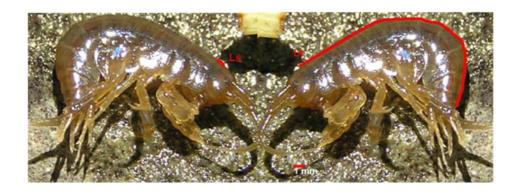


Figure 27: Red lines illustrate the measured length of first pereonal segment (Ls) and total body length (Lt) of G. fossarum (Vihtakari, 2008 with slight modifications).

Table 9: Selected concentration range for all tested compounds during experiments with amphipod G. fossarum.

μ <b>Μ</b> / μg L <sup>-1*</sup>	IMI	Confidor 200SL (a.i. IMI; % v/v)	6CNA
0.025	6.3	6.3 (0.000003)	3.9
0.05	12.7	12.7 (0.000006)	7.8
0.1	25.5	25.5 (0.000012)	15.7
0.2	51.1	51.1 (0.000025)	31.4
0.4	102.2	102.2 (0.00005)	62.8
0.6	153.3	153.3 (0.00007)	94.6
0.8	204.5	204.5 (0.0001)	126.2
1	255.6	255.6 (0.00012)	157.7
2	511.3	511.3 (0.0002)	315.5

<sup>\*</sup>To note: in the graphical representation of amphipods toxicity test all applied concentrations were shown as  $\mu g L^{-1}$  values.

### 2.4.4.1. Biochemical biomarkers assays in amphipods

Prior to individual homogenisation, excess chemicals present on the animal's surface were rinsed several times according to Jemec *et al.* (2007). Whole-body specimens were homogenised in 500 μL of ice cold PBS (100 mM; pH 7.0) for 3 min using a glass–glass Elvehjem–Potter homogeniser. The homogenate was sonicated on ice (5–10 s) and centrifuged for 15 min at 3,000 rpm and 4 °C. Freshly prepared clear supernatant was collected and kept on ice to be used for enzyme activities measurements.

**AChE activity** was determined using DTNB and acetylthiocholine iodide as substrate according to Ellman *et al.* (1961). The principle of this technique is the measurement of the rate of the production of thiocholine (an ACh analogue). Thiocholine, produced during the hydrolysis, reacts with DTNB and the final product of this reaction is a yellow anion of 5-thio-2-nitro-benzoic acid (Fig. 28).

$$\begin{array}{c} \text{enzyme} \\ \text{H}_2\text{O} \ + \ (\text{CH}_3)_3\text{N}^+\text{CH}_2\text{CH}_2\text{SCOCH}_3 & \longrightarrow & (\text{CH}_3)_3\text{N}^+\text{CH}_2\text{CH}_2\text{S}^- \ + \ \text{CH}_3\text{COO}^- \ + \ 2\ \text{H}^+ \\ \text{acetylcholine} & \text{thiocholine} \\ \text{(CH}_3)_3\text{N}^+\text{CH}_2\text{CH}_2\text{S}^- \ + \ \text{RSSR} & \longrightarrow & (\text{CH}_3)_3\text{N}^+\text{CH}_2\text{CH}_2\text{SSR} \ + \ \text{RS}^- \\ \text{thiocholine} & \text{dithiobis-} & \text{the yellow} \\ \text{nitrobenzoic acid} & \text{anion} \\ \\ \text{R} \ = \ \text{O}_2\text{N} & \longrightarrow & \text{OOC} \\ \end{array}$$

Figure 28: The principle of the Ellman technique (Ellman et al., 1961).

The reaction was followed on a Perkin Elmer Lambda 25 UV/VIS spectrophotometer at 412 nm for 8 min. Kinetic was calculated in the linear range and slope measured (the slope is the rate in absorbance units/min). Spontaneous substrate hydrolysis was assessed using two controls, a blank without ACh and a blank without the sample. The analyses were made in two replicates for each sample. AChE activity is expressed as  $\mu$ mol of substrate hydrolysed per minute per mg protein (based on  $\epsilon = 13600$  L cm<sup>-1</sup> mol<sup>-1</sup> for DTNB). The reaction mixture for AChE activity measurement was prepared as presented in Table 10.

Table 10: The reaction mixture used for AChE activity measurement in G. fossarum.

### G. fossarum

- 2380 μL of PBS (67 mM, pH 7)
- 100 μL of DTNB (0.01 M)
- 500 μL of sample
- 20 µL of acetylcholine iodide (0.075 M)

**CAT activity** was determined according to the method of Jamnik and Raspor (2003) by measuring the decrease in absorbance on a Perkin Elmer Lambda 25 UV/VIS spectrophotometer at 240 nm for 2 min due to the decomposition of  $H_2O_2$  ( $\epsilon = 40 \text{ L cm}^{-1} \text{ mol}^{-1}$ ). The specific activity of CAT was expressed as  $\mu$ mol of  $H_2O_2$  reduced per minute per mg protein. The methodology and the reaction mixture for CAT measurement are identical as for F11 cells (described in Section 2.5.2.).

**GST activity** was determined according to the protocol of Habig *et al.* (1974). The method is based on determination of the conjugated product dinitrophenyl-thioether at 340 nm produced from CDNB used as an artificial substrate and reduced glutathione. The conjugation is catalysed by GST isoenzymes (Fig. 29).

GSH + CI 
$$NO_2$$
 GST  $O_2N$   $O_2N$   $O_2N$  dinitrophenylthioether

Figure 29: The principle of the GST activity measurement (Habig et al., 1974).

The measurement of the sample was carried out on a Perkin Elmer Lambda 25 UV/VIS at 340 nm for 5 min. The reaction mixture without sample was used as control. Values were expressed as nmol of reduced glutathione and CDNB conjugate formed per min per mg protein ( $\varepsilon = 9600 \text{ L cm}^{-1} \text{ mol}^{-1}$  for CDNB). The reaction mixture for GST activity measure in *G. fossarum* is presented in Table 11.

Table 11: The reaction mixture used for GST activity measurement in G. fossarum.

# G. fossarum 900 μL of PBS (100 mM, pH 6.5) 25 μL of CDNB (60 mM) 25 μL of sample 50 of reduced glutathione (40 mM)

All the data relative to the enzymatic activity are normalised to the **total protein content** based on the method of Bradford (1976) (as described in Section 2.5.2).

LP was estimated *in vitro* after the formation of MDA, a major by-product of lipid peroxidation that reacts with TBA (Ortega-Villasante *et al.*, 2005), with slight modifications. Whole-body gammarids were rinsed, as described previously and homogenised individually in TCA-TBA-BTH reagent [15 % (w/v) TCA, 0.37 % (w/v) TBA, 1 M HCl, and 0.01 % BTH]. Samples were incubated at 90 °C for 30 min, then chilled at room temperature, added 1.2 mL of 1-butanol and centrifuged at 12,000 rpm for 10 min. Absorbance of the supernatant was measured at 535 and 600 nm, the final one to correct for the non-specific turbidity. Before the heating step, absorbance was measured at 280 nm for total protein concentration. These absorbance values of protein content were used to properly normalise absorbance values obtained for LP (Fig. 30).

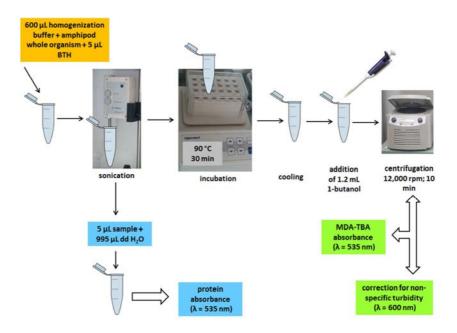


Figure 30: Schematic representation of lipid peroxidation determination in G. fossarum.

### 2.4.4.2. MXR inhibition assay

G. fossarum was exposed for 1 h to IMI, Confidor 200SL and 6CNA at 3  $\mu$ M and 30  $\mu$ M concentration (corresponding to: 0.7 and 7.6 mg L<sup>-1</sup> for IMI and Confidor 200SL, while for 6CNA to 0.4 and 4.7 mg L<sup>-1</sup>) and 0.75; 3 and 10  $\mu$ g L<sup>-1</sup> copper a.i. During this exposure to tested chemicals was also added a specific model substrate for P-gp called rhodamine B (Rh B). In our preliminary experimental setup the exposure of 1 h resulted in optimal experimental conditions and response.

To test for MXR activity and inhibition, an Rh B exclusion dye assay was adopted. As an indicator of membrane transport activity was used a fluorescent dye called Rh B. Under normal MXR function, amphipods exposed to Rh B will actively expel the dye from cells and the whole-body will have a low fluorescence. On the other hand, amphipods exposed to potential MXR inhibitors are less able to transport Rh B from inside the body, and show increased fluorescence. This increased fluorescence proves the accumulation of Rh B in cells as a consequence of transport activity inhibition. This method was already successfully applied in several studies including aquatic organisms (Smital and Kurelec, 1997; Kurelec *et al.*, 2000; Smital *et al.*, 2000). In addition, there is also available proof of presence of the MXR system in freshwater amphipods (Timofeyev *et al.*, 2007).

Briefly, five specimens of *G. fossarum* were exposed to each concentration of IMI, Confidor 200SL and 6CNA + Rh B (2.5  $\mu$ M) in Plastic Petri dishes (100 mm x 20 mm; 20 mL volume) covered in order to reduce water evaporation. The assays were conducted for 1 h in darkness, in a temperature and humidity-controlled chamber (12 ± 2 °C; 60 % humidity). Model inhibitors of MXR proteins, verapamil (VER; 10  $\mu$ M) and cyclosporine A (CA; 10  $\mu$ M) was run as a positive control alongside all treatments. After 1 h exposure amphipods were removed from their solutions and triple washed in fresh PBS to remove any external Rh B, weighed, transferred to flat-bottom tubes in 0.5 mL of distilled water and homogenised (for homogenisation procedure see Section 2.6.1). The homogenates were centrifuged at 3,000 rpm for 15 min, the supernatants carefully transferred to a 96-well flat-bottom dark microplate (100  $\mu$ L/well, each supernatant in triplicate), and the fluorescence of accumulated Rh B measured immediately.

The fluorescence of accumulated Rh B in whole-body of amphipods was measured on a microplate fluorometer (Tecan F200 Infinite multiplate reader-Tecan Austria GbmH, Austria) using 535 nm (excitation) and 590 nm (emission) filters. Due to the possible loss of the Rh B-fluorescence intensity caused by the direct exposure to light, glasses, vials/homogenates/supernatants and microplates were light protected with aluminium foil. Fluorescence readings for each well were then normalised to total animal weight and the MXR assay was repeated in three separated experiments.

# 2.5. Statistical analyses

All statistical tests were performed using STATISTICA 7 StatSoft software. Results from each exposure trial are presented in graphs as mean  $\pm$  standard error (SE). Statistical comparisons were conducted between control and exposure data using the Student's t-test or the Mann–Whitney rank sum test after the software direct choice of parametric or nonparametric data, respectively. In addition, multiple comparisons were analysed with the One-way ANOVA and Tukey post-test. p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*) were accepted as levels of statistical significance and shown in graphical representations.



### 3. RESULTS

### 3.1. In vitro model: F11 cells

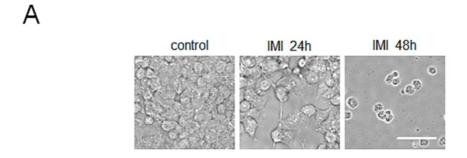
### 3.1.1. Imidacloprid cytotoxic effects

Cell viability MTT assay demonstrated limited IMI effects after 24 h and 48 h incubation at the whole micro-molar range of tested concentrations going from 1  $\mu$ M (0.25 mg L<sup>-1</sup>) up to 500  $\mu$ M (127.8 mg L<sup>-1</sup>) (p>0.05, n=9). Slightly reduced but transient effect on cell viability was noticed after 24 h exposure to IMI at 200  $\mu$ M (51.1 mg L<sup>-1</sup>) and 500  $\mu$ M (127.8 mg L<sup>-1</sup>) concentration (p<0.05). Due to the fact that no relevant and permanent adverse effect on lower concentration range was noticed we proceeded with testing even at higher concentrations reaching values of 1000  $\mu$ M (255.6 mg L<sup>-1</sup>) and 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>).

IMI evoked progressively a dose- and time-dependent lower metabolic activity of F11 cells and reduced survival compared to control especially at concentrations higher than 500  $\mu$ M (Fig. 31C). Viable cells after 24 h at 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>) were of 82.3  $\pm$  3.9 % (p<0.05, n=9), while after 48 h at 1000  $\mu$ M (255.6 mg L<sup>-1</sup>) and 4000  $\mu$ M were of 76.2  $\pm$  3.5 % and 56  $\pm$  3.8 %, respectively (p<0.001; n=9).

In general, a significant metabolic impairment was found at concentrations larger than 1000  $\mu M$  accompanied by a significant and progressive cell loss evident by bright field microscopy images (Fig. 31A).

To test F11 morphology after exposure to IMI, a panel of antibodies against cytoskeleton elements such as actin,  $\alpha$  tubulin and  $\beta$  type tubulin were applied. In Fig. 32 is shown the morphological characterisation with fluorescence microscopy of F11 cells exposed to IMI (4000  $\mu$ M) for 24 h that increases the presence of picnotic nuclei (DAPI staining), loss in plasma membrane integrity visualised with fluorescence WGA lectin and strong degradation of cytoplasmic protein components, like cytoskeleton elements (actin and tubulin). In addition to these initial, minor and evident signs of alteration of cytoskeleton, after 48 h IMI induced a drastic reorganisation of the cytoskeleton leading to cytoplasm and nuclear condensation, and finally reducing the overall adhesion of the cells to substrate (Fig. 31A).



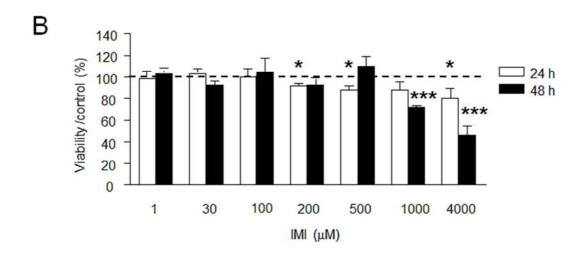


Figure 31: A. Bright field representative images of F11 cells in control and after 24 and 48 h IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) exposure. Scale bar: 50  $\mu$ m. B. Values of MTT assay in F11 cells after exposure to IMI for 24 and 48 h; data presented as percentage of control shown as dashed line (n=9). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

Further morphological characterisation of cell culture with serum and serum-free conditions in presence of IMI presented additional alterations. In control conditions cells did not have altered cytoskeleton and any significant signs of stress, while in treated cells there are present modified nuclear features and formation of protein aggregates (Fig. 33A). Also is visualised an interesting differentiation in absence of serum (as expected) with formation of processing and neuronal elongations (Fig. 33B).

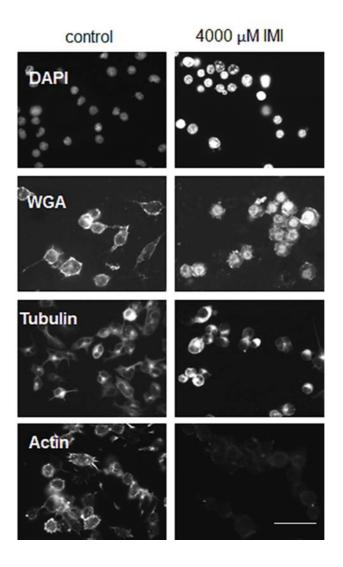


Figure 32: Representative microscopy images of F11 cells morphological characterisation in control conditions and after IMI exposure (4000  $\mu$ M; 1022.4 mg  $L^{-1}$ ) for 48 h. Cells were stained with different fluorescent markers, as indicated for each image. To note: stronger nuclear DAPI staining signal after treatment and resilient degradation of cellular proteins (actin staining). Scale bar: 100  $\mu$ m.

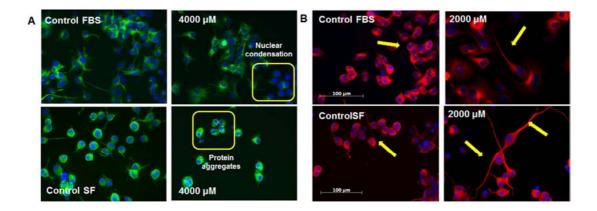
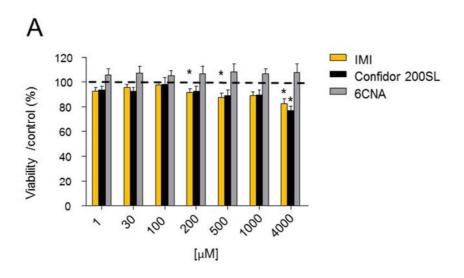


Figure 33: Morphology of F11 cells in control and after IMI exposure (4000  $\mu$ M; 1022.4 mg  $L^{-1}$ ) for 48 h in standard growing conditions (FBS) or in absence of serum (SF). A. Nuclei are counterstained with DAPI (blue),  $\beta$ -Tubulin III immunofluorescence staining is shown in green. To note: protein aggregates and nuclear condensation in F11 cells after IMI exposure (yellow rounded rectangles). B. Nuclei are counterstained with DAPI (blue),  $\beta$ -Tubulin III immunofluorescence staining is shown in red. To note: IMI stimulated processes elongation (arrows). Scale bar: 100  $\mu$ m.

To further cytotoxicity effects of IMI to neuronal cell survival, we compared potentially toxicity of IMI as its commercial formulation (Confidor 200SL - where IMI is the a.i.) and IMI's four TPs: 6CNA, desnitro-IMI, olefin-IMI and 5-hydroxy-IMI.

The negative control (mixture of known co-formulants: DMSO and NMP) did not have any adverse effects on F11 cell at all tested concentrations (data present in Annex section). Due to this fact, all values were compared to control (vehicle only; DMSO). Commercial formulation was found slightly more toxic than parent compound after 48 h of exposure at 4000  $\mu$ M; 1022.4 mg L<sup>-1</sup> a. i. (47.8  $\pm$  2.8 %; p<0.001 compared to control) (Fig. 34B). Interestingly, 6CNA decreased the F11 cells metabolic activity after 48 h already at 500  $\mu$ M (157.5 mg L<sup>-1</sup>) reaching the lowest viability of 72.2  $\pm$  3.5 % (p<0.01) at 4000  $\mu$ M (630 mg L<sup>-1</sup>). In the case of other IMI's TPs, it is important to emphasise that all significant effects on cell viability were noticed only at concentrations higher than 500  $\mu$ M for all tested TPs (Fig. 35A and 35B). After 24 h of exposure to 5-hydroxy-IMI at 4000  $\mu$ M (1086.4 mg L<sup>-1</sup>) the cell viability was significantly reduced and presented a value of 55.9  $\pm$  2.4% (p<0.01). This decreasing of cell viability was significantly enhanced after 48 h at the same concentration for all three TPs. The strongest negative effect and

drastically reduced cell survival at this highest concentration was induced by desnitro-IMI followed by 5-hydroxy-IMI and olefin-IMI. The percentage of viable cells after 48 h exposure to desnitro-IMI (4000  $\mu$ M; 988.5 mg L<sup>-1</sup>) was only of 19  $\pm$  2.7 %, while for olefin-IMI and 5-hydroxy-IMI was 24.6  $\pm$  1.04 % and 29.9  $\pm$  2.2 %, respectively (p<0.001). These adverse effects were stronger and more evident in case of IMI's TPs compared to both IMI as pure compound or as Confidor 200SL.



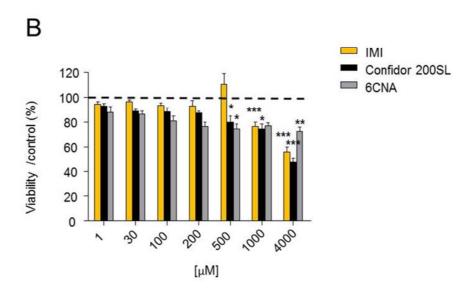
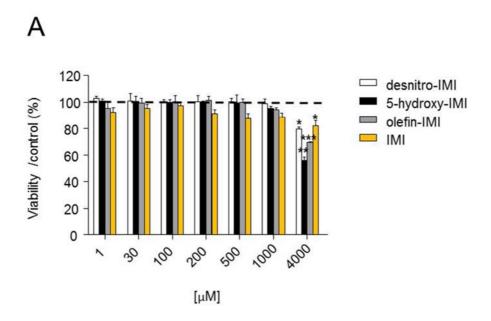


Figure 34: Values of MTT assay in F11 cells after exposure to IMI, Confidor 200SL and 6CNA for 24 h (A) and 48 h (B); data presented as percentage of control shown as dashed line (n=6). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).



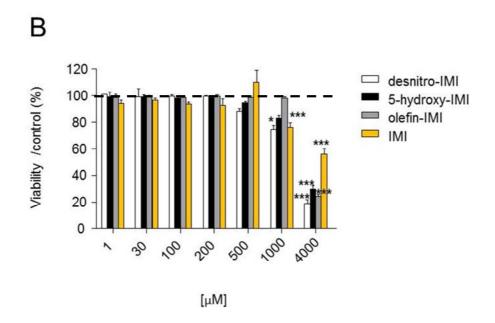


Figure 35: Values of MTT assay in F11 cells after exposure to IMI, desnitro-IMI, 5-hydroxy-IMI and olefin-IMI for 24 h (A) and 48 h (B); data presented as percentage of control shown as dashed line (n=6). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

# 3.1.2. p38, ERK MAPK and Nrf2 activation in response to imidacloprid

To further investigate the mechanisms of IMI toxicity in F11 cells and if acute IMI exposure was translated into rapid activation of downstream molecular pathways like p38 MAPK and ERK induction, we performed immunofluorescence time-course experiments (0, 5, 10, 30, 240 min) at 4000  $\mu$ m (1022.4 mg L<sup>-1</sup>) using specific antibodies against active (phosphorylated) form of kinases ERK 1,2 and p38 (Fig. 36C). In addition, IF experiments were performed on F11 treated cells exposed to different doses of IMI 400  $\mu$ M (102.2 mg L<sup>-1</sup>) and 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>) for 30 min, using again antibodies that specifically recognise activated form of p38 or ERK MAPK (Fig. 36A and 36B).

The p38 and ERK MAPK were phosphorylated and activated in response to in IMI in F11 cells. These experiments demonstrated that IMI activated these kinases in two different ways. In specific in case of p38 this activation was in a concentration dependent manner with significant induction after 30 min at 4000  $\mu$ M (p<0.01), while in case of ERK a significant and stronger activation response was present after 30 min IMI exposure already at 400  $\mu$ M (p<0.001) (Fig. 36A and 36B). Time course experiments suggested an early activation of p38 MAPK after few minutes (10 min) of IMI stimulation at 4000  $\mu$ M in the majority of the cells (92.2  $\pm$  1.3 %, p<0.01) (Fig. 36C). The p38 activation effect was further quantified and was persistent until 1 h of IMI exposure (162  $\pm$  0.01 % of the control, p>0.01). In addition, we observed also a rapid nuclear translocation of activated phosphorylated p38 MAPK after 10 min from IMI stimulation (4000  $\mu$ M) (Fig. 36C and 37). Nuclear translocation was quantified and was also persistent until 1 h with the highest values of p38 nuclear translocation after 10 min of IMI exposure (2  $\pm$  0.4 % fold increase respect to the control, p<0.01).

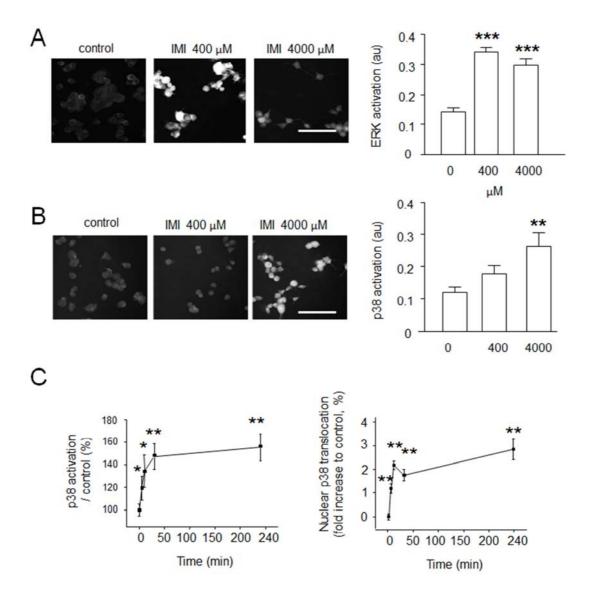


Figure 36: A, B. Representative fluorescence microscopy images of F11 cells in control conditions and in the presence of IMI 400  $\mu$ M (102.2 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) immunostained with antibodies anti-active Erk (A) or active p38 MAPK (B). Scale bar 50  $\mu$ m. Histograms quantify the grey values of immunopositive signals after background value subtraction. Data have been normalised on total cell number (counterstained with DAPI) and expressed as arbitrary units (AU). To note: different activation profile. C. Graphs quantify p38 signal intensity during the time after application of IMI at 4000  $\mu$ M to F11 cells (left) and nuclear translocation of activated phosphorylated p38 during time after IMI application (right) (n= 3), p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

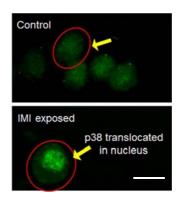


Figure 37: Representative microscopy images of translocated phosphorylated p38 MAPK in nucleus of F11 cells exposed to IMI (4000  $\mu$ M; 1022.4 mg  $L^{-1}$ ) for 10 min. Scale bar: 10  $\mu$ m.

During IMI experiments was explored the activation and response of Nrf2 to IMI. Our analyses indicated that IMI induced a stable and large expression of this transcription factor in a time dependent manner with the highest induction after 30 min of IMI exposure (p<0.05) (Fig. 38B). Experiments performed by immunofluorescence, suggested that IMI exposure increased the signal of the transcription Nrf2 factor, proposing the activation of potential rescue secondary responses in F11 cells confirmed by the increase in Nrf2 immunoreactivity (Fig. 38A).

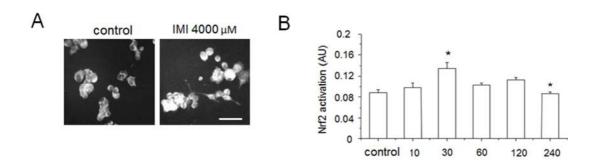


Figure 38: A. Representative example of F11 cells microscopy images analysed in control and after IMI incubation at 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) for 30 min. Cells are immunolabelled with anti-Nrf2 antibodies. Scale bar: 100  $\mu$ m B. The histogram describes the time course (10, 30, 60, 120 and 240 min) of the relative average quantification of Nrf2 immunoreactivity in control and in IMI-exposed cells (4000  $\mu$ M; 1022.4 mg  $L^{-1}$ ). Histograms quantify the grey values of immunopositive signals after background value subtraction. Data have been normalised on total cell number (counterstained with DAPI) and expressed as arbitrary units (AU) (n= 3), p < 0.05 (\*).

# 3.1.3. Imidacloprid effects on mitochondrial function and oxidative stress parameters

To further study IMI cytotoxicity, we proceeded with testing of oxidative stress pathways activation. For this purpose, we first performed mitochondria morphological imaging and then we quantified CAT enzymatic activity as well as LP levels to test on-going oxidative stress and possible oxidative damage.

In vivo time course imaging of living cells loaded with JC-1 probe (200  $\mu$ M; 15 min) and acutely exposed to IMI 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>) for 15 and 60 min was performed to quantify IMI-induced mitochondrial stress (Fig. 39A). Qualitative analysis of JC-1 staining indicated a decrease in MMP after IMI treatment. JC-1 dissociate into monomers that emit a green fluorescence pattern as MMP lowers in presence of IMI (4000  $\mu$ M). This decrease was already significant after 15 min from IMI application with a consequent loss of mitochondrial membrane function (0.7  $\pm$  0.05 fold decrease respect to control; p<0.05) (Fig. 39A).

To measure the effects on F11 cells of possible ROS intermediates produced after IMI exposure at 1000  $\mu$ M (255.6 mg L<sup>-1</sup>) and 4000  $\mu$ M for 24 and 48 h, we performed CAT enzymatic assays that revealed an increasing trend of CAT activity in IMI-exposed cells, although not significant (p>0.05) (Fig. 39B). TBARS assay performed after IMI exposure at 1000  $\mu$ M and 4000  $\mu$ M for 24 and 48 h demonstrated the activation of membrane LP (Fig. 39C) and showed a significant increase in the detection of intermediates of LP in F11 cell medium with respect to control samples. LP values were significantly increased already after 24 h exposure to 1000  $\mu$ M (6.03  $\pm$  1.2 % fold increase respect to control; p<0.01) and to 4000  $\mu$ M IMI (12.9  $\pm$  4.6 % fold increase respect to control; p<0.05). In general, during 24 or 48 h of IMI application, CAT activity in F11 cells showed still a dose-dependent larger increased activation, however not significant (n =3, p>0.05; Fig. 39B), while significantly higher levels of LP products (TBARS) were found released in cell growth medium exposed to IMI with respect to control (n=6, p<0.05; Fig. 39C).

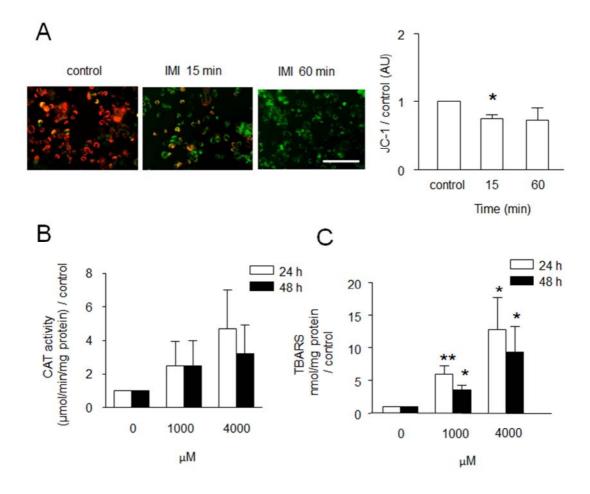
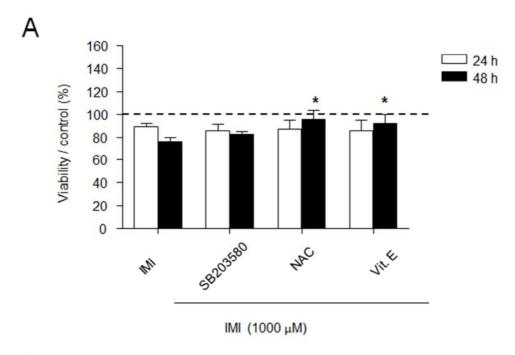


Figure 39: A. In vivo imaging of live cells exposed to IMI 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) 15 and 60 min. Representative microscopy images show JC-1 fluorescence of control (red) and IMI-treated F11 cells (green). Scale bar 100  $\mu$ m. Histogram represent JC-1 green-red fluorescence signal ratio already significant after 15 min from IMI application. Fluorescence values were expressed as the ratio of fluorescence intensity of J-aggregates (red) to fluorescence intensity of monomers (green); ratio results are presented as fold increase/decrease of the control values (JC-1/control in arbitrary units – AUs) (n=3). B. Catalase activity in F11 cells tested 24 or 48 h after IMI application - 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) (n =3). C. Quantification of TBARS products released in cell growth medium by cell exposed to IMI 1000  $\mu$ M and 4000  $\mu$ M for 24 and 48 h (n=3). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

### 3.1.4. Effects of different tested inhibitors on imidacloprid-mediated cytotoxicity

The role of oxidative stress and p38 induction in IMI cytotoxicity was further confirmed by additional cell survival measurements (Fig. 40A and 40B) performed on cells exposed to 1000  $\mu$ M (255.6 mg L<sup>-1</sup>) and 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>) IMI for 24 and 48 h cultivated in the presence of N-acetyl cysteine (NAC; 1 mM, 30 min preapplication), mixed tocopherols (1 mM, 30 min pre-application) or with the p38 inhibitor SB203580 (10 µM, 30 min pre-application). These inhibitors were tested separately to block possible IMI effects or selected molecular pathways and was noticed that their presence significantly reduced the cell loss. With previous data was demonstrated that MAPK signal transduction pathways as well as IMI-induced oxidative stress and altered oxidative stress parameters are possibly associated to IMI cytotoxic effects. To additionally prove the necessary and sufficient role of these mechanisms, a series of MTT cell viability tests were performed on IMI-exposed cells in the presence of pharmacological inhibitors like p38 MAPK inhibitor SB203580, the broad-range anti-oxidant NAC or LP inhibitor mixture of tocopherols known as vitamin E. These compounds were co-applied with IMI to test their efficacy in protection from IMI endpoint toxic effects after 24 or 48 h incubation (Fig. 40A and 40B).

Oxidative stress inhibitors NAC and vitamin E were sufficient to protect from 1000  $\mu$ M and 4000  $\mu$ M (48 h) IMI exposure (p<0.05 and p<0.01 compared to IMI exposed cells at same time and concentration) (Fig. 40A and 40B). Vitamin E resulted highly effective after 24 h in presence of the highest concentration of IMI (4000  $\mu$ M) and viable cells reached values of 128  $\pm$  17.6 % (p<0.05 compared to IMI exposed cells). A significant protection from 4000  $\mu$ M IMI toxicity after 24 (p<0.05) and 48 h (p<0.01) was found also blocking p38 pathways with inhibitor SB203580 (Fig. 40B). Generally, these experiments demonstrated a complex dose- and time-dependent scenario, which demonstrated that oxidative stress inhibitors seem sufficient in some cases to significantly improve the F11 cell survival and to lower the toxic impact of IMI.



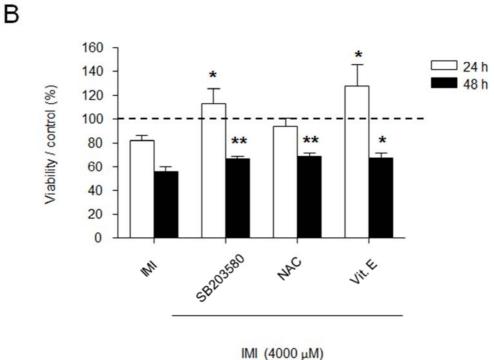


Figure 40: Values of MTT assay in F11 cells after 24 and 48 h exposure to 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) (A) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI (B) in presence of p38 MAPK inhibitor SB203580, N-acetyl-cysteine (NAC) or mixed tocopherols (vitamin E); data presented as percentage of control shown as dashed line (n=4). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*) compared to IMI-exposed cells.

Toxicity data relative to F11 cells and measured parameters are summarised in Table 12. The lowest concentration of tested chemicals that caused significantly different cell responses from that in the control group was considered as LOEC value of tested compound, while the highest tested concentration at which no significant differences from control were found was the NOEC point. The actual safe or no effect concentration lies between the obtained NOEC and LOEC values for specific biomarker. All obtained data are sorted based on type of response, tested chemical and time of exposure/measurement. Toxicity differed considerably among the tested compounds and among test methods and endpoints. The NOEC and LOEC values for the different chemicals and endpoints ranged from 31.5 to >1022.4 mg L<sup>-1</sup> (Table 12).

In summary, based on the survival endpoint, the IMI's TP 6CNA clearly results as the most biologically active towards neuronal cells. The toxicity trend (according to cell survival after 48 h) for all tested compound was as follows: 6CNA > Confidor 200SL > IMI > desnitro-IMI > olefin-IMI > 5-hydroxy-IMI. In this toxicity trend is important to notice that desnitro-IMI after 48 h presented significantly higher adverse effect on cell survival compared to IMI (as pure compound or commercial formulation). It reduced the neuronal cell viability of more than 3.3 fold compared to IMI effect alone.

Activation of phosphorylated ERK resulted as the most sensitive endpoint with an early response at IMI concentrations of 400  $\mu$ M (102.2 mg L<sup>-1</sup>). Cellular CAT activity was not induced after IMI exposure, but showed an increasing trend. However, elevated cellular ROS production caused consequent peroxidative damage of lipid membrane which was confirmed with elevated values of LP. In general, toxicity tests performed on neuronal cells demonstrated an overall cellular dynamics related to induced oxidative stress with activated Nrf2 factor and MAPK kinases as well as elevated LP.

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Table 12: NOEC and LOEC values obtained for applied in vitro models and measured endpoints.

	Chemical  IMI  (mg L <sup>-1</sup> )											
Endpoint			Confidor 200SL (a. i. IMI mg L <sup>-1</sup> )		<b>6CNA</b> (mg L <sup>-1</sup> )		desnitro-IMI (mg L <sup>-1</sup> )		<b>olefin-IMI</b> ( mg L <sup>-1</sup> )		5-hydroxy-IMI (mg L <sup>-1</sup> )	
VII VIII O	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC
IN VITRO (F11 cells)												
Cell survival (24 h)	25.6	51.1	255.6	1022.4	630	>630	247.1	988.5	253.6	1014.1	271.6	1086.4
Cell survival (48 h)	127.8	255.6	51.1	127.8	31.5	78.7	123.6	247.1	253.6	1014.4	271.6	1086.4
ERK induction (30 min)	<102.2	102.2	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
p38 induction (30 min)	102.2	1022.4	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Nrf2 induction (24 h)	<511.2	511.2	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
MMP alterations (15 min)	<1022.4	1022.4	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
CAT (24 h)	1022.4	>1022.4	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
CAT (48 h)	1022.4	>1022.4	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
LP (24 h)	<255.6	255.6	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
LP (48 h)	<255.6	255.6	n. t.	n. t.	n. t.	n. t.	n. t.	<i>n. t.</i>	n. t.	n. t.	n. t.	n. t.

Abbreviations: n. t. not tested

Table 12: NOEC and LOEC values obtained for applied in vitro models and measured endpoint. (continued)

	Chemica	l										
Endpoint	$\frac{\mathbf{IMI}}{(mg\ L^{-1})}$		<b>Confidor 200SL</b> (a. i. IMI mg L <sup>-1</sup> )		$\begin{array}{c} \mathbf{6CNA} \\ (mg\ L^{-1}) \end{array}$		desnitro-IMI (mg L <sup>-1</sup> )		olefin-IMI ( mg L <sup>-1</sup> )		5-hydroxy-IMI (mg L <sup>-1</sup> )	
	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC
IN VITRO (S. typhimurium)												
Base pair substitution (48 h)	255.6	>255.6	255.6	>255.6	157.5	>157.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Frame shift mutation (48 h)	255.6	>255.6	255.6	>255.6	157.5	>157.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.

Abbreviations: n. t. not tested

### 3.1.5. HPLC and LC-MS analysis of imidacloprid presence in F11 cells

In order to test IMI toxicity on a mammalian cellular model, we tested concurrently the presence of IMI and its TPs in the cell culture medium through HPLC-DAD analyses, while through LC-MS analyses was tested the presence of these compounds inside cells and its capability to enter cellular compartments. These analyses were performed only on the highest concentration of 4000 µM; 1022.4 mg L<sup>-1</sup> that induced some significant effects on cell survival. High concentration of tested compounds was also selected to ensure possible detection of TPs formed during the incubation period of 48 h. With higher amounts of IMI (up to 1022.4 mg L<sup>-1</sup>) if the TPs are formed is possible to have them at a detectable/measurable level, otherwise the concentrations were be too low even for highly sensitive methods.

In Fig. 41A are shown representative absorbance traces of HPLC-DAD analysis of cell culture medium containing IMI (4000  $\mu$ M; 1022.4 mg L<sup>-1</sup>) as fresh solution (0 h) without cells (blue line), solution of IMI incubated for 48 h without cells (black line) and solution of IMI incubated for 48 h with cells (red line). Incubation of IMI for 48 h at 37 °C reduced the peak signal at  $55.7 \pm 3.2$  % with cells and at  $67.8 \pm 1.2$  % without cells (n=3) compared to control (fresh and not incubated solution of IMI) expressed as C/C<sub>0</sub> [%]. The actual concentration reduction expressed in mg L<sup>-1</sup> was: for fresh IMI at 0 h (983.4  $\pm$  mg L<sup>-1</sup>), for incubated medium (48 h) containing IMI without cells (685.2  $\pm$  7.7 mg L<sup>-1</sup>) and with cells (561.8  $\pm$  14.4 mg L<sup>-1</sup>). These data demonstrated a significant degradation of IMI in absence of cells (after 48 h in dark at 37 °C) and even a stronger IMI disappearance through possible biotransformation processes in presence of cells under the same conditions (p<0.05). Due to the applied high concentrations of IMI it was expected the formation and detection of possible TPs. However, at the moment through LC-MS analyses this metabolites released in culture medium were not detected.

In addition, cell lysate were also qualitatively analysed for the presence of IMI and its potential TPs (i.e. metabolites) with LC-MS. Cell lysate chromatograms showed the presence of IMI in the cell lysate (red line), which was confirmed with the extracted ions for the peak at 6.8 min (Fig. 41B and 41C). Also in this case, no emerging of additional peaks was present that could suggest possible biological

transformation of IMI in presence of mammalian cells within 48 h incubation period and these issues are still under investigation.

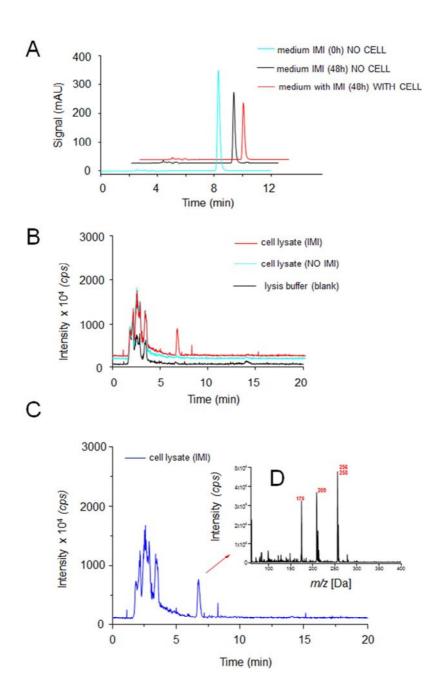


Figure 41: A. Representative peaks of HPLC analysis of cell culture medium samples containing IMI (4000  $\mu$ M, 1022.4 mg  $L^{-1}$ ). Fresh IMI (0 h) without cells (blue line), incubated IMI (48 h) without cells (black line) and incubated IMI (48 h) with cells (red line). B. LC-MS total ion chromatograms of cell lysate exposed to IMI (4000  $\mu$ M; 1022.4 mg  $L^{-1}$ ) for 48 h (red line) overimposed to total ion chromatograms of cell lysate in control group without IMI (blue line) and lysis buffer alone as blank (black line). C. Total ion chromatogram of cell lysate treated with IMI with extracted ions for peaks at 6.8 min (1D) (n=3).

# 3.2. In vitro model: Salmonella typhimurium

In the *S. typhimurium* reverse mutation assay with metabolic activation IMI tested as pure compound, as Confidor 200SL and its TP 6CNA did not cause any mutations. The determination of mutagenicity induced by IMI and its TP was performed using two bacterial strains (TA98 and TA100) and addition of metabolic S9 fraction in Ames test (ISO 16240, 2005).

The results of the mutagenicity conducted with S9 fraction in *Salmonella* tested strains are presented in Table 13. Aminofluorene (2-AF) and methyl methanesulfonate (MMS) were used as the positive control for mutagen action. The maximum tested doses for each compound was chosen based on non-toxic effect of this dose on bacterial growth. Results revealed that all three tested compounds produced no mutagenicity (at least > or = to two-fold background - control only) colonies with metabolic activation.

It is important to mention that only after Confidor 200SL exposure a minor slight increase of the number of TA100 revertants was noticed, which indicated possible potential for mutagenic impact. The numbers of spontaneous revertants of TA100 is ~100 and for Confidor 200SL were present values of 174 and 183 (but only in two replicates) and were not statistically significant. In summary, no specific mutagenic potential or effect was noticed at the highest tested concentration (Table 12 and 13). These data are in concomitance with previous studies with IMI that did not show any mutations, with or without metabolic activation, in the Ames test (JMPR, 2001).

Table 13: Mutagenicity test responses after treatment with IMI, Confidor 200SL and 6CNA with addition of S9 metabolic fraction in Ames test (n=6).

Chemicals	<u>His<sup>+</sup> revertant/plate<sup>a</sup></u> Doses – μg/plate (μM)		Number of revertants TA98	Number of revertants TA100
Positive control (2AF; MMS)	20 μg; 2μL/plate		>2000	>2000
Control (no chemicals)			$19.2 \pm 1.4$	122 ± 2
IMIDACLOPRID	0.053	(30)	22.5 ± 1.5	$128.5 \pm 22.5$
	0.178	(100)	25 ± 6	$139.5 \pm 12.5$
	1.78	(1000)	$27.5 \pm 0.5$	$132 \pm 7$
CONFIDOR 200SL	0.265	(30)	21 ± 1	$150 \pm 3.5$
	0.89	(100)	$26.5 \pm 3.5$	$158 \pm 16$
	8.9	(1000)	$19.5 \pm 2.5$	161.5 ± 21
6CNA	0.033	(30)	$26.5 \pm 3.5$	$122.5 \pm 6.5$
	0.11	(100)	18 ± 1	$131.5 \pm 4.5$
	1.11	(1000)	$20.5 \pm 0.5$	102 ± 1

<sup>&</sup>lt;sup>a</sup> Data represent the mean revertants  $\pm$  standard error of data from two independent experiments, each having three replicates (n = 6). The numbers of revertants that are > or = two-fold above that of negative control (without tested chemicals addition) are regarded as the positive response.

#### 3.3. *In vivo* models

In the following sections are presented the results relative to comparative toxicity testing of IMI, Confidor 200SL and four IMI's TP performed on different *in vivo* laboratory bioassay using: *V. fischeri*, *D. subspicatus* and *G. fossarum*.

## 3.3.1. Toxicity testing with marine luminescent bacteria Vibrio fischeri

Toxicity testing of aqueous solutions for three IMI metabolites (5-hydroxy-IMI, desnitro-IMI and olefin-IMI) was performed and the results revealed the highest toxicity in case of olefin-IMI. The toxicity trend for these three compounds was olefin IMI > desnitro IMI > 5-hydroxy IMI. The results are presented in Table 14.

Table 14: Inhibition V. fischeri bacterial luminescence after 30 min exposure to 5-hydroxy-IMI, desnitro-IMI and olefin-IMI aqueous solutions.

Concentration chemical	Inhibition of luminescence %								
$[mg\ L^{-1}]$	olefin-IMI	olefin-IMI desnitro-IMI 5-hydroxy-IM							
0.39	$2.25 \pm 2.66$	$4.27 \pm 0.58$	$6.00 \pm 0.98$						
0.78	$2.13 \pm 0.36$	$9.74 \pm 0.58$	$7.13 \pm 0.37$						
1.56	$6.74 \pm 0.18$	$4.81 \pm 0.25$	$6.53 \pm 1.1$						
3.12	$7.06 \pm 2.05$	$4.96 \pm 0.58$	$8.71 \pm 0.65$						
6.25	$10.99 \pm 2.53$	$5.76 \pm 1$	$8.88 \pm 0.15$						
12.5	$12.5 \pm 2.73$	$9.38 \pm 0.05$	$10.13 \pm 0.31$						
25	$20.39 \pm 0.67$	$14.99 \pm 0.41$	$10.73 \pm 0.06$						
50	$27.74 \pm 1.58$	$19.79 \pm 0$	$19.54 \pm 0.28$						
100	$44.33 \pm 0.56$	$36.6 \pm 0.68$	$34.54 \pm 0.09$						

On the basis of results listed above, the dose response curves for all three TPs were derived. For all three compounds, the inhibition of luminescence with tested concentrations was too low and in this case only  $EC_{20}$  values were obtained. The 30 min  $EC_{20}$  for olefin-IMI was of 25.08 mg  $L^{-1}$ , for desnitro-IMI of 48.23 mg  $L^{-1}$  and for 5-hydroxy-IMI of 51.04 mg  $L^{-1}$  based on graphical extrapolation from obtained curves (Fig. 42A, B and C). Also, based on obtained  $\Gamma$  values for EC calculation (presented in Annexes), additional graphs with gamma values and its linear regression are shown in Fig. 42a and comparable 30 min  $EC_{20}$  values were obtained

(EC<sub>20</sub> for olefin-IMI was of 22.57 mg  $L^{-1}$ , for desnitro-IMI of 41.56 mg  $L^{-1}$  and for 5-hydroxy-IMI of 46.01 mg  $L^{-1}$ ).

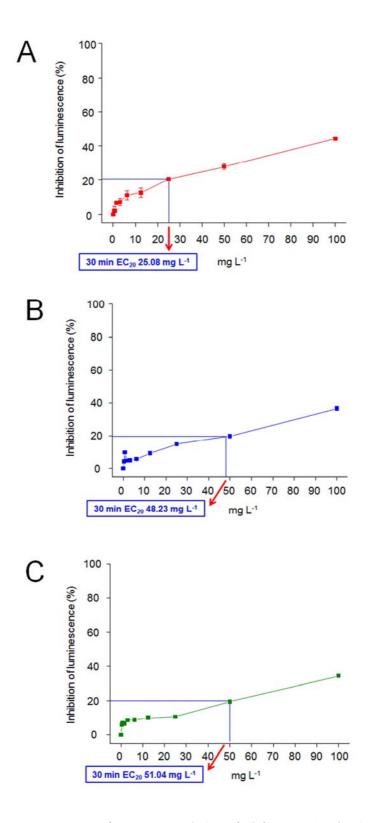


Figure 42: Dose response curve for aqueous solution of olefin-IMI (A), desnitro-IMI (B) and 5-hydroxy-IMI (C) for V. fischeri luminescent bacteria within 30 minutes of exposure (n=2).

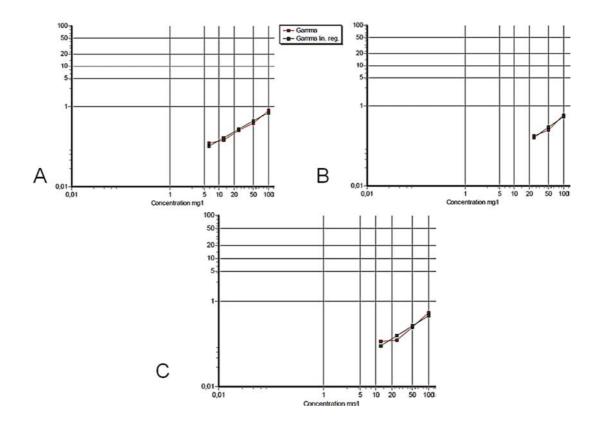


Figure 42a: Gamma values ploted against their corresponding chemical concentration (A) olefin-IMI, (B) desnitro-IMI, (C) 5-hydroxy-IMI.  $EC_{20}$  value is given by the point of intersection with the x-axis at  $\Gamma = 0.25$ .

Data on the toxicity of all tested compounds are presented in Table 15 with corresponding EC or IC values. Data relative to IMI, Confidor 200SL and 6CNA are obtained from previous studies and are listed for better comparison of toxic effects within all tested compounds. Based on all EC<sub>50</sub> and EC<sub>20</sub> data, 6CNA resulted as the most biologically active and toxic for *V. fischeri* followed by Confidor 200SL and IMI. From three additionally tested IMI's TP, olefin-IMI toxicity could deserve some further consideration and testing.

 $11^{2}$ 

Table 15: EC and IC values obtained for applied in vivo models and measured endpoints.

Endpoint	Chemical  IMI (mg L-1)	Confidor 200SL (a. i. IMI mg L <sup>-1</sup> )	<b>6CNA</b> (mg L <sup>-1</sup> )	desnitro-IMI (mg L <sup>-1</sup> )	olefin-IMI (mg L <sup>-1</sup> )	5-hydroxy-IMI (mg L <sup>-1</sup> )	
IN VIVO (V. fischeri)  Luminescence inhibition (30 min)	<61.9 $IC_{50}=61.9^{a}$ $IC_{50}=80^{b}$ $IC_{50}=226^{d}$ (159-322)	<56 IC <sub>50</sub> =56 <sup>a</sup>	<15.1 EC <sub>50</sub> =15.1°	<48.23 EC <sub>20</sub> =48.23	<25.08 EC <sub>20</sub> =25.08	<51.04 EC <sub>20</sub> =51.04	

<sup>&</sup>lt;sup>a</sup>Tišler et al., 2009

<sup>&</sup>lt;sup>b</sup> Dell'Arciprete et al., 2009

<sup>&</sup>lt;sup>c</sup> Žabar, 2012

<sup>&</sup>lt;sup>d</sup> Kungolos et al., 2009

# 3.3.2. Water quality parameters and stability study with green microalgae and crustacean amphipod

Water quality parameters were measured for all water samples from the sampling site and during toxicity tests with *D. subspicatus* and *G. fossarum*. No significant changes were observed through whole experimental trial (n=10). Mean values were as follows: pH 7.9  $\pm$  0.1, temperature 14.7  $\pm$  0.3 °C and water conductivity of 378.3  $\pm$  21.7  $\mu$ S cm<sup>-1</sup>. The water had average oxygen concentration of 9.8  $\pm$  0.2 mg L<sup>-1</sup> and saturation of 95.8  $\pm$  2.3 %. Mean values of TOC and TN at the water source location were 8.7  $\pm$  0.1 and 0 mg L<sup>-1</sup>, respectively. Moreover, dissolved oxygen concentration during whole experimental trial was between 70 % and 80 % of saturation. These were all acceptable conditions for toxicity test (EPA, 1996).

Our experiments showed no significant changes in concentration of IMI and 6CNA in test solutions during 24 h (amphipods) and 96 h (algae) exposure (Table 16). The actual exposure concentrations of both chemicals did not differ by more than  $3.4 \pm 0.3$  % (for concentrations in tests with amphipods) and by  $15.8 \pm 0.4$  % (for concentrations in tests with algae) from the initial concentrations. IMI and 6CNA concentrations were consistent over time in all tests (Table 16). Therefore the results are given in nominal concentrations, as suggested by ISO 10706 (2000).

#### 3.3.3. Algae toxicity test

Algal chronic toxicity revealed a high toxic potential of 6CNA at the highest concentration (Fig. 43C). 6CNA induced some perceivable alterations in algae growth, causing slight and temporary inhibition effects at lower doses (4.7 and 15.7 mg  $L^{-1}$ ) already after 24 h compared to control (p<0.05) (Fig. 43C). The highest dose of 6CNA extensively suppressed the algal growth. 6CNA induced acidification of the algal medium (pH up to 5.5  $\pm$  0.1; n=3). In all other groups, pH did not deviate significantly from the initial values as in the case of 6CNA at the highest dose. Overall 6CNA effects were stimulatory on algae growth. Major stimulatory effect of 6CNA was observed at 31.5 mg  $L^{-1}$  (48 h) reaching 176.4  $\pm$  3.4 % and stayed significantly increased also after 72 h compared to control (p < 0.001) (Fig. 43C). It

was not possible to calculate the  $IC_{50}$  value for IMI due to its low inhibitory effects within the entire range of tested concentrations (Fig. 43A). Furthermore, the toxicity of Confidor 200SL ranged from 27.9 % up to 49.72 % (Fig. 43B). Inhibition of algal growth was significant at 127.8 and 255.6 mg L<sup>-1</sup> compared to control (p<0.01). Higher toxicity of Confidor 200SL was possibly induced by known co-formulants present in the commercial formulation which contributed as a major part to toxicity for algae. These co-formulants (DMSO and NMP) alone induced a significant inhibition of 82.3 % and 89.7 % (at 0.06 and 0.12 %; v/v) compared to control (p<0.001) (Fig. 43B).

Table 16: Mean  $\pm$  standard error detected concentrations expressed as  $\mu g L^{-1}$  and  $mg L^{-1}$  of IMI and 6CNA in aqueous samples for the 24 h G. fossarum and 96 h D. subspicatus static toxicity tests (n=3).

Nominal concentration (µg L <sup>-1</sup> )	G. foss Dark Т [µg]	=22°C	Nominal concentration (mg L <sup>-1</sup> )	D. subs Light T [mg	
<u>IMI</u>	0 h	24 h	<u>IMI</u>	0 h	96 h
102.2	$105.5 \pm 2.5$	$99.7 \pm 0.7$	7.6	$7.5 \pm 0.1$	$6.4 \pm 0.1$
153.3	$154.7\pm0.7$	$148.5 \pm 1.4$	25.6	$26.3 \pm 0.5$	$21.9 \pm 0.5$
204.5	$203.9 \pm 1.8$	$198.1 \pm 0.5$	51.1	$51.4 \pm 1.2$	44 ±1.4
255.6	254.2± 1. 6	$250.8 \pm 0.4$	127.8	$127.4\pm0.7$	103.6 ±2.2
511.3	$511.7 \pm 0.18$	$481.2 \pm 0.6$	255.6	$255.1 \pm 0.8$	240.4 ±2.9
6CNA	0 h	24 h	6CNA	0 h	96 h
62.8	$62.4 \pm 0.5$	$63.3 \pm 0.9$	4.7	$4.5 \pm 0.1$	$4.1 \pm 0.1$
94.6	$93.5 \pm 0.8$	$92 \pm 0.8$	15.7	$14.8 \pm 0.5$	$14\pm0.2$
126.2	$127.3 \pm 0.4$	$120 \pm 0.9$	31.5	$29.9 \pm 0.9$	$28.7 \pm 0.6$
157.7	$157.4 \pm 0.9$	$152.6 \pm 1.1$	78.7	$77.1 \pm 1.3$	$71.1 \pm 0.8$
315.5	$315.7 \pm 0.3$	$310.1 \pm 1.2$	157.5	$156.1 \pm 0.8$	122.3 ±2.6

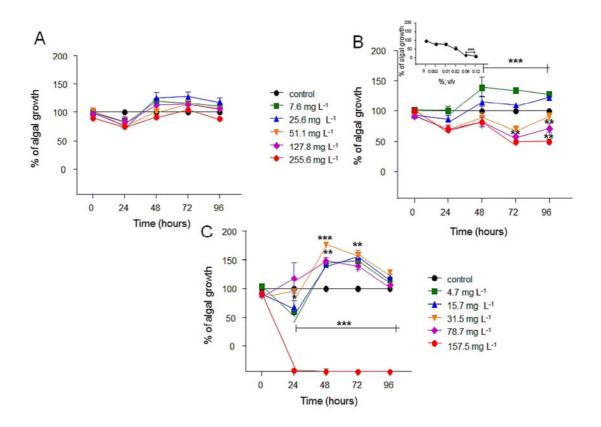


Figure 43: D. subspicatus % of algal growth compared to control after exposure to IMI (A) Confidor 200SL (B) and 6CNA (C) at 24, 48, 72 and 96 h. The inside graph represents exposure to negative control (known co-formulants only). Data are reported as mean  $\pm$  standard error (n = 3). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).

### 3.3.4. Amphipods toxicity test

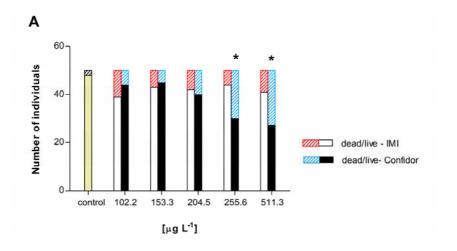
#### 3.3.4.1. Survival rate and behavioural alterations

After 24 h of acute toxicity test, monitored in all groups were: (1) the number of dead amphipods (mortality) and (2) the number of immobile/paralysed or recently moulted amphipods. Only male adult specimens were used for laboratory tests. All specimens presented a mean total body length of  $12.35 \pm 0.25$  mm and mean weight of  $0.029 \pm 0.002$  g. Individuals which sex was not possible to determine were classified as juveniles and not used for this research.

The negative control (mixture of known co-formulants: DMSO and NMP) did not have any adverse effects on *G. fossarum* at all tested concentrations (data present in Annex section). Due to this fact, all values were compared to control (only stream

water). Furthermore, concentrations of all tested compounds lower than 102.2 µg L<sup>-1</sup> for IMI and 62.8 µg L<sup>-1</sup> for 6CNA did not induce significant effects compared to control (data present in Annex section).

Average mortality in control groups was between  $2.2 \pm 1.1$  and  $4.3 \pm 1.9$  % in all bioassays. Our data demonstrated slight toxicity of IMI with minor changes in mortality rate (Fig. 44A). IMI induced only  $22.3 \pm 5.09$  % of dead organisms at  $102.2 \ \mu g \ L^{-1}$ . Commercial formulation Confidor 200SL demonstrated an increased effect on mortality, especially at higher concentrations. Percentages of dead organisms at 255.6 and 511.3  $\mu g \ L^{-1}$  of a. i. reached  $40 \pm 5.7$  % and  $45.5 \pm 7.3$  %, respectively (Fig. 44A). This increased mortality was significant for the both concentrations (p<0.05). On the contrary, 6CNA showed an overall low toxicity, ranging from  $8.6 \pm 1.9$  % up to  $14.1 \pm 1.1$  % (at 62.8 and  $315.5 \ \mu g \ L^{-1}$ , respectively; Fig. 44B).



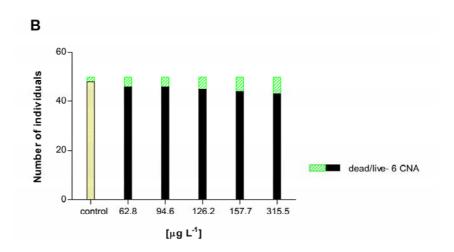


Figure 44: Mortality rate of G. fossarum after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). (n=50). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

At 511.3  $\mu$ g L<sup>-1</sup> of IMI and Confidor 200SL was present a high number of inactive animals with only respiration movements. These values were of 76.6  $\pm$  6.6 % for IMI and of 90  $\pm$  5.7 % for Confidor 200SL (p<0.001; compared to control) (Table 17). It is also interesting to report the number of animals that underwent moult (leaving the entire old *exuvia*) after treatment with tested compounds, apparently stimulatory effect on moult processes was due to the action of transformation product 6CNA. Number of moulted amphipods after 24 h exposure to 6CNA at 315.5  $\mu$ g L<sup>-1</sup> was of 56.6  $\pm$  3.3 % (p<0.001) (Table 17). Number of moulted animals was minor after 24 h of exposure to IMI and Confidor 200SL at 511.3  $\mu$ g L<sup>-1</sup> (23.3  $\pm$  3.3 % and 13.3  $\pm$  3.3 %, respectively; p>0.05). 6CNA seemed to induce overall hyperactivity and rapid swimming (with numerous sideways and back-and-forth movements)

which affected  $80 \pm 5.7$  % of total treated gammarids at 315.5 µg L<sup>-1</sup>. 6CNA (compared to control; p<0.001). Numbers of counted individuals which presented the described behavioural characteristic are summarised in Table 17. It is important to emphasise that this data need further quantification with technologies that allow a more detailed analyses and recording of behavioural patterns, also including the fact of high inter-individual variability which make difficulties in data interpretation after statistical analyses (Malev *et al.*, 2012).

Table 17: Percentage of immobile/paralysed, hyperactive and moulted individuals of G. fossarum (% of total treated animals) exposed to IMI, Confidor 200SL and 6CNA for 24 h. Data are expressed as mean  $\pm$  standard error (n=30). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).

	%	%	%
Nominal	Immobile/paralysed	Hyperactive	Moulted
concentration	concentration individuals		individuals
$(\mu g L^{-1})$			
<u>IMI</u>			
Control	none		none
102.2	16.6 ± 3.3 (**)	None for all groups	$10 \pm 5.7$
153.3	$16.6 \pm 8.8$		$13.3 \pm 3.3$
204.5	$13.3 \pm 3.3$		$23.3 \pm 8.8$
255.6	43.3 ± 3.3 (***)		26.6 ± 3.3 (**)
511.3	76.6 ± 6.6 (***)		23.3 ± 3.3 (**)
Confidor 200SL			
Control	none		none
102.2	23.3 ± 3.3 (**)	None for all groups	$6.6 \pm 3.3$
153.3	33.3 ± 3.3 (**)		$13.3 \pm 3.3$
204.5	46.6 ± 14.5 (**)		$13.3 \pm 8.8$
255.6	56.6 ± 3.3 (***)		$10 \pm 0$
511.3	90 ± 5.7(***)		13.3± 3.33
6CNA			
Control		none	None
62.8	None for all groups	16.6 ± 3.3 (**)	20 ± 5.7 (**)
94.6		23.3 ± 3.3 (**)	33.3 ± 3.3 (**)
126.2		43.3 ± 3.3 (***)	43.3 ± 12 (**)
157.7		43.3 ± 3.3 (***)	46.6 ± 3.3 (***)
315.5		80 ± 5.7 (***)	56.6 ± 3.3 (***)

#### 3.3.4.2 Effects on enzyme activities and lipid peroxidation

Results of enzyme activities were expressed per protein content, since changes in the protein were not significant as a result of 24 h exposure to all tested compounds.

In this study a possible indirect effect of IMI on AChE activity in neonicotinoid exposed gammarids was tested as a biomarker of the cholinergic system. G. fossarum exposed to IMI displayed no significant changes of AChE activity at all concentrations (data not presented in graph). The AChE values at all exposure concentrations of IMI ranged between  $70.6 \pm 7.8$  and  $78.2 \pm 11.6$ µmol/min/mg proteins (p>0.05; compared to control). CAT activity was not modified after IMI exposure (Fig. 45A). The values ranged between  $22.04 \pm 1.5 \,\mu\text{mol/min/mg}$ protein for control and  $28.4 \pm 8.6 \,\mu\text{mol/min/mg}$  protein at 255.6  $\,\mu\text{g L}^{-1}$ . Commercial formulation induced a moderate change in CAT at 511.3 µg L<sup>-1</sup> a. i. going up to  $48.06 \pm 9.7 \, \mu \text{mol/min/mg}$  protein compared to control (p<0.05). Values of CAT activity in the case of exposure to 6CNA reached 48.9 ± 6.7 µmol/min/mg protein already at 157.7 µg L<sup>-1</sup> (p<0.001) (Fig. 45B). After exposure to Confidor 200SL two different outcomes for GST activity at 255.6 and 511.3 µg L<sup>-1</sup> were evident (Fig. 46A). At 255.6 µg L<sup>-1</sup> was present an observable, but statistically not significant decrease in GST activity (p = 0.053). The values of GST went from control values of  $419.1 \pm 101.8$  nmol/min/mg protein to  $286.8 \pm 92.71$  nmol/min/mg protein at 255.6μg L<sup>-1</sup>. Higher concentration of Confidor 200SL (511.3 μg L<sup>-1</sup> of a. i.) induced an increase of GST activity up to 831.4 ± 117.2 nmol/min/mg protein (p<0.05). IMI and 6CNA exposure provoked no significant changes in GST activity compared to control (p>0.05) (Fig. 46A and 46B, respectively). IMI induced at 102.2 µg L<sup>-1</sup> an increase in LP levels (Fig. 47A). This increase was 2.7-fold higher in contrast to the control group (p<0.01). On contrary, Confidor 200SL induced significant rise of TBARS only at higher dose (255.6  $\mu$ g L<sup>-1</sup> of a. i.; p<0.05). This increase was lower than the significant peak induced by IMI at 102.2 µg L<sup>-1</sup> (Fig. 47A). No significant effect of 6CNA on LP increase was noted after 24 h at all concentrations (Fig. 47B). However, it was detected a significant decrease of LP values at 315.5 µg L<sup>-1</sup> compared to control (p<0.001).

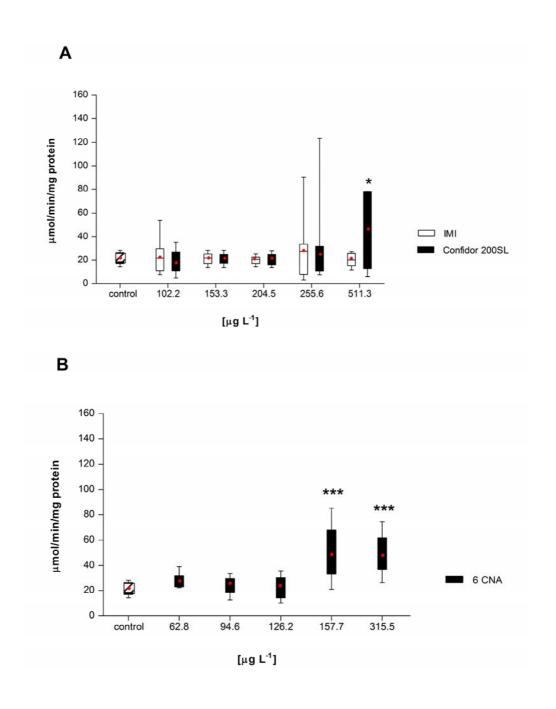


Figure 45: Whole-body CAT activity ( $\mu$ mol/min/mg protein) of G. fossarum measured after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75 % of all readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $^{\bullet}$ ). (n=10). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).

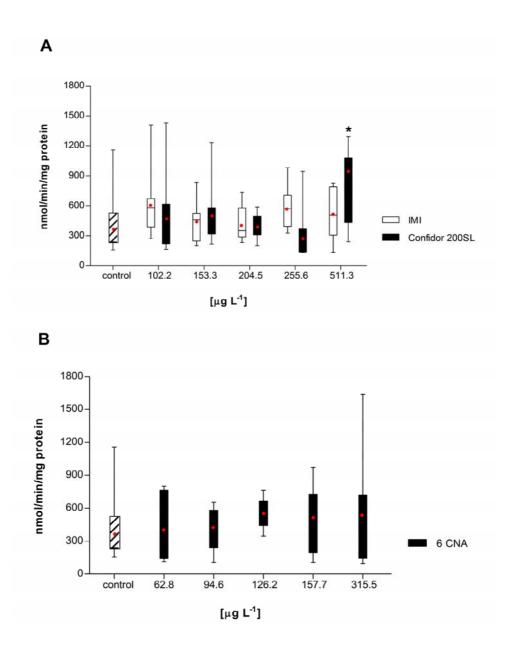


Figure 46: Whole-body GST activity (nmol/min/mg protein) of G. fossarum measured after 24 h of exposure to IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75 % of all readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $^{\bullet}$ ). (n=10). p < 0.001 (\*\*\*), p < 0.01 (\*\*), and p < 0.05 (\*).

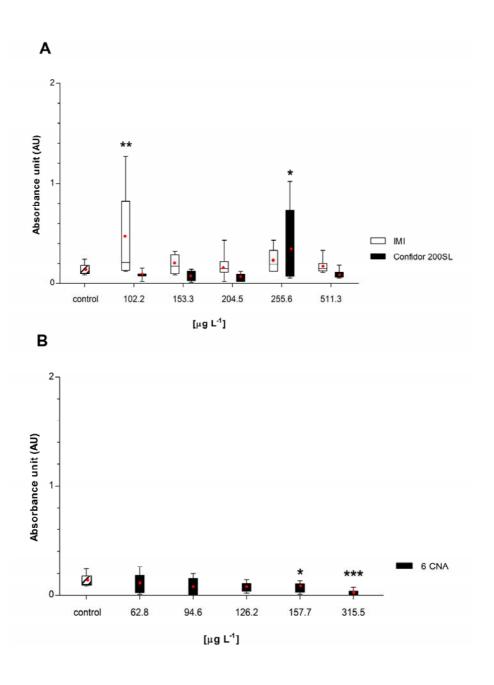


Figure 47: Whole-body LP of G. fossarum (expressed in absorbance units of TBARS products) measured after 24 h of exposure IMI or Confidor 200SL (A) and 6CNA (B). The boxes contain 75 % of all readings, the symbols represent minimum and maximum values ( $^{\perp}$ ) and the mean value ( $^{\bullet}$ ). (n=10). p < 0.001 (\*\*\*), p < 0.01 (\*\*\*), and p < 0.05 (\*).

#### 3.3.4.3 Effects on MXR mechanism

Contaminants in ecosystems can occur as complex mixture, so it is important to consider their possible synergistic, additive or antagonistic action and to understand the interactions between stressor to identify the appropriate application of laboratory studies to the environment (Herbrandson *et al.*, 2003). To test possible synergistic effect was selected copper sulphate (CuSO<sub>4</sub>) which can be found as a solution in most running waters. CuSO<sub>4</sub> is in large amounts used as fungicide in agriculture and could interact with neonicotinoid insecticide (i. e. IMI).

The testing of MXR potential in *G. fossarum* was performed by the accumulation version of the exclusion dye assay (using Rh B as a model MXR substrate). Obtained data are shown in Fig. 48. The animals were exposed to IMI or Confidor 200SL (3 or 30  $\mu$ M; 0.7 or 7.6 mg L<sup>-1</sup>) and 6CNA (3 or 30  $\mu$ M; 0.4 or 4.7 mg L<sup>-1</sup>) and copper sulphate (0.75; 3 and 10  $\mu$ g L<sup>-1</sup> copper a.i.), as pure compounds and mixtures.

In the presence of high concentrations of MXR inhibitors (20  $\mu$ M VER and 10  $\mu$ M CA, respectively) the accumulation of Rh B in the whole body of amphipods increase of 112.3  $\pm$  30.1 % (VER) and 78.4  $\pm$  22.2 % (CA) in comparison to control (100  $\pm$  0.0 %). VER resulted as a better competitive inhibitor, which contends with active sites for the dye and the animals were more sensitive to its presence.

No inhibitory effect of IMI and its commercial formulation was observed vs. control. Interestingly, lower concentration of 6CNA had inhibitory effect on membrane transporters causing accumulation of Rh B ( $205.2 \pm 49.9 \%$ ; p<0.05).

Rh B accumulation in whole body exposed to mixture of IMI and copper (10  $\mu$ g L<sup>-1</sup>) as mixture was of 194.3  $\pm$  82.8 %, while for amphipods exposed to 6CNA and copper (10  $\mu$ g L<sup>-1</sup>) was of 196.4  $\pm$  17.4 % higher in comparison to control (p<0.05). This inhibitory effect was relevant and significant showing dye accumulation and possible synergistic effect of IMI with copper in *G. fossarum*. It is important to note high inhibitory effect of 6CNA alone which exclude possible synergism with copper as in case of IMI. (Fig. 48).

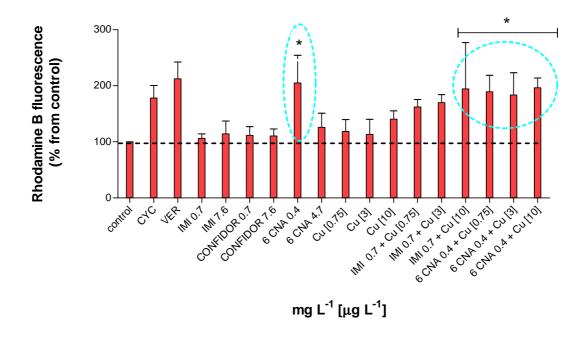


Figure 48: Inhibitory potential of IMI, Confidor 200SL and 6CNA (0.7 or 7.6 mg  $L^{-1}$ ) and copper (3 and 10  $\mu$ g  $L^{-1}$ ) on the accumulation of fluorescent Rh B dye in G. fossarum. Results are expressed in fluorescence units (f.u.) normalised on total animal wet body weight expressed as percentage average of n=3 separated experiments. p < 0.05 (\*).

IMI, Confidor 200SL and 6CNA toxicity to *D. subspicatus* and *G. fossarum* in 24 h (acute) and 96 h (sub-chronic) exposure was also expressed as NOEC and LOEC values based on measured parameters – endpoints (Table 15).

It is clearly shown a different effect mode of action of IMI/Confidor 200SL and 6CNA on freshwater algae growth. IMI tested alone as pure compound showed no significant effects on algae, while Confidor 200SL, due to known co-formulants present in the marketed mixture induced significant adverse and inhibitory effect on algal growth. On contrary, 6CNA had a stimulatory and hormetic effect that induced strong algal proliferation at all concentrations  $\leq$  78.7 mg L<sup>-1</sup>. These specific effects and highly diverse algae responses after exposure to IMI and its TPs confirmed measurement of algae growth as a suitable biomarker for neonicotinoid toxicity testing.

In amphipods was confirmed also a different toxic mechanism of IMI/Confidor 200SL and 6CNA. In specific this effect was noticed on level of physiological biomarker such as behavioural alterations. IMI and Confidor 200SL

induced mostly amphipod immobilisation, while 6CNA was affecting through amphipod hyperactivity and increased moult frequency. Based on high-level (physiological) measured endpoint such as behaviour (24 h) the toxicity trend of tested compound could be presented as: Confidor 200SL< IMI < 6CNA, while based on mortality this trend was slightly modified: IMI < 6CNA < Confidor 200SL.

The NOEC and LOEC values for the different chemicals and endpoints in amphipods ranged from 62.8 to 7600 µg L<sup>-1</sup> of all tested chemicals (Table 15 continued). In general, was shown a complex reaction to the tested chemicals (i.e. 6CNA) with specific behavioural response and induced oxidative stress. In addition, were noticed possible effect on membrane level; peroxidative damage of lipid membrane as well as effect on level of membrane transporters (e.g. MXR endpoint). This latter biomarker results as a useful early signal for fast screening short-term (1 h) acute neonicotinoid exposure at higher concentrations, which needs further considerations and standardisation.

Table 15a: NOEC and LOEC values obtained for in applied vivo models and measured endpoints. (continued)

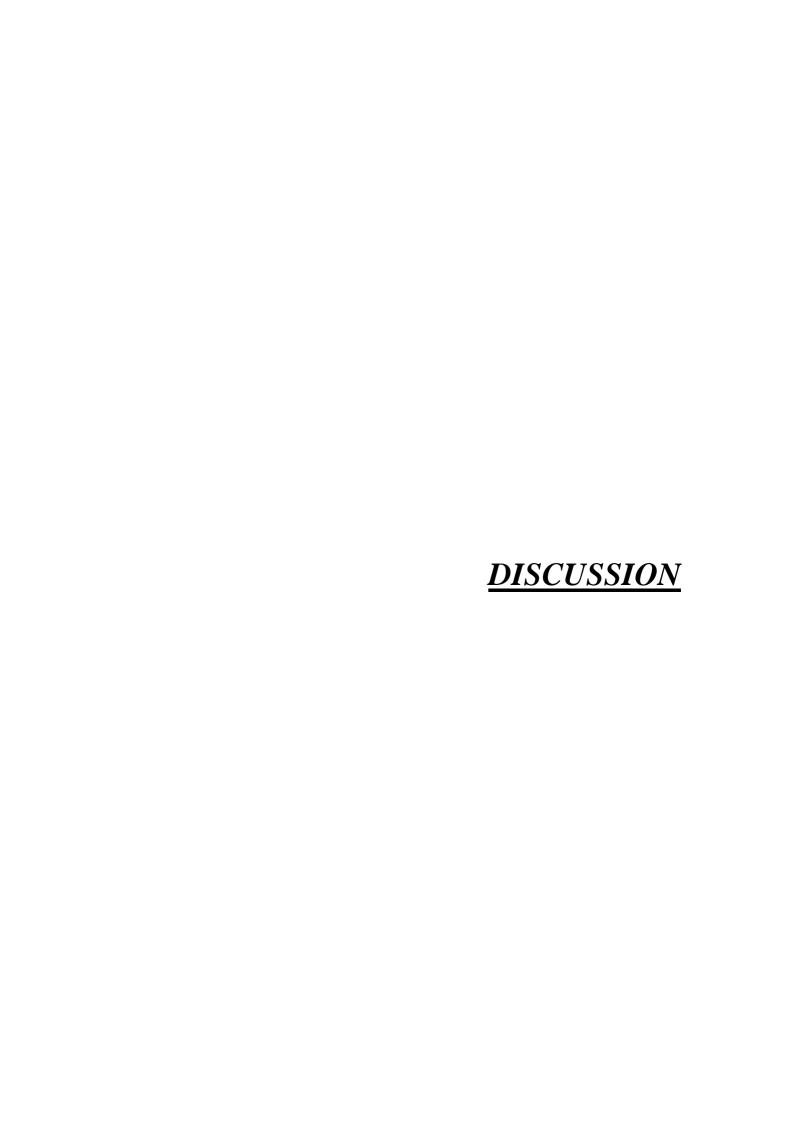
	Chemical											
Endpoint		MI g L <sup>-1</sup> )		or <b>200SL</b> M mg L <sup>-1</sup> )		CNA ng L <sup>-1</sup> )		itro-IMI $ag L^{-1}$ )		<b>iin-IMI</b> <i>ig L</i> <sup>-1</sup> )		roxy-IMI
•	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC
IN VIVO (D. subspicatus)												
Algae growth inhibition (24 h)	255.6	>255.6	<255.6	255.6	<4.7	4.7	n. t.	n. t.	n. t.	n. t.	n. t.	n. t
Algae growth proliferation (24 h)	n. e.	n. e.	n. e.	n. e.	n. e.	n. e.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Algae growth inhibition (48 h)	255.6	>255.6	<25.6	25.6	<157.5	157.5	n. t.	n. t	n. t.	n. t.	n. t.	n. t.
Algae growth proliferation (48 h)	<25.6	25.6	<7.6	7.6	15.7	31.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Algae growth inhibition (72 h)	255.6	>255.6	51.1	127.8	<157.5	157.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Algae growth proliferation (72 h)	<25.6	25.6	<7.6	7.6	15.7	31.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Algae growth inhibition (96 h)	255.6	>255.6	51.1	127.8	<157.5	157.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Algae growth proliferation (96 h)	255.6	>255.6	<7.6	7.6	<31.5	31.5	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.

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Table 15a: NOEC and LOEC values obtained for applied in vivo models and measured endpoints. (continued)

	Chemical											
	IM		Confid	lor 200SL		6CNA		itro-IMI		n-IMI	5-hydi	roxy-IMI
	(in μg			$II$ in $\mu g L^{-1}$ )		in $\mu g L^{-1}$ )		$\mu g L^{-1}$ )		$g L^{-1}$		$ug L^{-1}$ )
Endpoint	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC	NOEC	LOEC
IN VIVO (G. fossarum)												
Mortality (24 h)	511.3	n. e.	204.5	255.6	315.5	n. e.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
Behavioural alterations (24 h)	204.5	102.2 nmobile) 255.6 noulted)	<102.2	102.2 (immobile)	<62.8 <62.8	62.8 (hyperactive) 62.8 (moulted)	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
AChE (24 h)	511.3	n. e.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
CAT (24 h)	511.3	n. e.	255.6	511.3	126.2	157.7	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
GST (24 h)	511.3	n. e.	255.6	511.3	315.5	n. e.	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
LP (24 h)	<102.2	102.2	204.5	255.6	126.2	157.5 (decreased LP)	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.
MXR (1 h)	7.6 mg L <sup>-1</sup>	n. e.	7.6 mg L	-1 n. e.	<0.4 mg	L-1 0.4 mg L-1	n. t.	n. t.	n. t.	n. t.	n. t.	n. t.

Abbreviations: n. e. not evaluated; n. t. not tested



#### 4. Discussion

#### In vitro

The utilisation of *in vitro* test models such as immortalised neuronal cell lines is growing of importance in human and environmental toxicology. All interactions between chemicals and biota occur firstly on the level of cell. Therefore, the use of cell culture in assessment and detection of possible toxic effects of chemical (e.g. pesticides) can be useful (Segner and Braunbeck, 1998). Great amounts of different pesticides, especially insecticides are used worldwide and 5 % of whole population (mainly agricultural workers) are directly exposed to them. According to recent reports, this part of population is calculated to be of 2.6 million persons (David, 2004; Najafi *et al.*, 2010). Chronic exposure to pesticides has been associated in humans to several diseases of nervous system, including neurodegenerative diseases and peripheral neuropathies in adult and childhood (Eskenazi *et al.*, 2008). Prolonged pesticides affecting of neurotransmitter release or receptors operation in human nervous system may lead to synaptic plasticity, long-lasting hypersensitisation or induction/attenuation of molecular mechanisms for neuronal cell survival.

All these effects present a novel area of research which needs further clarifications because there are still a limited number of studies that are referring to IMI in terms of both contaminant and as potential hazard for the additive/cumulative effects on human health. Recently, research in this field is concentrating on IMI and its TPs-mediated toxic effects with biochemical and molecular approaches performed on suitable neuronal cell lines (Table 5 and 6). When dealing with the issue of IMI effects on signalling pathways one should mention the well-known fact that both inorganic and organic chemicals can stimulate ROS production (Viarengo *et al.*, 2002).

IMI and its TPs also elicit acute intracellular responses, particularly in relation to signal integration pathways in mammalian cells. In neuroblastoma cells, low levels of these compounds (i.e. IMI and desnitro-IMI) activate the extracellular-regulated MAPK kinase cascade (in specific ERK) via nAChRs and intracellular calcium mobilisation, leading to possible attenuation of neuronal functions (Tomizawa and Casida, 2002). In our study, F11 cells treated with IMI showed short-

term and transient p38 and ERK MAPK activation, while inhibition of p38 MAPK, blocked and attenuated IMI-induced cell death proving its involvement in oxidative stress pathway. Pesticides are known to induce oxidative stress by generation of free radicals that might cause LP, alternations in membrane fluidity and DNA damage. El-Gendy et al. (2010) observed a marked increase in the activities of the antioxidant enzymes SOD and CAT after IMI treatments in adult mice. Since these enzymes are the first lines of defence against the oxygen free radicals, such increase in the activities has been attributed to the defence mechanisms against oxidative stress in the process of attempted cellular repair. In the same study, IMI treatment to mice indicated a marked increase in the hepatic LP. These results are parallel to the data of Khan and Kour (2007); Yu et al. (2008); Duzguner and Erdogan (2010) and Kapoor et al. (2011) with increased LP values, altered SOD or CAT activity and decrease in GSH content suggesting that induction of oxidative stress is perhaps the central way by which IMI exerts its cytotoxic effects (Khan and Kour, 2007; El-Gendy et al., 2010). Our data demonstrated a moderate toxicity of IMI, however associated with activation of stress oxidative responses. A possible accumulation of intracellular ROS induced by IMI-exposure led to increased levels of LP and slightly improved CAT activity. These results indicate that the increase in oxidative stress has a potential crucial role in the neurotoxicity of IMI in F11 cells.

Oxidative stress is one of the most important pathways leading to neuronal cell death (Lee *et al.*, 2011). This phenomenon is commonly observed when free radicals, such as ROS or RNS overcome cellular antioxidant system and induce cellular damage. ROSs and subsequent oxidative damage lead to disruption of cellular membrane integrity and mitochondrial dysfunctions. In F11 cells, IMI (at higher concentration) affected also mitochondria causing substantial decrease of MMP and consequent loss of mitochondrial activity. In this study was observed that incubation of F11 cell with IMI induced oxidative stress and that IMI-induced cell death was diminished by pre-treatment with antioxidants (i.e. NAC known as vitamin C and vitamin E). NAC can directly and rapidly scavenge free radicals or quench their reactivity and can act by up-regulating endogenous antioxidant defences. It also, fights off several pollutants by stimulating enzymes that detoxify the cell/organism (El-Gendy *et al.*, 2010). Vitamin E is also a naturally occurring lipid soluble antioxidant which protects cells against membrane LP and expression of peroxidation

products including MDA (Jaine *et al.*, 2000). Pre-treatment with antioxidant compounds decreases the generation of ROS preventing the pesticide instability in the antioxidant enzyme activities (Niki *et al.*, 1995; El-Gendy *et al.*, 2010). Our results have revealed that the pre-treatment of neurons with NAC and vitamin E detoxifies IMI-induced neuronal toxicity due to protection of cells against IMI-induced toxic effects (inhibition of mitochondrial activity, LP and ROS action). Presented data are in accordance with those of Verma *et al.* (2007) and El-Gendy *et al.* (2010) who confirmed that pre-treatment with vitamins E and C protected rats from chlorpyrifos-induced oxidative stress and diminished its toxic effects.

Interestingly, the IMI-dependent effects were different in the F11 serum growing line with respect to starved neurons (differentiated), suggesting a different molecular action *via* specific elements in the cell, presumably related to a mature neuronal phenotype. Starved neurons were more susceptible to IMI effect with 50 % of cell loss already at 340 µM (86.7 mg L<sup>-1</sup>) (still under investigation). A detailed functional study of the effect of IMI on F11 sensitisation is yet to be explored and may be achieved by functional studies like calcium imaging or patch clamp on IMI-treated cells. A relevant morphological difference was evident between control cells and those exposed to high dose of IMI. These differences were observed as an important change in the monolayer cell density when compared to control as opposed to distinguished morphological changes and damage, which were also visible. Despite the subjectivity involved identifying morphological alterations, all fluorescence/light microscopy analyses and methods applied in this research were definitely a useful helper in verifying the result of the cytotoxicity test (Holler *et al.*, 2000).

The F11 cellular model used in this study was sensitive to IMI activation and showed consequent signs of stress. The cytotoxicity of IMI appeared to be concentration-dependent in the cell viability test using an MTT assay. IMI resulted as moderately toxic to F11 cells with a cell loss of ~ 50 % only at concentrations higher than 4000  $\mu$ M (1022.4 mg L<sup>-1</sup>) after 48 h of exposure. In case of Confidor 200SL this same effect after 48 h was observed at concentrations lower than 4000  $\mu$ M a. i. IMI. No significant effect on cell survival were observed at concentrations lower than 200  $\mu$ M (51.1. mg L<sup>-1</sup>) for IMI and 500  $\mu$ M (127.8 mg L<sup>-1</sup>) for Confidor 200SL. In a study of Skandrani *et al.* (2006) with differentiated dopaminergic neuronal cell line

SH-SY5Y, 50 % of cell death was reached after 72 h exposure at 2.5-fold lower concentration of IMI (1565 µM; 400 µg mL<sup>-1</sup>) and 5-fold lower concentrations of Confidor 200SL (782 µM; 200 µg mL<sup>-1</sup>). Even if in both studies, cytotoxicity was related to relatively high tested concentration, it is evident that commercial formulations were more aggressive to biological models than the pure compound, when compared to similar concentrations of active principle. Similar observations were reported for marketed mixtures of various pyrethroids and carbamates (Bonatti et al., 1994; Skandrani et al., 2006). Information on inert ingredients is largely treated as business secret, but these chemicals have been shown to exert additive or synergistic toxicity, due to their specific mode of action or by increasing the bioavailability and effectiveness of the a. i. (Beggle et al., 2010). Co-formulants and additives used in pesticide formulation may present themselves as very active, biologically or chemically, and can be also toxic. At the moment more than 500 additives are used as active ingredients (Cox and Surgan, 2006; De Silva et al., 2010). Moreover, it is rare for the population to be exposed to insecticides as pure ingredients and the increase in toxicity of formulation raises more concerns and should be further studied.

It is still unclear if, the IMI evoked cytotoxicity, is a direct or indirect effect. One possibility is that IMI, acting as agonist of nAChRs (Tomizawa and Casida, 2003) causes a persistent neuronal depolarisation and as a consequence it may induce the release of molecules or peptides (ATP, glutamate or CGRP) with associated toxic effects as well as characteristic features of mitochondrial intrinsic apoptotic pathway such as activation of caspase-9 and -3 and nuclear condensation observed by Hoescht (DAPI) staining in our study. However, pre-incubation with a specific antagonist of nAChR α3 subunit which are homogenously highly expressed in F11 cells (mecamylamine; MEC, Arias *et al.* 2010) prevented only partially IMI cytotoxicity, suggesting more complex effects that originate also in a nAChR-independent manner and that is impossible to exclude multiple molecular targets of IMI.

Besides IMI are relevant also IMI's TPs (metabolites) which have been discussed in other sections of this manuscript. Since all of the *in vivo* toxicology studies on IMI involve the generation of TPs (i. e. metabolites), the potential toxicity of the metabolites should be included in the available toxicity data. IMI goes through degradation to a large number of metabolites formed by multiple pathways, both

alternative and sequential. For example, after oral administration of <sup>14</sup>C-methylene labelled IMI in rats, 95 % of the administered dose was absorbed, with an estimated half-life of 35 min. The maximum concentration of <sup>14</sup>C-methylene labelled IMI was reached in blood plasma within 2.5 h after oral or intravenous exposure in rats (Klein and Karl, 1990). Approximately 80% of the administered IMI was excreted in the urine, and 72 % of the urinary secretion was identified as primary metabolite 6CNA and its glycine conjugate (28 % of the total), olefin and 5-hydroxy-IMI (as 30 % of total), and unmodified IMI (15 % of total). In rats exposed orally to <sup>14</sup>C-IMI labelled at the 4- and 5- carbon of the imidazolidine ring (Klein and Brauner, 1991), the following metabolites were detected in the urine 48 h after administration: 5hydroxy-IMI: 13.7 - 14.7 %, olefin-IMI: 7.7-9.1 % and IMI: 6.9 - 14.2 %. Through biotransformation reactions, IMI forms a large number of metabolites (both ionised and non-ionised form) which represent activation and detoxification products relative to action at the nAChRs and possibly more toxic to mammalians than their parent compound (Schulz-Jander et al., 2002; Skandrani et al., 2006). Due to these issues, also IMI TPs were tested during our study for their potential adverse effect on neuronal cell survival. Obtained data relative to TPs cytotoxicity demonstrated significant effects only at highest doses (4000 µM; 1022.4 mg L<sup>-1</sup>) as parent compound, but this effect was stronger and induced overall cell death, especially in presence of desnitro-IMI (as expected). From a toxicological point of view, the metabolite desnitro-IMI is of particular relevance: (1) this metabolite displays a nicotinic receptor mode of action with a evidently higher toxicity to mammals than IMI, (2) it is a major metabolite produced in vitro with human liver microsomes and (3) besides 6CNA, desnitro-IMI is identified as the major degradation product of IMI in the environment (CDPR, 2006).

Furthermore, it was performed also a mutagenic potential screening of IMI, Confidor 200SL and 6CNA with the use of ISO standardised Ames test (ISO 16240, 2005), to provide a full picture of neonicotinoids effects on molecular and cellular mechanisms. Since the introduction of IMI on the insecticide market, its genotoxicity has been studied in different non-target test organisms. Studies showed evidence that exposure to IMI does not increase the micronucleus (MN) frequency in Vicia faba seeds, even though it shows a dose - response for DNA damage induction in earthworms Eisenia fetida (Zang et al., 2000). Increased genotoxicity (MN formation in rats and mutation induction in a bacterial assay) was found in studies which used IMI in combination with an organophosphate (Karabay and Oguz, 2005). Furthermore, one study suggests that IMI and other pesticides may render an organism more susceptible to DNA damage; Shah et al. (1997) found that IMIexposed calf thymus cells had significantly increased amount of damaged DNA than unexposed control cells. On contrary, in vivo studies with orally administrated IMI induced no DNA damage. Negative results were obtained in Chinese hamster bone marrow (Herbold, 1989a, b), in mouse germ cells (Volkner, 1990) and no induction of micronuclei formation in mouse (Herbold, 1988). Also, IMI did not cause any DNA damage in bacterial spores (Watanabe, 1990). In our case, neither IMI/Confidor 200SL nor 6CNA presented any indications of possible mutagenic potential.

Different studies relative to IMI performed on mammalian organism models (i.e. rats and mice) were mainly focused on evaluation of LD<sub>50</sub>, NOEC and LOEC values for conventional endpoints such as: growth, reproduction, survival and behavioural patterns in case of IMI exposure (Table 5 and 6). In general, IMI resulted as moderately toxic to mammals via the oral route of exposure which poses the greatest toxic threat (Mulye, 1996; PMRA, 2001). There are present several studies and a great amount of data on human risk assessment for IMI based on this already established NOEC (NOEL; NOAEL) or LOEC (LOEL; LOAEL) values for conventional endpoints which are summarised in several reports such as: SERA (2006), CDPR (2006) and CCME (2007). Based on these data, U.S. EPA has derived an acute and chronic reference doses (RfD) for IMI of 0.14 and 0.057 mg/kg/day, respectively (US EPA, 2003). Acute and chronic RfDs are well-documented and used directly for all longer term exposures to IMI. Chronic RfD value is based on a NOAEL of 5.7 mg/kg/day in rats (Eiben and Kaliner, 1991; Eiben, 1991) and an uncertainty factor of 100 - two factors of 10 for interspecies and intra-species variability. Acute RfD value is based on a LOAEL of 42 mg/kg in rats (Sheets, 1994) and an uncertainty factor of 300 - a factor of three for extrapolating NOAEL from LOAEL, and two factors of 10 for interspecies and intra-species variability. For workers, the upper range of exposures during the normal broadcast application lead to hazard quotients (HQ) of 1.1, while for general public the highest HQ for nonaccidental exposures is 1.5. HQ for accidental exposures associated with spills into a small body of water result in HQs that range from 1.1 (adult male consuming fish) to 15 (child consuming 1 L of contaminated water) (SERA, 2005).

Few data exist on toxicity threshold of pesticides (as pure ingredient, commercial formulation or TP) in humans or specific cell models. Therefore the present study was designed to implement and provide more information regarding IMI and its TPs toxicity on neuronal cell line through examination of several molecular, biochemical and cellular biomarkers (Fig. 49). In addition were provided data on development of NOEC/LOEC values not merely related to conventional endpoints but also to other measured molecular biomarkers. These *in vitro* results support the possibility of potential toxic effects of IMI and its TPs that may induce chronic sensitisation or long-lasting modulation of sensory neurons and molecular/biochemical alteration. This process is of primary importance in several

chronic disorders (typically involving sensory neurons), like skin hypersensitisation, airways hypersensitisation and chronic pain, which might be relevant for occupational exposure. IMI (as nicotine) may induce neuronal plasticity thus changing the sensitivity to chronic sub-threshold levels of this insecticide. Since high potential impact on human health by chronic exposure to pollutants, new molecular findings on alterations at cellular level after IMI/TPs exposure is environmentally and medically relevant, especially if considered for agricultural occupational exposure and toxicology.

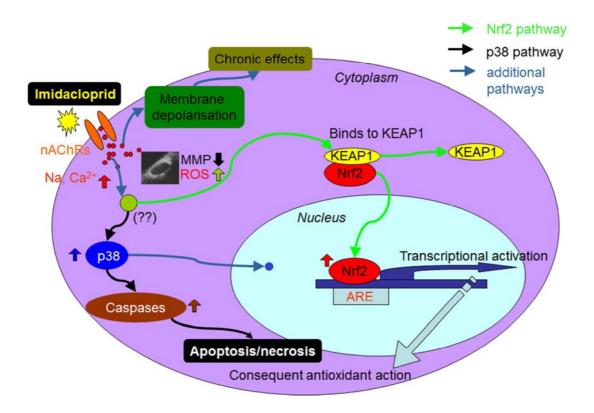


Figure 49: Schematic diagram describing the possible cytotoxic model of IMI and its activated molecular pathways in human neuroblastoma F11 cells. (YNM, 2010 with slight modifications).

#### In vivo

One problematic of the *in vitro* assays is that they only give information about cellular level, so it is important to integrate the results with different *in vivo* organism models. In our case besides mammalian cell model, were used crustacean *G. fossarum*, freshwater algae *D. subspicatus* and the bacteria *V. fischeri*. Different toxicants act differently and not all living organisms are equally sensible. Consequently, single bioassays will never provide an overall picture of the environmental quality and a representative multiple test battery should be used (Bierksen *et al.*, 1998). The use of multi-species laboratory experiments with organisms from different taxonomic and trophic level could provide a possible helpful solution.

Chronic testing was performed on freshwater microalgae D. subspicatus. Generally, it appears that algae are some orders of magnitude less sensitive to IMI than arthropod species and exhibiting no effects of IMI on their growth rate (ISO 8692, 2004; Gagliano and Bowers, 1991). Tišler et al. (2009) determined for D. subspicatus an IC<sub>50</sub> (72 h) for IMI a. i. at 389 mg L<sup>-1</sup> (in comparison highest applied concentration in this study was 255 mg L<sup>-1</sup>). In addition, Kungolos et al. (2009) showed that the inhibitory effect of IMI on Pseudokirchneriella subcapitata growth at a concentration of 1000 mg L<sup>-1</sup> was lower than 50 %. Data presented in this research confirmed the same action of IMI as pure compound causing no significant adverse effects on algal growth (Malev et al., 2012). On the contrary, Confidor 200SL was highly toxic to algae due to the presence of co-formulants which started to inhibit their growth already at 0.003 v/v %. On the other hand, 6CNA induced algal growth and proliferation after a 96 h period of exposure at all concentrations, while at the highest dose (157.5 mg L<sup>-1</sup>) already after 24h induced a significant inhibition and algae death. Presumably the algal growth was inhibited because of the dissociation of the carboxylic group present in 6CNA (Ruiz et al., 2011). This issue induced acidic changes in pH of the algal media and adversely influenced the sensitive microalgae. Algae as primary producers contribute substantially to aquatic habitats and their sensitivity to Confidor 200SL and 6CNA found in this study could cause environmental problems (Malev et al., 2012).

6CNA is a final transformation product formed in environment that does not act as nicotinic agonist but may also contribute to the toxicity effects (Nauen et al., 2001). 6CNA contains the 6-chloropyridinyl moiety and based on its structural/chemical consideration may be of toxicological significance. This transformation product is included in the tolerances established for the IMI residues, although should be considered on its own as other IMI TPs in order to recognise additional IMI-toxicity effects. IMI TPs were already thoroughly presented in the Introduction section, however is important to emphasise again the importance of their formation and presence in environments as they can have possible adverse effects on non-target organisms. Photo-transformation and biotransformation in the aqueous environment appear to be significant transformation routes (Stevens and Halarnkar 1996, as reviewed by Mulye 1997). For example, in a study of Spiteller (1993) was examined radio-labelled IMI degradation in a 30 d laboratory study using water and sediment collected from a pond. By the end of the experiment, 67.6 % of labelled IMI was present in the water (64.0 % as parent IMI and 3.6 % as TPs). In the sediment, 20.4 % was as parent IMI, 0.7 % as TPs, and 8.2 % was as bound residues. On other hand, in soil all processes depend on the soil type and imidazolidin-<sup>14</sup>C labelled IMI presents a maximum mineralisation to CO<sub>2</sub> of 8.8 % or 14 % after incubation for 12 weeks (Anderson 1995, as reviewed in Mulye 1997). Koskinen et al. (2000) found that the bioavailability of IMI decreased in different soils with ageing as indicated by decreases in the amounts of the extracted IMI and increases in the amounts of the bound portion (until 21 d). It is generally noticed that the total <sup>14</sup>C recoveries decrease by time, going from 92.9–96.2, 77.1–88.4 to 53.5–62.4 % of the applied IMI at 7, 14 and 21 d after application, respectively (El-Hamady et al., 2008). Although <sup>14</sup>C-TPs of IMI are rarely detected in soil extracts, the loss of radioactivity must have been due to degradation, strong binding of TPs to the soil matrix and evolution of <sup>14</sup>CO<sub>2</sub>. In soil extracts is present unchanged <sup>14</sup>C-IMI but also <sup>14</sup>C-labelled TPs even if in low percentage/quantity (El-Hamady et al., 2008). Until now, only incomplete data on environmental fate and behaviour of such TPs are available, so there is a strong need to further analyse also intermediate degradation products with respect to their environmental behaviour and potential for non-target organism exposure.

In addition to the tested 6CNA a well-known primary IMI degradation product we performed a screening of three more IMI TPs (desnitro-IMI, 5-hydroxy-IMI and olefin-IMI) with ISO standard test based on *V. fischeri* (ISO 11348, 2007). Based on the collected results was possible to establish olefin-IMI as the most harmful to luminescent bacteria (30 min EC<sub>20</sub> 25.08 mg L<sup>-1</sup>). However, it was not possible to calculate EC<sub>50</sub> values for all three TPs due to very low inhibitory effect on bacterial luminescence for given dilution levels. This issue made also impossible to classify the tested compounds within specific toxicity categories based on the EC<sub>50</sub> values established by the European Commission Directive 93/67/EEC ("very toxic to aquatic organisms" [EC<sub>50</sub>  $\leq$  1 mg L<sup>-1</sup>], "toxic" [EC<sub>50</sub> in the range of 1–10 mg L<sup>-1</sup>], and "harmful" [EC<sub>50</sub> in the range of 10–100 mg L<sup>-1</sup>].

Acute toxicity (24 h) of IMI and 6CNA was evaluated on the freshwater amphipod G. fossarum. After exposure to the highest dose of IMI (511.3  $\mu g L^{-1}$ ) and 6CNA (315.5 µg L<sup>-1</sup>), an overall low mortality was noticed. Most significant effect, as in algae, was observed in case of Confidor 200SL. Increased mortality induced by Confidor 200SL supports the idea that major side effects could be caused by additives such as DMSO and NMP. These known co-formulants mixture alone induced no toxicity in amphipods, while the combined action of IMI and coformulants increased the toxicity of the commercial formulation. In the case of another amphipod crustacean Hyalella azteca, Stoughton et al. (2008) confirmed its higher sensitivity to formulated product than to technical IMI. Also, other tests have shown formulated pesticides to be more toxic to aquatic organisms (Beggel et al., 2010; Chen et al., 2010; Kaczala et al., 2011). These supplementary substances in commercial formulations often represent the highest proportion in pesticide mixtures, so even a minor concern regarding their toxicity and possible synergistic effects with other ingredients should be considered (Tobiassen et al., 2003; Surgan, 2005). Additional studies will be needed to address the potential effect of additives, but such studies are not easily feasible since identity and quantity of other ingredients is most often regarded as confidential information and therefore rarely revealed in easily accessed literature or product labels.

Neonicotinoids are agonist of nAChRs (Tomizawa and Casida, 2003) and do not exert a direct inhibition of the AChE activity as for example organophosphates. In our study we tested possible indirect inhibitory effects on freshwater amphipods

exposed to neonicotinoids. This measurement was also performed on gills of neonicotinoid exposed mussels and showed an interesting outcome with 'U-shape dynamics' of AChE activity (Dondero *et al.*, 2010). In this study Dondero *et al.* observed significant inhibition at the lowest and at the intermediate tested concentration. On contrary, in our case the outcome of IMI effect on AChE activity presented no indirect effect or changes at all exposure concentrations compared to control group (Malev *et al.*, 2012).

It is well-known that pesticides can induce oxidative stress by the generation of ROS, which can induce oxidant-mediated effects (such as increased activities of antioxidant enzymes) and oxidant-mediated toxicities (such as oxidation of lipids) (Zama et al., 2007). Only a few previously published data are available regarding the IMI-induced oxidative stress and these merely relate to mammalian model organisms. These studies showed a slight increase in intracellular ROS and nitric oxide production after IMI exposure (Costa et al., 2009; Duzguner and Erdogan, 2010). A study of Lukančič et al. on G. fossarum demonstrated that IMI influenced not only the respiration but also the electron transport system (ETS) activity (Lukančič et al., 2010). This effect was a consequence of different processes, including oxidative stress. Partial damage to the inner mitochondrial membrane by lipid peroxidation possibly impaired the function of ETS. For better understanding of ROS involvement in the toxicity mechanisms of neonicotinoids, antioxidant enzyme activity, detoxifying GST mechanism and lipid peroxidative damage were monitored in amphipods. In this study, CAT activity after Confidor 200SL and 6CNA exposure at highest doses was significantly increased and indicated action of the protection mechanisms involved in cellular repair processes. El-Gendy et al. (2010) reported a similar increase of CAT after neonicotinoid exposure, but again only in IMI-treated mice. Enhanced GST activity after Confidor 200SL exposure reflects the detoxification processes in treated gammarids and this induction may be due to the glutathione dependent enzyme system that provides major protection against xenobiotic agents. A recent study on the mosquito, Aedes aegypti, demonstrated that exposure to IMI increased glutathione transferase mRNA levels as well as other genes coding for antioxidant proteins (Riaz et al., 2009). In addition was also noticed a slight decrease in antioxidant enzyme GST after exposure to Confidor 200SL (at  $255.6~\mu g~L^{\text{--}1}$  of a. i.). This decrease of the GST activity, although not significant was

evident with 1.5-fold lowered GST activity at 255.6 µg L<sup>-1</sup> of a.i. in Confidor 200SL. This decrease could be interpreted as being overwhelmed by conspicuous ROS production. An additional explanation of enzyme's indirect inhibition is related to their binding with ROS produced also during pesticide metabolism. Metabolism of IMI involves many processes of hydroxylation, i.e. the hydroxylation of the imidazolidine ring at position 4 or 5 leading to the formation of hydroxylated compounds and subsequent loss of important amounts of hydroxyl radicals (Sur and Stork, 2003). Concurrently, with slightly diminished GST activity increased lipid peroxidation levels occurred (at the same exposure concentration of Confidor 200SL). IMI and Confidor 200SL exposure provoked an increase of LP in amphipods. During IMI exposure LP increase occurred at 102.2 µg L<sup>-1</sup> and was represented by a similar-to-hormetic effect. This increase was induced at lower concentrations of IMI and not at higher doses as expected. On the other hand, Confidor 200SL induced an increase of TBARS products, which was highest at  $255.6~\mu g~L^{-1}$  of a. i. Higher TBARS levels at  $255.6~\mu g~L^{-1}$  suggested that exposure to Confidor 200SL resulted in a different time-course of cellular ROS generation or in a possible direct lipid oxidation due to the interactive action of co-formulants and IMI. It is important to notice potentially different toxicity pathways or time-course effects of the parent compound and its transformation product that were observed during this study. After a 24 h exposure 6CNA provoked strong induction of antioxidant enzyme CAT, while its effect was completely absent on the LP, probably due to highly active CAT. On the contrary, Confidor 200SL altered all parameters confirming its higher toxicity compared to active ingredient.

Behaviour is considered as a useful tool in ecotoxicology since is one of the early warning indicators of toxicant stress (Pestana *et al.*, 2009). During experimental pesticide exposure analysed individual biochemical biomarkers should be linked to behavioural responses whenever this is possible (Hellou, 2011). In this study individuals with modified behaviour were counted. During exposure, animals treated with 511.3 µg L<sup>-1</sup> (IMI) exhibited an increase in immobility and inactivity that can be a direct IMI effect on neuro-muscular acetylcholine receptors provoking impairment of locomotion and food filtration, with consequent animal starvation and difficulties in ventilation (Férnandez-Casalderréy *et al.*, 1994). Alternatively, 6CNA at the highest dose induced rapid movements and animal hyperactivity, as well as

disorientation. This disoriented behaviour was also shown in non-target organisms, such as *Apis mellifera*. Honey bees treated with IMI were confused and failed to return to their homing site (Bortolotti *et al.*, 2003). Hyperactivity in swimming may also be linked to an avoidance response towards present chemicals (Roast *et al.*, 2000). Interestingly, short-term 6CNA exposure stimulated amphipods moulting processes. Moulting is an essential physiological process for crustaceans controlled by the neuroendocrine system, on which different toxicants, such as pesticides, can act (Waddy *et al.*, 2002). Moreover, moulted or recently moulted animals could be more susceptible to pesticide action.

This *in vivo* research on freshwater amphipod, confirmed the importance of testing commercial formulations of IMI and IMI's transformation products as they interfere with pure compound safety characteristics. Our present results show that commercial formulation of IMI and its by-product 6CNA exert oxidative stress in freshwater amphipods as well as negative effects on algae growth. The induction of CAT, GST and LP levels demonstrates that exposure of *G. fossarum* to Confidor 200SL leads to peroxidation of membrane lipids and triggers antioxidant and detoxifying cellular mechanisms. Amphipods exposed to 6CNA experienced mainly the activation of catalase scavenging protection mechanism. In general, the major toxic effects were due to the commercial formulation Confidor 200SL both in case of algae and amphipods. This issue is relevant as these marketed mixtures are the one applied directly in the environment and should be further monitored.

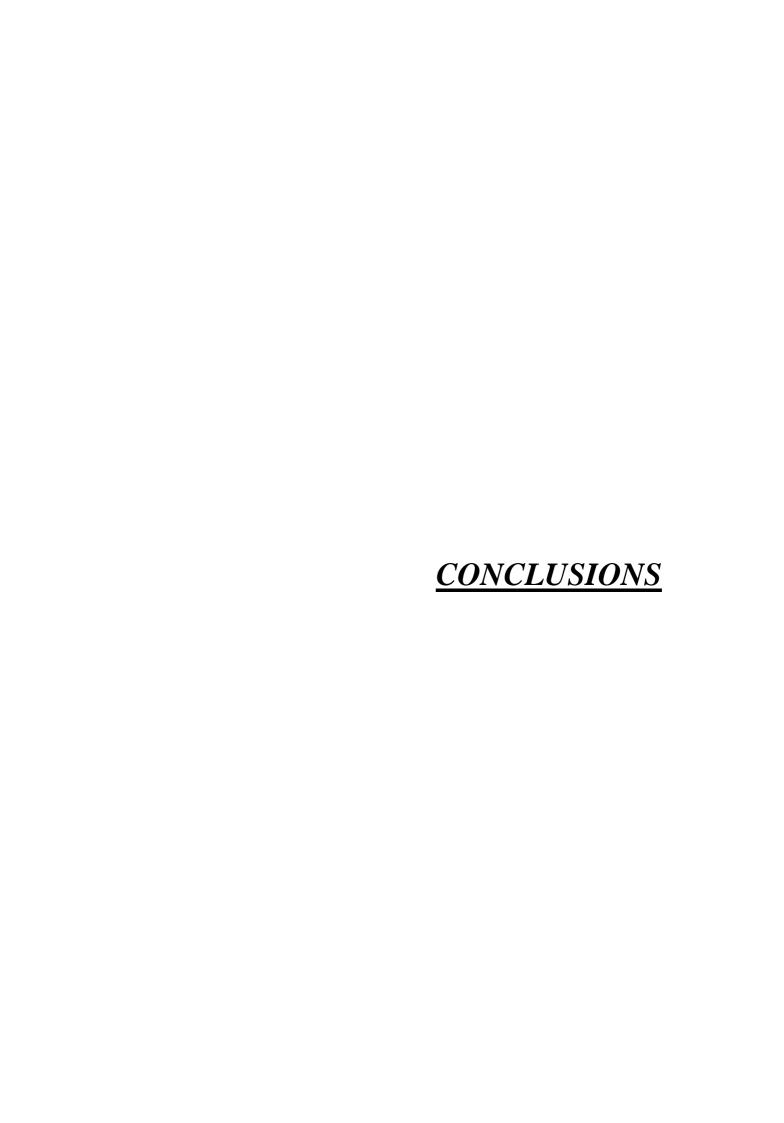
Different studies and available toxicity data relative to IMI are used for the derivation of specific toxicity values used in environmental risk assessment (SERA, 2006; CCME, 2007. These data separate dose-response assessment in six classes of organisms: terrestrial mammals, birds, non-target terrestrial invertebrates, fish, as well as aquatic invertebrates, and algae which were tested in our study. Acute toxicity values of IMI for aquatic invertebrates have a wide range going from a NOAEC (96 h) of 0.00035 mg L<sup>-1</sup> for Hyalella azteca (England and Bucksath, 1991). to a NOAEC (96 h) of 145 mg L<sup>-1</sup> for eastern oyster (Wheat and Ward, 1991). On the basis of these studies, NOAEC values of 0.00035 mg L<sup>-1</sup> and 145 mg L<sup>-1</sup> are chosen to evaluate acute exposure of sensitive and tolerant aquatic invertebrate species, respectively. On other hand, longer-term studies that assessed reproduction, growth and survival demonstrated Mysidopsis bahia as the most sensitive species, with a NOAEC value of 0.000163 mg L<sup>-1</sup> IMI for growth and reproduction endpoints (Ward, 1991), and Daphnia magna as the most tolerant species with a NOAEC (21 d) for immobility of 1.8 mg L<sup>-1</sup> (Young and Blake, 1990). A 19-week microcosm study conducted as part of EPA's pesticide registration requirements for IMI confirms the sensitivity of amphipods and midges observed in laboratory studies (Moring et al., 1992). In addition, on the basis of Moring et al. (1992) 19 week microcosm study a recommended regulatory NOAEC for IMI of 0.006 mg L<sup>-1</sup> was suggested. For freshwater algae risk assessment, NOEC for sensitive species was set at of 6.69 mg L<sup>-1</sup> based on the study with *Navicula pelliculosa* (Hall, 1996), while for tolerant species was of 119 mg L<sup>-1</sup> based on the study with Selenastrum capricornutum (Gagliano and Bowers, 1991).

As in the human health risk assessment, the environmental risk assessment uses predictive environmental scenarios for an accidental spill that involves the contamination of a small water bodies with IMI. The HQs in these cases for sensitive aquatic invertebrates are extremely high, going from 500 to over 50,000. Although, the probability of such spills may be isolated, these HQ clearly suggest that the greatest risk in the event of an accidental spill will be to aquatic non-target invertebrates (SERA, 2005).

The wish to have estimates of no effect toxicant concentrations is highly desirable and needed, especially in case of non-target organisms. The NOEC, however, can be easily misleading, because of its problematic statistical properties. The interpretation of results from toxicity experiments frequently uses NOECs which are then also applied in human and environmental risk assessments. However this use is under constant criticism due to a substantial statistical problem associated to this concept. Although an alternative is necessary, none of proposed alternative methods to characterise toxic effects is still generally accepted. Therefore, in this study as a selective and supportive tool for evaluation of IMI adverse effects on either *in vitro* or *in vivo* or biological system were evaluated also NOECs and LOECs.

Present *in vivo* study was designed to provide additional information on toxic effects of IMI and its TPs on non-target aquatic organisms and to determine threshold concentrations (i.e. NOECs/LOECs) of these compounds through examination of different endpoints/biomarkers (such as: antioxidant enzymes, LP, survival, growth and behaviour). Interestingly, some recent studies tried to correlate and compare *in vitro* cell toxicity data with the *in vivo* toxicity thresholds reported in literature for aquatic organisms; fish or invertebrates. In particular, human cells kept in culture appeared less sensitive to insecticides than above-mentioned aquatic species. In a study of Skandrani *et al.* (2006) was reported that IMI's LOEC values for fish or invertebrates were of 1.2 and 1.8 μg mL<sup>-1</sup> while in neuronal and pulmonary human cell lines LOECs were about 100-300 times higher. Similar deductions and significant differences between LOEC values for neuronal cell lines and the ones for invertebrates were confirmed also in our study. Consequently, these organisms seem to be useful for monitoring of aquatic pesticide pollution.

Toxicity testing of pesticides in aquatic environment should be performed on whole organism models at different levels of biological organisation allowing a better comparison of the data. In addition, applying a battery of bioassays and of measured biomarkers we can reflect different species susceptibility to xenobiotics and also the complexity of entire ecosystems. In conclusion, with the increasing use of IMI, this insecticide could be available in the environment during concurrent periods and could pose a possible cumulative risk to non-target organisms (invertebrates, algae, bacteria as well as humans).



#### 5. Conclusions

- Neuronal cell model applied in this research demonstrated an oxidative stress cellular pattern due to IMI action.
- *V. fischeri* luminescence inhibition confirmed as a suitable biomarker; 6CNA resulted clearly as the most biologically active.
- Algae showed chemically and biologically diverse action of Confidor 200SL

   (a. i. IMI) and its TP 6CNA; interestingly Confidor 200SL had inhibitory,
   while 6CNA stimulatory effect on algae growth (hormesis).
- In amphipods as in algae IMI and 6CNA expressed different modes of action; 6CNA affected adversely on whole-cell level not only on neuronal system.
- Behaviour alterations resulted as promising physiological biomarker for neonicotinoid exposure.
- MXR mechanisms (effect on cell membrane level) resulted as a promising and sensitive molecular biomarkers for neonicotinoid exposure; with future standardisation could be a possible fast screening biomarker to range substances based on their biological potential.
- IMI toxicity testing should be performed on whole organism models (*in vivo*) at different levels of biological organisation applying a battery of bioassays and a multi-biomarker approach.
- *In vivo bio* assays should be integrated with *in vitro* (cell model) assays as they give information about IMI effects on cellular level.

- IMI transformation product should be considered on its own in order to
  recognise additional IMI-toxicity effects and it is reasonable to test these
  products for hazard identification purposes. A separate assessment of IMI
  TPs needs to be made, as the physicochemical properties, persistence and
  toxicity of these compounds may be very different from those of their parent
  compounds. They may also be present at concentrations similar to or even
  exceeding the parent compound
- Commercial formulations containing IMI should be also evaluated during comparative toxicity test for potential adverse effect to non-target groups as they are the one applied in the environment and should be considered in detail for human health and environmental risk assessment.
- All toxicity tests relative to IMI should last long enough to compromise the formation of transformation products (degradation products/metabolites) or pesticide "ageing".

#### 6. ANNEXES

2.1384

1.4488

1.7205

1.3339

1.4097

1.3155

1.4671

Table 18: Absorbance values and % of cell survival of MTT assay experiments with F11 cells after exposure to IMI, Confidor 200SL and 6CNA for 24 h (A) and 48 h (B). (n=6).

#### A MTT assay (F11 cells) Absorbance values 550 nm % of cell survival IMIDACLOPRID IMIDACLOPRID 24 h 24 h control 1 uM 30 μΜ 100 μM 200 μM 500 μM 1000 μM 4000 μM 30 uM 100 uM 200 uM 500 uM 1000 uM 4000 uM 1 mM1.534 1.2434 100 98,2269 1.5068 1.3068 1.3998 1.3826 1.1818 1.3166 85,189 91,2516 90,1304 77,0404 85,8279 81,0561 1.4745 1.3035 1.2791 1.2776 1.1758 1.2189 1.1707 1.1063 100 88.4028 86.7481 86.6463 79,7423 82.6653 79,3964 1.4268 1.1481 1.0415 1.1523 1.1423 1.1001 1.2058 1.1793 100 80.4668 72,9955 80.7611 80.0603 77.1026 84.5108 82,6535 1.6777 1.7092 1.5328 1.5499 1.6467 1.668 1.5104 1.4771 100 101.878 91.3632 92.3824 98.1522 99,4218 90.028 88.0432 1.5832 1.452 1.3738 1.2797 1.3913 1.4154 1.485 1.3022 91.713 86.7736 80.83 87.879 89.4012 93,7974 82.2511 100 1.4884 1.4124 1.3464 1.3946 1.3614 1.3949 1.4903 1.5373 100 94.8938 90,4596 93.6979 91.4673 93.7181 100.128 103.154 CONFIDOR 200SL CONFIDOR 200SL 24 h 100 uM 200 uM 500 uM 1000 uM 4000 uM 30 uM 100 μM 200 μM 500 μM 1000 μM 4000 μM control 1 uM 30 uM control 1 uM 81.1591 1.2182 1.2041 1.2117 1.1894 1.1734 1.081 0.98868 0.96044 100 98.8426 99.4664 97.6359 96.3224 88.7375 78.8409 1.182 1.0828 1.173 1.1372 1.0042 1.0058 1 0896 0.8223 91.6074 99.2386 96.2098 84.9577 85.0931 92.1827 69.5685 100 1.2351 1.1546 1.1779 1.1173 1.1461 0.989 1.0639 0.96711 100 93,4823 95.3688 90.4623 92,7941 80.0745 86,1388 78.3022 1.2656 1.2455 1.3733 0.84955 79.5591 109.845 108.51 1.2783 1.0069 1.3902 1.1739 100 101.003 98.4118 92,7544 67.1263 1.3187 80.0085 1.4856 1.3172 1.2965 1.2479 1.3374 1.282 90.1347 79.9175 78.6616 75.7129 1.6482 81.1431 77.7818 1.2956 1.2447 1.1904 1.4922 1.3501 1.237 1.3573 1.1784 100 96.0713 91.8802 115.174 104.207 95,477 104.762 90.954 6CNA 6CNA 24 h 24 h control $1 \, \mu M$ $30\,\mu M$ $100~\mu M - 200~\mu M$ 500 μΜ 1000 μΜ 4000 μΜ control $1~\mu M$ 30 µM $100 \ \mu M - 200 \ \mu M - 500 \ \mu M - 1000 \ \mu M - 4000 \ \mu M$ 1.2166 1.0528 1.1022 1.1092 1.0563 1.0339 1 0903 100 86.5362 90.5967 91.1721 86.8239 84.9827 1.1184 91.9283 1.0164 0.99646 0.90174 1.0546 1.0396 1.0072 1.0609 1.1034 100 98.0382 88.719 103.758 102.283 99.0948 104.378 108.56 0.8212 0.9781 0.93719 0.91524 0.97308 1.0766 1.001 1.1433 100 119.106 111.452 118.495 131.101 139,223 114.124 121.895 1.3485 1.2475 1.4424 1.2992 1.2058 1.2908 1.3183 1.2134 108.096 115.623 104.144 96.6573 103.471 105.675 97.2665 100 1 4264 1 5517 1 7926 1 7254 1 7999 1 5655 1 4491 1 5748 100 108 784 125 673 120 962 126 185 109 752 101 591 110 404 1.1497 1.1687 1.3349 1.2354 1.2732 1.4061 1.3183 1.1636 114.221 105.707 98.3743 108.942 120.313 112.801 99.5636 B MTT assay (F11 cells) Absorbance values 550 nm % of cell survival IMIDACLOPRID IMIDACLOPRID 48 h 48 h 2.5901 2 6651 2 5532 2.5003 2 487 2 4935 2 5575 2 115 100 102 896 98 5753 96 533 96 0195 96 2704 98 7414 81 6571 2.7437 2.4492 2.5065 2.413 1.7082 89.2663 91.3547 87.9469 62,259 2.4206 2.4868 2.1467 100 88.2239 90.6367 78.2411 91.5468 2.7055 2.6162 2.7093 2.5411 2,4299 2.3528 2.4768 1.9808 100 96.6993 100.14 93.9235 89.8133 86,9636 73.2138 2 3015 2.0315 2 1576 2 2086 2 3867 1 9303 1 6966 1 4175 100 88 2685 93 7476 95 9635 103 702 83 8714 73 7171 61 5903 1.9748 85.5354 56.6061 2.3395 2.1931 2.3032 2.0011 2.1586 2.1216 1.3243 100 93.7423 98.4484 92.2676 90.686 84.4112 2.3424 2.4123 2.2934 2.1754 95.0842 97.9217 96.8541 88.3053 2.4635 2.386 1.9841 100 93.0952 80.5399 CONFIDOR 200SL CONFIDOR 200SL 48 h 48 h control 1 mM30 uM 100 uM 200 uM 500 uM 1000 uM 4000 uM control 1 mM30 uM 100 uM 200 uM 500 uM 1000 uM 4000 uM 2.6443 2.3459 100.93 88.7153 95.1745 87.0022 87.6338 85.8412 47.9295 2.6689 2.5167 2.3006 2.3173 2.2699 1.2674 100 2.6555 2.4446 2.244 2.3475 2.2865 1.0427 92.058 59.292 1.5745 1.5675 100 84.5039 88.4014 86.1043 59.0284 39.2657 2 4745 2 3363 2 1728 2 2437 2 1497 2 1873 1 6504 1.0091 100 94 415 87 8076 90 6729 86 8741 88 3936 66 6963 40.78 2.5724 2.4159 2.4132 2.3581 2.3107 2.1604 2.0212 1.4177 93.9162 93.8112 91,6693 89.8266 83.9838 78.5725 55.112 100 73.3876 2.5165 2.1481 2.2779 1.8468 2.0286 1.7089 1.907 1.3817 85.3606 90.5186 80.612 67.9078 2 4527 2.1931 2 1741 2.239 2.289 2.2346 1 9663 1.2072 100 89.4157 88.6411 91.2872 93.3257 91.1078 80.1688 49.2192 6CNA 6CNA 48 h 48 h control 1 uM 30 uM 100 uM 200 uM 500 uM 1000 uM 4000 uM control 1 uM 30 uM 100 uM 200 uM 500 uM 1000 uM 4000 uM 2.6133 2.4252 2.2529 1.7692 92.8022 86.209 70.5468 2.0113 1.997 1.8436 1.9181 100 76.964 76.4168 73.3976 67.6998 2 4027 2 267 2 1852 2.0776 1 9032 1 8666 2.0029 1 9233 100 94 3522 90 9477 86 4694 79 2109 77 6876 83 3604 80 0474 2.0372 1.5718 1.7937 1.6244 84.8409 79.8392 65,4589 74.7001 67.6495 2.4012 1.9171 1.8864 1.6498 100 78,5607 68,7073 2.4101 2.2093 2.1409 2.108 2.0925 2.0274 1.8436 91.6684 90.5357 88.8303 87.4653 86.8221 76.4948 2 3035 2 2244 2.0992 2.0868 1.8623 1 9106 1 7915 1 8783 100 96 5661 91 1309 90 5926 80 8465 82 9433 77,773 81 5411

100 67,7516

80.4574

62,3784

65,9231

61.518

68,6074

59.8906

1.2807

Table 19: Absorbance values and % of cell survival of MTT assay experiments with F11 cells after exposure to desnitro-IMI, 5-hydroxy-IMI and olefin-IMI for 24 h (A) and 48 h (B). (n=6).

Λ		

#### MTT assay (F11 cells)

Absorbance values 550 nm

desnitro-I	MI							desnitro-IMI 24 h	
	1 μM	30 μΜ	100 μM	200М	500 uM	1000 uM	4000 uM	control 1 μM 30 μM 100 μM 200 μM 500 μM 1000 μM 4000 μM	Л
3.1422	3.3636			3.1657	3.1371	3.0605	2.8288	100 103.166 100.628 100.32 100.287 100.291 98.8086 79.604	
3.1643	3.1928	3.1719	3.2005	3.1662	3.175	3.1731	2.2558	100 103.100 100.028 100.32 100.287 100.291 98.8080 79.004	
3.178	3.1928	3.1719		3.1798	3.2273	3.1379	2.4655	100 103.127 97.8399 90.2020 97.0397 90.1829 93.8343 80.730	
3.178	3.334	3.123	3.1746	3.1798	3.137		2.4633	100 97.8900 97.2498 98.1267 97.0731 97.3449 97.2800 69.162	
						3.1657			
3.421	3.321	3.165	3.134	3.145	3.123	3.1662	2.345	100 102.22 95.7506 98.418 97.0383 96.1798 97.0597 87.227	
3.12	3.432	3.187	3.2413	3.1234	3.1367	3.0798	2.213	100 101.821 97.0383 96.0878 96.4251 95.7506 97.0751 71.897	2
5-hvdroxy	-IMI							5-hydroxy-IMI	
24 h								24 h	
	1 μM	30 μΜ	100 μM	200 uM	500 uM	1000 uM	4000 μM	control 1 μM 30 μM 100 μM 200 μM 500 μM 1000 μM 4000 μM	Л
3.1572	3.1883	3.2085	3,1854	3,1646		2.9085	1,326	100 100.65 100.136 99.9621 100.184 99.4603 95.1825 55.997	
3.2061	3.2628	3.1484	3.1594	3.1731	3.1192	3.0806	2.095	100 102.983 99.3719 99.7191 100.152 98.4503 97.232 66.123	
3.1416	3.1156	3.1609	3.1565	3.1847	3.1641	3.0579	1.9015	100 98.3366 99.7664 99.6276 100.518 99.8674 96.5155 60.016	
3.1567	3.111	3.201	3.147	3.1567	3.137	3.165	1,425	100 98.1915 101.032 99.3277 99.6339 99.0121 99.8958 44.976	
3.1546	3.123	3.128	3.124	3.111	3.123	2.937	1.982	100 98.5702 98.728 98.6018 98.1915 98.5702 92.6996 62.557	
3.1456	3.147	3.18	3.112	3.201	3.1367	3.0657	2.1784	100 99.3277 100.369 98.223 101.032 99.0026 96.7617 68.756	
			*****						-
olefin-IMI	ſ							olefin-IMI	
24 h	-							24 h	
control	1 uM	30 μΜ	100 μM	200 uM	500 uM	1000 uM	4000 μM	control 1 μM 30 μM 100 μM 200 μM 500 μM 1000 μM 4000 μM	Л
3.166	2.882			3.2659		3.1732	2.3752	100 95.4491 99.5583 100.194 101.466 99.8261 94.439 69.554	
3.0261	3.096	3.178	3.1759	3.1777	3.1593	3.0514	2.0955	100 100.843 103.513 103.445 103.504 102.904 99.3898 68.254	4
3.0183	3.0741	3.1852	3.167	3.1791	3.106		2.1256	100 100.129 103.748 103.155 103.549 101.168 88.9766 69.234	
3.176	2.782	3.1786		3.2659	3.2219	3.1743	2.2752	100 90.615 103.533 103.969 106.376 104.943 103.393 74.107	
3.1261	3.076	3.198	3.159	3.1872	3.1494	3.1514	2.0945	100 100.191 104.165 102.895 103.813 102.582 102.647 68.221	
3.0983	3.1441	3.1652	3.167	3.1797		2.75147	2.1275	100 102.409 103.096 103.155 103.569 101.494 89.6205 69.296	
5.0705	5.1441	5.1052	3.107	5.1171	3.110	2.73147	2.12/3	100 102.407 103.070 103.133 103.307 101.474 07.0203 07.270	.,

#### В

### MTT assay (F11 cells)

Absorbance values 550 nm

ue	SIIII	10-	LIVI	.1
48	h			

control	1 μΜ	30 μΜ	100 μM	200 μΜ	500 μΜ	$1000 \mu M$	4000 μΜ
3.2228	3.3682	3.1855	3.2054	3.2321	2.133	1.0293	0.66286
3.1998	3.1926	3.155	3.1863	3.1699	3.1673	2.894	0.56907
3.2029	3.1781	3.1889	3.1786	3.174	3.1722	3.2313	0.59862
3.2328	3.546	3.1335	3.2051	3.2341	2.103	1.0193	0.6686
3.1548	3.1846	3.134	3.1663	3.1674	3.153	2.794	0.5407
3.2089	3.1651	3.1619	3.1762	3.1274	3.1922	3.1263	0.58262

#### 5-hydroxy-IMI

48 n							
control	1 μΜ	$30  \mu M$	$100  \mu M$	200 μΜ	500 μΜ	$1000~\mu\text{M}$	$4000~\mu M$
3.1807	3.0974	3.2092	3.1785	3.1871	2.8908	2.9582	0.77138
3.3291	3.2984	3.19	3.1679	3.1814	3.1819	2.2017	0.71477
3.1511	3.1738	3.1522	3.1501	3.1471	3.0852	2.8612	1.0027
3.1827	3.094	3.292	3.1685	3.181	2.8708	2.9762	0.778
3.3143	3.283	3.123	3.1699	3.1904	3.1769	2.2019	0.71457
3.1501	3.1728	3.1502	3.1461	3.151	3.0452	2.7822	1.2027

#### olefin-IMI

control	1 μM	30 μΜ	100 μM	200 μΜ	500 μM	$1000 \mu M$	$4000 \mu M$
3.187	3.2021	3.1601	3.1535	3.2824	3.1692	3.1554	0.66549
3.3054	3.1553	3.1953	3.1409	3.1365	3.1832	3.1525	0.71075
3.1267	3.1098	3.155	3.1691	3.1557	3.1498	3.1275	0.9881
3.1101	3.161	3.182	3.1409	3.108	3.0925	3.0274	0.8436
3.3035	3.1853	2.992	3.1868	3.2623	3.106	3.1915	0.8782
3.1384	3.155	3.1205	3.1339	3.1097	3.155	3.1671	0.6807

#### % of cell survival

#### desnitro-IMI

40 II								
control	1 μM	30 μΜ	$100 \mu M$	200 μΜ	500 μM	$1000~\mu M$	$4000  \mu M$	
100	101.178	99.0016	99.4265	99.4857	88.0214	74.3296	19.0177	
100	99.5044	98.3326	99.3081	98.7969	98.7159	90.1979	17.7363	
100	99.0525	99.3891	99.0681	98.9247	98.8686	100.711	18.6573	
100	110.519	97.6625	99.894	100.798	65.5446	31.7687	20.8384	
100	99.2551	97.678	98.6847	98.719	98.2702	87.0812	16.8521	
100	98.6473	98.5476	98.9933	97.4723	99.492	97.4381	18.1586	

#### 5-hydroxy-IMI

control	1 μΜ	30 μΜ	$100  \mu M$	200 μΜ	500 μΜ	$1000\;\mu\text{M}$	$4000~\mu M$
100	99.055	98.8666	98.2983	98.496	94.7935	83.0264	29.9025
100	102.425	99.0591	98.3728	98.792	98.8076	68.3694	32.1925
100	98.556	97.8853	97.8201	97.7269	95.8047	88.8489	31.1369
100	96.078	102.227	98.3915	98.7796	89.147	92.42	35.0403
100	101.947	96.9785	98.4349	99.0715	98.6523	68.3756	32.1835
100	98.525	97.8232	97.6959	97.848	94.5626	86.3957	37.3475

#### olefin-IMI

ontrol	1 μM	30 μM	100 μM	200 μM	500 μM	1000 μM	4000 μM
100	98.4209	98.87	98.3824	99.5374	98.7847	98.0903	24.5796
100	98.4073	99.6549	97.9582	97.821	99.2775	98.32	22.1668
100	96.9883	98.398	98.8377	98.4198	98.2358	97.5403	30.8168
100	98.5851	99.2401	97.9582	96.9321	96.4487	94.4184	26.3102
100	99.343	93.3143	99.3898	101.744	96.8698	99.5363	27.3893
100	98.398	97.322	97.7399	96.9852	98.398	98.7754	21.2296

Table 20: Absorbance values and % of cell survival of MTT assay experiments with F11 cells after pre-treatment (30 min) with p38 inhibitor SB203580, vitamin C (NAC) and mixture of tocopherols (vitamin E) in presence of 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 24 h (A) and 48 h (B). (n=6).

#### $\mathbf{A}$

MTT	assav	/F11	colle	۱
NI I	assav	(FII	ceus	)

MITASSA	y (F11 ceus	)		
Absorbar	nce values 5	50 nm		
p38 inhib 24 h	itor (SB203	580)		
control	DMSO	DMSO+NMP	SB203580+ 1000 μM IMI	SB203580+ 4000 µM IMI
0.372	0.41	0.427	0.439	0.431
0.539	0.476	0.473	0.359	0.539
0.387	0.395	0.421	0.379	0.442
0.455	0.492	0.41	0.315	0.543
0.455	0.41	0.41	0.315	0.54
0.487	0.45	0.42	0.439	0.53
vitamin E	2			
24 h	DMGO	DMGO MM	THE THEORY MAN	: F. 1000 M.B.C
control	DMSO	DMSO+NMP	vitamin E+1000 μM IMI	vitamine E+4000 μM IMI
0.372	0.41	0.427	0.415	0.431
0.539	0.476	0.473	0.433	0.539
0.387	0.395	0.421	0.259	0.442
0.455	0.492	0.41	0.395	0.543
0.455	0.41	0.41	0.415	0.543
0.487	0.45	0.42	0.395	0.44
NAC				
24 h				
control	DMSO	DMSO+NMP	NAC+1000 μM IMI	NAC+4000 µM IMI
0.372	0.41	0.427	0.369	0.431
0.539	0.476	0.473	0.483	0.539
0.387	0.395	0.421	0.329	0.442
0.455	0.492	0.41	0.342	0.543
0.455	0.41	0.41	0.342	0.45
0.487	0.45	0.42	0.369	0.53
% of cell	survival			
p38 inh 24 h	nibitor (SB2	03580)		
control	DMSO	DMSO+NMP	SB203580+ 1000 μM IMI	SB203580+ 4000 μM IMI
100	93.56	97.48	100.22	101.63
100	108.67	107.99	81.9	149.38
100	90.18	96.11	86.25	92.21
100	112.32	93.06	71.91	108.4
100	93.56	93.06	71.91	151.2
100	93.30			
100	94	94.01	100.22	149.5
vitamine 24 h	E			
control	DMSO	DMSO+NMP	vitamin E 1000 vM IMI	vitamina E : 4000M IMI
			vitamin E+1000 μM IMI	vitamine E+4000 μM IMI
100	93.56	97.48	94.74	117.41
100	108.67	107.99	98.85	174.79
100	90.18	96.11	59.13	129.5
100	112.32	93.06	90.18	90.16
100	93.56	93.06	94.74	90.16
100	94	94.01	59.13	117.6
NAC				
24 h				
control	DMSO	DMSO+NMP	NAC+1000 μM IMI	NAC+4000 µM IMI
100	93.56	97.48	84.24	96.72
100	108.67	107.99	110.2	81.9
100	90.18	96.11	75.11	112.2
100	112.32	93.06	78.02	84.42
100	93.56	93.06	78.02	82
100	94	94.01	84.24	112.8

B

#### MTT assay (F11 cells)

#### Absorbance values 550 nm

38 inhibitor () 48 h	3 <b>D2</b> 03300)			
control	DMSO	DMSO+NMP	$SB203580 + 1000 \mu M IMI$	SB203580+ 4000 μM IM
1.162	1.296	1.423	1.059	0.801
1.287	1.193	1.202	1.024	0.825
1.24	1.202	1.183	0.947	0.73
1.24	1.198	1.235	1.043	0.828
1.164	1.201	1.236	1.045	0.835
1.288	1.294	1.424	1.05	0.802
vitamine E				
48 h				
control	DMSO	DMSO+NMP	vitamin E+1000 μM IMI	vitamine E+4000 μM IMI
1.162	1.296	1.423	0.936	0.864
1.287	1.193	1.202	1.202	0.876
1.24	1.202	1.183	1.36	0.677
1.24	1.198	1.235	1.049	0.819
1.164	1.201	1.236	1.2	0.822
1.288	1.294	1.424	0.93	0.823
NAC 48 h				
control	DMSO	DMSO+NMP	NAC+1000 μM IMI	NAC+4000 μM IMI
1.162	1.296	1.423	1.37	0.803
1.102	1.193	1.202	1.234	0.75
1.24	1.202	1.183	0.915	0.868
1.24	1.198	1.235	1.2	0.875
1.164	1.198	1.236	1.2	0.88
		1.424	1.2	
1.288	1.294	1.424	1.3	0.87
6 of cell survi	val	1.424	1.3	0.87
	val	1.424	1.5	0.87
6 of cell survi	val			
6 of cell survi 38 inhibitor ( 48 h control	val SB203580) DMSO	DMSO+NMP	SB203580+ 1000 μM IMI	SB203580+ 4000 μM IMI
6 of cell survi 38 inhibitor (s 48 h control 100	val SB203580) DMSO 105.19	DMSO+NMP 115.5	SB203580+ 1000 μM IMI 85.95	SB203580+ 4000 μM IMI 67.07
38 inhibitor (3 48 h control 100 100	DMSO 105.19 96.83	DMSO+NMP 115.5 97.56	SB203580+ 1000 μM IMI 85.95 83.11	SB203580+ 4000 μM IMI 67.07 69.08
38 inhibitor (348 h control 100 100 100	DMSO 105.19 96.83 97.56	DMSO+NMP 115.5 97.56 96.02	SB203580+ 1000 μM IMI 85.95 83.11 76.86	SB203580+ 4000 μM IMI 67.07 69.08 61.12
38 inhibitor (348 h control 100 100 100 100 100 100	DMSO 105.19 96.83 97.56 97.24	DMSO+NMP 115.5 97.56 96.02 100.24	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65	DMSO+NMP 115.5 97.56 96.02 100.24 100.3	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02
38 inhibitor (3 48 h control 100 100 100 100 100 100 100	DMSO 105.19 96.83 97.56 97.24	DMSO+NMP 115.5 97.56 96.02 100.24	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33
38 inhibitor (3 48 h control 100 100 100 100 100 100	DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08
38 inhibitor (3 48 h control 100 100 100 100 100 100 vitamine E	DMSO 105.19 96.83 97.56 97.24 97.65	DMSO+NMP 115.5 97.56 96.02 100.24 100.3	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08
38 inhibitor (3 48 h control 100 100 100 100 100 vitamine E 48 h	DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02
38 inhibitor (3 48 h control 100 100 100 100 100 100 vitamine E 48 h control	DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08
38 inhibitor (3 48 h control 100 100 100 100 100 100 vitamine E 48 h control 100	DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7 DMSO+NMP 115.5	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92 vitamin E+1000 μM IMI 75.97	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34
38 inhibitor (3 48 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.11	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7 DMSO+NMP 115.5 97.56	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92 vitamin E+1000 μM IMI 75.97 97.56	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35
38 inhibitor (3 48 h control 100 100 100 100 100  100 100 100 100 1	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56 97.24	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7 DMSO+NMP 115.5 97.56 96.02	SB203580+ 1000 μM IMI  85.95  83.11  76.86  84.65  84.7  84.92   vitamin E+1000 μM IMI  75.97  97.56  110.38	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57
38 inhibitor (3 48 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7 DMSO+NMP 115.5 97.56 96.02 100.24	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68
38 inhibitor (3 48 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56 97.24 97.65	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7 DMSO+NMP 115.5 97.56 96.02 100.24 100.3	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7
38 inhibitor (3 48 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.11  DMSO 105.19 96.83 97.56 97.24 97.65 105.11	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7 NAC+4000 μM IMI
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02  NAC+1000 μM IMI 111.2	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7  NAC+4000 μM IMI 67.23
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.1 DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02  NAC+1000 μM IMI 111.2 100.16	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7  NAC+4000 μM IMI 67.23 62.8
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.11  DMSO 105.19 96.83 97.56 97.24 97.65 105.1  DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02  NAC+1000 μM IMI 111.2 100.16 74.26	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7  NAC+4000 μM IMI 67.23 62.8 72.68
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.11  DMSO 105.19 96.83 97.56 97.24 97.65 105.1  DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02  NAC+1000 μM IMI 111.2 100.16 74.26 97.4	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7 68.7  NAC+4000 μM IMI 67.23 62.8 72.68 73.26
38 inhibitor (348 h control 100 100 100 100 100 100 100 100 100 10	DMSO 105.19 96.83 97.56 97.24 97.65 105.11  DMSO 105.19 96.83 97.56 97.24 97.65 105.1  DMSO 105.19 96.83 97.56 97.24 97.65 105.1	DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02 100.24 100.3 100.7  DMSO+NMP 115.5 97.56 96.02	SB203580+ 1000 μM IMI 85.95 83.11 76.86 84.65 84.7 84.92  vitamin E+1000 μM IMI 75.97 97.56 110.38 85.146 97.7 76.02  NAC+1000 μM IMI 111.2 100.16 74.26	SB203580+ 4000 μM IMI 67.07 69.08 61.12 69.33 70.02 67.08  vitamine E+4000 μM IMI 72.34 73.35 56.68 68.57 68.7 68.7  NAC+4000 μM IMI 67.23 62.8 72.68

Table 21: Mean values relative to p38 signal intensity quantification (grey values of immunopositive signals after background – BG subtraction) after exposure to 400  $\mu$ M (102.2 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 30 min. (n=3).

p38

	$\boldsymbol{\mathcal{A}}$	7 T'	TT	$\sim$	•
	,,	/₩.	ı K	•	•
•	$\boldsymbol{\smile}$		LAN	$\boldsymbol{\smile}$	_

	No. of cell	whole	BG1	BG2	BG3	mean BG	
1	16	25.712	21.679	25.237	21.927	22.94767	2.76433 0.172771
2	23	26.841	24.943	19.467	26.104	23.50467	3.33633 0.145058
3	27	28.423	24.413	31.475	19.748	25.21200	3.21100 0.118926
4	27	24.797	28.104	19.543	20.658	22.76833	2.02867 0.075136
5	20	25.758	18.341	29.615	16.768	21.57467	4.18333 0.209167
6	30	26.34	23.735	27.089	16.871	22.56500	3.77500 0.125833
7	32	25.21	26.114	25.141	18.485	23.24667	1.96333 0.061354
8	22	22.358	25.208	24.409	13.508	21.04167	1.31633 0.059833
9	46	30.138	26.102	28.557	17.929	24.19600	5.94200 0.129174
10	24	30.523	23.897	32.656	26.857	27.80333	2.71967 0.113319

#### 400 μM IMI

	No. of cell	whole	BG1	BG2	BG3	mean BG	
1	16	27.04	18.366	31.274	22.559	24.06633	2.97367 0.185854
2	34	34.9	31.417	37.82	24.962	31.39967	3.50033 0.102951
3	20	28.307	29.774	21.316	21.529	24.20633	4.10067 0.205033
4	30	26.041	27.992	19.943	26.747	24.89400	1.14700 0.038233
5	29	29.458	20.048	33.697	24.991	26.24533	3.21267 0.110782
6	24	25.858	13.657	26.757	17.289	19.23433	6.62367 0.275986
7	18	25.142	17.169	27.379	16.638	20.39533	4.74667 0.263704
8	16	23.301	25.353	14.14	24.927	21.47333	1.82767 0.114229
9	17	26.934	20.588	27.739	19.138	22.48833	4.44567 0.26151
10	19	26.829	18.284	19.587	30.582	22.81767	4.01133 0.211123

## *4000* μM IMI

	No. of cell	whole	BG1	BG2	BG3	mean BG		
1	42	28.964	22.312	27.366	20.829	23.50233	5.46167	0.13004
2	38	35.155	29.39	32.309	22.476	28.05833	7.09667	0.186754
3	29	34.938	32.034	25.658	24.459	27.38367	7.55433	0.260494
4	18	38.737	30.588	29.085	23.939	27.87067	10.86633	0.603685
5	28	30.375	29.084	28.796	20.626	26.16867	4.20633	0.150226
6	22	32.466	25.677	23.85	29.177	26.23467	6.23133	0.283242
7	25	61.177	59.357	50.707	50.939	53.66767	7.50933	0.300373
8	30	32.237	27.696	24.61	29.89	27.39867	4.83833	0.161278
9	21	29.744	19.644	24.006	29.196	24.28200	5.46200	0.260095
10	30	31.173	18.466	29.62	18.311	22.13233	9.04067	0.301356

Table 22: Mean values relative to ERK signal intensity quantification (grey values of immunopositive signals after background – BG subtraction) after exposure to 400  $\mu$ M (102.2 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 30 min. (n=3).

## **ERK**

#### **CONTROL**

	No. of cell	whole	BG1	BG2	BG3	mean BG
1	82	17.859	7.47	7.851	6.712	7.34433 10.51467 0.089565
3	60	8.141	7.795	7.91	5.491	7.06533 1.07567 0.117756
5	40	7.809	7.541	6.988	5.336	6.62167 1.18733 0.165542
6	36	10.303	7.513	6.01	7.13	6.88433 3.41867 0.191231
8	50	14.061	7.661	6.214	8.032	7.30233 6.75867 0.146047
9	46	9.938	5.832	6.823	5.955	6.20333 3.73467 0.134855

## *400* μ**M IMI**

	No. of cell	whole	BG1	BG2	BG3	mean BG		
1	25	17.301	7.338	11.473	10.733	9.84800	7.45300	0.39392
2	25	15.521	10.637	9.896	6.623	9.05200	6.46900	0.36208
3	28	16.977	8.395	10.743	7.638	8.92533	8.05167	0.318762
4	31	15.139	11.125	11.962	7.527	10.20467	4.93433	0.329183
5	22	14.614	7.062	11.473	7.528	8.68767	5.92633	0.394894
6	36	17.299	7.927	11.208	7.754	8.96300	8.33600	0.248972
7	25	13.434	9.962	10.009	7.064	9.01167	4.42233	0.360467
9	29	20.264	9.411	8.373	11.119	9.63433 1	10.62967	0.332218
10	30	22.399	12.439	7.767	10.807	10.33767	12.06133	0.344589

## *4000* μM IMI

	No. of cell	whole	BG1	BG2	BG3	mean BG
1	27	12.439	10.613	10.322	7.08	9.40500 3.63400 0.348333
2	51	18.657	11.775	9.057	11.478	10.43067 8.243333 0.204523
3	32	12.094	10.851	9.828	6.577	9.08533 3.00867 0.283917
4	34	14.964	10.127	10.333	6.579	9.01300 5.95100 0.265088
5	36	17.834	9.403	10.838	6.581	8.94067 8.89333 0.248352
5	27	12.439	10.813	10.322	7.08	9.40500 3.03400 0.348333
6	51	18.687	11.755	8.059	11.478	10.43067 8.25633 0.204523
7	52	18.687	11.855	8.059	11.578	10.43067 8.25633 0.20059
8	27	13.6	10.101	9.423	10.246	9.92333 3.67667 0.367531
9	41	18.574	14.768	13.448	15.932	14.71600 3.85800 0.358927
10	46	13.594	11.74	10.286	11.54	11.18867 2.40533 0.243232

Table 23: Mean values relative to Nrf2 signal intensity quantification (grey values of immunopositive signals after background – BG subtraction) after exposure to 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 30 and 240 min (n=3).

#### Nrf2

f2					
		con	trol; 24h		
		cell brightness			
	cell No.	Nrf2(mean)	cell BG1	cell BG2	cell BG3
1 2	42 55	44.937 49.608	34.889 32.332	35.784 31.295	35.254 30.006
3	54	77.607	49.516	50.382	48.519
4	70	60.476	43.369	36.046	44.796
5	68	56.913	42.287	33.698	37.458
6	32	86.482	70.93	70.099	72.001
7	22	88.226	66.632	72.145	68.774
8	46	55.404	44.973	36.903	37.184
9	54	53.72	41.746	32.399	46.98
10	42	65.884	56.385	53.043	55.499
		Mean BG	Nrf2	Nrf2/cell	
		35.309 31.211	9.628 18.397	0.229238095 0.334490909	
		49.47233333	28.13466667	0.521012346	
		41.40366667	19.07233333	0.272461905	
		37.81433333	19.09866667	0.280862745	
		71.01	15.472	0.4835	
		69.18366667	19.04233333	0.865560606	
		39.68666667	15.71733333	0.341681159	
		40.375	13.345	0.24712963	
		54.97566667	10.90833333	0.259722222	
		4000 ր	ι <b>M</b> ; 30 min		
	cell No.	cell brightness Nrf2(mean)	cell BG1	cell BG2	cell BG3
1	36	56.109	48.124	36.679	46.437
2	25	58.418	41.141	46.341	42.499
3	58	67.952	39.791	39.398	53.114
4	32	109.803	97.237	104.235	99.341
5	24	53.038	46.358	37.933	37.836
6	46	62.136	41.516	49.104	41.237
7	35	58.025	40.584	49.52	39.922
8	31	52.866	36.302	42.351	38.895
9	26	54.562	35.297	40.729	40.679
10	25	63.755	45.383	51.91	48.82
		Mean BG 43.74666667	Nrf2 12.36233333	Nrf2/cell 0.343398148	
		43.7400007	12.30233333	0.60364	
		44.101	23.851	0.411224138	
		100.271	9.532	0.297875	
		40.709	12.329	0.513708333	
		43.95233333	18.18366667	0.395297101	
		43.342	14.683	0.419514286	
		39.18266667	13.68333333	0.441397849	
		38.90166667	15.66033333	0.602320513	
		48.70433333	15.05066667	0.602026667	
		<b>4000</b> μ	M; 240 min		
	II NI-	cell brightness		II BC2	II DC2
1	cell No.	Nrf2(mean) 86.641	cell BG1 75.978	cell BG2 85.3	cell BG3 85.209
2	43	70.849	52.303	49.266	57.792
3	21	64.554	54.483	47.043	50.365
4	17	61.688	51.361	53.95	53.338
5	25	68.01	55.05	59.536	53.832
7	41	69.22	51.557	51.641	49.535
8	30	141.842	132.231	138.322	118.34
9	27	64.673	61.644	50.217	50.077
10	26	68.804	65.998	55.201	62.475
		Mean BG	Nrf2	Nrf2/cell	
		82.16233333	4.478666667	0.319904762	
		53.12033333	17.72866667	0.412294574	
		50.63033333	13.92366667	0.663031746	
		52.883	8.805	0.517941176	
		56.13933333	11.87066667	0.474826667	
		50.911	18.309	0.446560976	
		129.631	12.211	0.407033333	
		53.97933333	10.69366667	0.396061728	

7.579333333

0.291512821

61.22466667

Table 24: Mean values relative to JC-1 signal intensity quantification (JC-1 green-red signal ratio) and % of cells with higher MMP (red signal) after exposure to 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 15 and 60 min (n=3).

JC-1

Control				Control			
sample n.	ALL CELLS	red signal	% cells (higher MMP)	sample n.	ALL CELLS	red signal	% cells (higher MMP)
1	110	24	21.8	1	61	6	9.83
2	106	36	33.96	2	72	13	18.05
3	89	26	29.21	3	81	13	16.04
4	139	64	46.04	4	60	13	18.33
5				5			
	115	60	52.17		100	12	12
6	105	54	51.42	6	97	18	18.55
7	109	38	34.86	7	76	16	21.05
8	118	88	74.57	8	67	25	37.31
9	131	48	36.64	9	69	20	28.98
10	47	27	57.44	10	70	10	14.28
1	54	29	53.7	1	64	9	14.06
2	53	17	32.07	2	66	40	60.6
3	97	28	28.86	3	86	43	50
4	125	63	50.04	4	82	38	46.34
5	118	72	61	5	114	54	47.36
4000M. 15	••			4000 Mr. 15			
4000 μM; 15				4000 μM; 15			
sample n.	ALL CELLS	red signal	% cells (higher MMP)	sample n.	ALL CELLS	red signal	% cells (higher MMP)
1	12	7	58.3	1	94	19	20.21
2	49	12	24.48	2	95	30	31.57
3	75	4	5.33	3	109	13	11.92
4	35	15	42.85	4	37	1	2.7
5	55	25	45.45	5	103	17	16.5
6	36	16	44.44	6	108	9	8.33
7	43	13	30.23	7	92	18	19.56
8	42	13	30.95	8	76	18	23.68
9	43	13	30.23	9	46	20	43.47
10	79	14	17.27	10	81	14	17.28
1	108	10	9.25	1	73	7	9.58
2	47	14	29.78	2	51	10	19.6
3	31	12	38.7	3	98	16	16.32
4	36	12	33.33	4	44	20	45.45
5	19	8	42.1	5	87	25	28.73
4000 μM; 60	) min			4000 μM; 60	min		
sample n.	ALL CELLS	red signal	% cells (higher MMP)	sample n.	ALL CELLS	red signal	% cells (higher MMP)
1	52	12	22.64	1	99	9	9.09
2	42	18	42.85	2	29	10	34.48
3	45	17	37.77	3	63	11	17.46
4	24	9	37.5	4	38	13	34.21
5	17	6	35.29	5	69	12	17.39
6	14	11	78.57	6	83	5	6.02
7	35	13	37.14	7	108	11	10.18
8	20	6	30	8			
					83	6	7.22
9	31	9	29.03	9	56	10	17.85
10	43	13	30.23	10	90	7	7.77
1	60	39	65	1	90	6	6.66
2	57	23	40.35	2	60	5	8.33
3	90	30	33.33	3	61	6	9.83
4	69	34	49.27	4	82	10	12.19
5	74	24	32.43	5	69	7	10.14

Table 25: CAT activity in F11 cells exposed to 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 24 and 48 h. Representative absorbance (AU) values of whole 2 min measurement (selected for every 30 s) and data normalised on total protein content ( $\mu$ mol/min/mg protein). (n=3).

CAT							
24 h							
Control				(	Calculated valu	ues normalised on pro	tein content
time	AU	AU	AU	24 h	μmol/min	mg protein sample	μmol/min/mg prot
0 s	0.008508	0.4167	0.1138	control	0.051	0.0257	1.98
30 s	0.007056	0.41096	0.11353	1000 μM	0.038	0.0256	1.48
60 s	0.00676	0.40603	0.11289	4000 μM	0.15	0.0184	8.15
90 s	0.007455	0.40235	0.11262				
120 s	0.006561	0.40038	0.11225	control	0.028	0.0862	0.32
1000 μM IMI				1000 μM	0.294	0.0604	4.86
time	AU	AU	AU	4000 μM	0.184	0.048	3.83
0 s	0.014257	0.30639	0.06278	•			
30 s	0.014281	0.30271	0.06139	control	0.04	0.0429	0.93
60 s	0.013046	0.29939	0.06156	1000 μM	0.025	0.049	0.51
90 s	0.013548	0.29705	0.06026	4000 μM	0.043	0.0448	0.88
120 s	0.012263	0.29519	0.05979				
<b>4000 μM IMI</b>							
time	AU	AU	AU				
0 s	0.166809	0.24196	0.05851				
30 s	0.163384	0.23991	0.05749				
60 s	0.16239	0.23788	0.05719				
90 s	0.160803	0.23615	0.05735				
120 s	0.160877	0.23494	0.05687				
48 h							
Control				(	Calculated vali	ues normalised on pro	tein content
time	AU	AU	AU	48 h	μmol/min	mg protein sample	µmol/min/mg prot
0 s	0.033807	0.16851	0.19837	control	0.04	0.032	1.25
30 s	0.03209	0.1638	0.19618	1000 μM	0.014	0.0378	0.37
60 s	0.032174	0.16417	0.19524	4000 μM	0.005	0.0242	0.2
90 s	0.032248	0.16719	0.1947				
120 s	0.032269	0.1637	0.19429	control	0.049	0.041	1.19
1000 μM IMI				1000 μΜ	0.073	0.0388	1.88
time	AU	AU	AU	4000 μM	0.133	0.02	6.65
0 s	0.017433	0.09331	0.03723	•			
30 s	0.017026	0.09245	0.03701	control	0.034	0.111	0.3
60 s	0.016976	0.09228	0.03673	1000 μM	0.139	0.024	5.79
90 s	0.016976	0.09249	0.03656	4000 μM	0.076	0.022	3.45
120 s	0.016057	0.09218	0.03612				
4000 μM IMI							
time	AU	AU	AU				
0 s	0.018725	0.09583	0.1073				
30 s	0.018338	0.09563	0.10771				
60 s	0.018511	0.09403	0.10556				
90 s	0.018593	0.09385	0.10419				
120 s	0.018522	0.09373	0.10348				

Table 26: Absorbance (AU) values for LP measurements in F11 cells exposed to 1000  $\mu$ M (255.6 mg  $L^{-1}$ ) and 4000  $\mu$ M (1022.4 mg  $L^{-1}$ ) IMI for 24 and 48 h and data normalised on total protein content (nmol//mg protein of MDA). (n=3).

LP							
24 h				48 h			
	AU	AU	AU		AU	AU	AU
	535 nm	600 nm	535-600 nm		535 nm	600 nm	535-600 nm
control	0.02	0.012	0.008	control	0.006	0.01	0.005
control	0.005	0.001	0.004	control	0.01	0.007	0.003
control	0.009	0.002	0.007	control	0.005	0.001	0.004
control	0.006	0.001	0.005	control	0.003	0	0.003
control	0.003	0.001	0.002	control	0.008	0.003	0.005
control	0.007	0.001	0.006	control	0.002	0	0.002
1000 μΜ	0.031	0.01	0.021	1000 μM	0.019	0.007	0.012
1000 μΜ	0.032	0.011	0.021	1000 μΜ	0.02	0.006	0.014
1000 µM	0.051	0.003	0.048	1000 μM	0.005	0.001	0.004
1000 µM	0.022	0.004	0.018	1000 μM	0.005	0.001	0.004
$1000 \mu M$	0.045	0	0.045	1000 μM	0.013	0.004	0.011
1000 μΜ	0.014	0.001	0.013	1000 μΜ	0.011	0.002	0.009
4000 μΜ	0.46	0.005	0.455	4000 μM	0.102	0.01	0.092
$4000 \mu M$	0.41	0.003	0.407	4000 μM	0.423	0.01	0.413
$4000 \mu M$	0.077	0.002	0.075	4000 μM	0.014	0.002	0.012
$4000 \mu M$	0.059	0.005	0.054	4000 μM	0.058	0.006	0.052
$4000 \mu M$	0.209	0.003	0.206	4000 μM	0.013	0.004	0.009
4000 μΜ	0.313	0.004	0.309	4000 μM	0.015	0.002	0.013
MDA nmol/ml me	edium			MDA nmol	/ml mediı	ım	
	control	1000 µM	4000 µM		control	1000 µM	4000 µM
	1.05	2.4	18.45		1.05	2.4	18.45
	0.375	2.7	56.25		0.375		56.25
	0.75	0.75	2.25		0.75		2.25
	0.375	0.45	9		0.375		9
	1.05	2.55	1.8		1.05		1.8
	0.27	1.8	2.55		0.27		2.55
normalised MDA				normalised	_	/mg protein	
	15.69	13.333	87.857		15.69		87.857
	8.09	24.545	274		8.09		274
	3.703	7.5	16.071		3.703		16.071
	1.923	7.32	45		1.923		45
	19.02	14.488	7.826		19.02		7.826
	1.313	14.285	19.318		1.313	14.285	19.318
protein (mg)	0.2255	0.2505	0.067		0.0660	0.010	0.21
	0.2355	0.2595	0.067		0.0669		0.21
	0.33	0.2595	0.124		0.04635		0.205
	0.3525	0.1575	0.2445		0.2025		0.14
	0.1815	0.1785	0.23		0.195	0.0614	0.2

Table 27: Values of luminescence measurements ( $L_o$  and  $L_e$ ) and % of inhibition in V. fischeri exposed for 30 min to aqueous solutions of desnitro-IMI, olefin-IMI and 5-hydroxy-IMI.

## V. fischeri

#### desnitro-IMI

Geometric dillution

Concentration: 100 mg L<sup>-1</sup>

$mg L^{-1}$	G-value	Lo	Le			% Inhibition average	Valid
	K	349,10	323,10	318,2368	-1,5		1,53
		397,70	357,00	362,5402	1,5		Yes
0.39	512	370,30	325,10	337,5626	3,7		0,58
		363,30	315,10	331,1814	4,9	4,27	Yes
0.78	256	336,40	272,30	306,6596	11,2		0,58
		364,30	304,60	332,093	8,3	9,74	Yes
1.5625	128	344,00	299,30	313,5877	4,6		0,25
		345,00	298,60	314,4993	5,1	4,81	Yes
3.125	64	361,20	311,00	329,2671	5,5		0,58
		339,00	295,50	309,0297	4,4	4,96	Yes
6.25	32	355,10	308,30	323,7064	4,8		1,00
		316,10	268,70	288,1543	6,8	5,76	Yes
12.5	16	338,40	279,70	308,4828	9,3		0,05
		329,10	271,70	300,005	9,4	9,38	Yes
25	8	323,00	249,10	294,4443	15,4		0,41
		312,30	243,20	284,6902	14,6	14,99	Yes
50	4	302,10	220,90	275,392	19,8		0
		312,80	228,70	285,146	19,8	19,79	Yes
100	2	332,30	190,00	302,9221	37,3		0,68
		339,30	198,20	309,3032	35,9	36,60	Yes

#### olefin-IMI

Geometric dillution

Concentration: 100 mg  $L^{-1}$ 

$mg L^{-1}$	G-value	Lo	Le			% Inhibition average	Valid
	K	667,50	695,70	667,7769 -	-4,2		4,18
		719,50	689,70	719,7984	4,2		no
0.39	512	703,00	668,70	703,2916	4,9		2,66
		688,30	691,40	688,5855 -	-0,4	2,25	Yes
0.78	256	672,40	660,80	672,6789	1,8		0,36
		656,10	640,00	656,3721	2,5	2,13	Yes
1.5625	128	691,30	646,20	691,5867	6,6		0,18
		698,60	650,50	698,8898	6,9	6,74	Yes
3.125	64	735,80	699,20	736,1052	5,0		2,05
		716,10	651,10	716,397	9,1	7,06	Yes
6.25	32	728,40	667,00	728,7021 8	8,5		2,53
		730,40	631,90	730,7029	13,5	10,99	Yes
12.5	16	726,50	655,80	726,8013	9,8		2,73
		707,60	600,10	707,8935	15,2	12,50	Yes
25	8	761,30	601,20	761,6158	21,1		0,67
		689,80	554,00	690,0861	19,7	20,39	Yes
50	4	712,30	503,70	712,5954	29,3		1,58
		708,10	523,10	708,3937	26,2	27,74	Yes
100	2	690,00	388,10	690,2862	43,8		0,56
		701,40	386,70	701,6909	44,9	44,33	Yes

Table 27: Values of luminescence measurements ( $L_o$  and  $L_e$ ) and % of inhibition in V. fischeri exposed for 30 min to aqueous solutions of desnitro-IMI, olefin-IMI and 5-hydroxy-IMI (continued).

## V. fischeri

## 5-hydroxy-IMI

Geometric dillution Concentration: 100 mg/L

$mg L^{-1}$	G-value	Lo	Le	Mean	% Inhibition average	Valid
	) K	801,50	841,60	836,9317 -0,6	· ·	0,56
		825,90	857,60	862,4104 0,6		Yes
0.39	512	815,70	809,00	851,7595 5,0		0,98
		801,40	778,40	836,8273 7,0	6,00	Yes
0.78	3 256	831,70	803,40	868,4668 7,5		0,37
		841,90	819,70	879,1177 6,8	7,13	Yes
1.5625	5 128	839,60	829,10	876,716 5,4		1,10
		841,40	811,50	878,5956 7,6	6,53	Yes
3.125	5 64	864,10	829,60	902,2991 8,1		0,65
		855,00	809,20	892,7968 9,4	8,71	Yes
6.25	5 32	820,10	781,60	856,354 8,7		0,15
		856,30	813,40	894,1543 9,0	8,88	Yes
12.5	5 16	862,60	812,30	900,7328 9,8		0,31
		841,70	787,10	878,9088 10,4	10,13	Yes
25	5 8	824,50	769,10	860,9485 10,7		0,06
		861,60	802,60	899,6886 10,8	10,73	Yes
50	) 4	836,40	700,30	873,3746 19,8		0,28
		860,00	725,00	898,0178 19,3	19,54	Yes
100	2	876,20	598,10	914,934 34,6		0,09
		858,00	587,30	895,9294 34,4	34,54	Yes

Table 27a: Gamma values and gamma linear regression applied for EC-calculations in V. fischeri test.

## Olefin-IMI

Conc. mg/l	% Inhibition	Gamma	log Gama Y	log conc.	Gamma Lin. Reg	EC	Concentration mg/l
0,39	2,2549	0,02			0,02	20	22,577
0.78	2,1301	0.02			0.03	30	49.93
1,5625	6.7432	0.07			0.04	40	95.517
3,125	7.0641	0.08			0.07	50	173.217
6.25	10.9945	0.12	-0.908	0.796	0.10	60	314,351
12,50	12,4982	0.14	-0.845	1.097	0.17	70	601,678
25.00	20.3914	0.26	-0.592	1,398	0.27	80	1328.985
50.00	27,7358	0.38	-0.416	1.699	0.43		
100,00	44,3336	0.80	-0.099	2,00	0,69		

#### Desnitro-IMI

Conc. mg/l	% Inhibition	Gamma	log Gama Y	log conc.	Gamma Lin. Reg	EC	Concentration mg/l
0,39	4,2739	0.04			0,0046	20	41,563
0.78	9,7416	0.11			0.01	30	78,125
1.5625	4.8058	0.05			0.02	40	130,856
3,125	4,963	0.05			0.03	50	210,059
6.25	5.7554	0.06			0.05	60	337,396
12.50	9,3826	0.10			0.09	70	565,362
25.00	14,9869	0.18	-0.754	1.398	0.16	80	1061,635
50.00	19,7913	0.25	-0.608	1,699	0.29		
100,00	36,599	0.58	-0,239	2.00	0.53		

## 5-hydroxy-IMI

Conc. mg/l	% Inhibition	Gamma	log Gama Y	log conc. X	Gamma Lin. Reg	EC	Concentration mg/l
0,39	6,0011	0,06			0,01	20	46,01
0.78	7,1255	0.08			0.01	30	92,821
1,5625	6,5339	0,07			0.02	40	164,721
3,125	8,7103	0,10			0.03	50	278,822
6,25	8,8803	0,10			0.05	60	472,265
12,50	10,1318	0,11	-0,948	1.097	0.09	70	838,478
25,00	10,7298	0,12	-0.92	1,398	0,16	80	1689,669
50,00	19,5417	0.24	-0.615	1,699	0,27		
100,00	34,5386	0,53	-0,278	2,00	0,45		

Table 28: Values of fluorescence measurement and % of algal growth in D. subspicatus exposed for 0, 24, 48, 72 and 96 h to IMI, Confidor 200SL and 6CNA.

#### D. subspicatus

0 h						
Fluorescence measurments(FU) mg L <sup>-1</sup>	7.6	25.6	51.1	127.8	255.6	
positive control	431	371	387	387	408	
	408	434	374	425	362	
	375	383	374	376	338	
control	363	388	376	391	401	
	437 404	389 414	388 379	358 201	382 428	
	404	414	319	201	426	
imidacloprid	365	353	412	366	359	
	399 378	406 378	397 366	387 370	359 318	
# 1	200	255	2.10	255	400	
confidor 200SL	380 400	355 373	340 350	355 352	403 390	
	380	334	345	340	360	
6CNA	405	330	332	368	277	
	455	395	356	373	234	
Average	444	402	395	393	254	
positive control (mg L <sup>-1</sup> )	0	7.6	25.6	51.1	127.8	255.6
Mean	379.933	404.667	396	378.333 99.5789	396	369.333
Survival % SD	100	106.51 7.40878	104.229 8.80456	1.97549	104.229 6.76696	97.21 9.36257
imidacloprid						
Mean Survival %	379.933 100	380.667 100.193	379 99.7543	391.667 103.088	374.333 98.5261	345.333 90.8931
SD	0	4.51557	6.97863	6.17455	2.93485	6.2304
confidor 200SL Mean	379.933	386.667	354	345	349	384.333
Survival %	100	101.772	93.1742	90.8054	91.8582	101.158
SD 6CNA	0	3.03922	5.13754	1.31602	2.08912	5.80443
Mean	417.333	434.667	375.667	361	378	388.333
Survival % SD	100	104.153 6.29573	90.016 9.51351	86.5016 7.6189	90.5751 3.16983	93.0511 9.86899
3D	Ü	0.29373	9.31331	7.0189	3.10963	9.00099
24 h						
24 h  Fluorescence measurments(FU) mg L <sup>-1</sup>	7.6	25.6	51.1	127.8	255.6	
Fluorescence measurments(FU)	328	108	72	68	39	
Fluorescence measurments(FU) $mg L^{-1}$						
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control	328 521 204	108 131 83	72 71 73	68 95 61	39 29 39	
Fluorescence measurments(FU) $mg L^{-1}$	328 521	108 131	72 71	68 95	39 29	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control	328 521 204 579	108 131 83 647	72 71 73 546	68 95 61 593	39 29 39 530	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control	328 521 204 579 761	108 131 83 647 587	72 71 73 546 649	68 95 61 593 557	39 29 39 530 620	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control	328 521 204 579 761 567 465 485	108 131 83 647 587 543 435 478	72 71 73 546 649 551 427 458	68 95 61 593 557 104 548 417	39 29 39 530 620 488 445 438	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control	328 521 204 579 761 567	108 131 83 647 587 543	72 71 73 546 649 551	68 95 61 593 557 104	39 29 39 530 620 488	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control	328 521 204 579 761 567 465 485 478	108 131 83 647 587 543 435 478 433 541	72 71 73 546 649 551 427 458 439	68 95 61 593 557 104 548 417 456	39 29 39 530 620 488 445 438 387	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control  imidacloprid	328 521 204 579 761 567 465 485 478	108 131 83 647 587 543 435 478 433	72 71 73 546 649 551 427 458 439	68 95 61 593 557 104 548 417 456	39 29 39 530 620 488 445 438 387	
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL	328 521 204 579 761 567 465 485 478 647 529 513	108 131 83 647 587 543 435 478 433 541 486 435	72 71 73 546 649 551 427 458 439 400 404 412	68 95 61 593 557 104 548 417 456 415 414 375	39 29 39 530 620 488 445 438 387 418 393 358	
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control  imidacloprid	328 521 204 579 761 567 465 485 478	108 131 83 647 587 543 435 478 433 541 486	72 71 73 546 649 551 427 458 439	68 95 61 593 557 104 548 417 456	39 29 39 530 620 488 445 438 387 418 393	
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL  6CNA	328 521 204 579 761 567 465 485 478 647 529 513	108 131 83 647 587 543 435 478 433 541 486 435	72 71 73 546 649 551 427 458 439 400 404 412	68 95 61 593 557 104 548 417 456 415 414 375 2596	39 29 39 530 620 488 445 438 387 418 393 358	
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average	328 521 204 579 761 567 465 485 478 647 529 513	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800	39 29 39 530 620 488 445 438 387 418 393 358 341 40	255.6
Fluorescence measurments(FU)  mg L <sup>-1</sup> positive control  control  imidacloprid  confidor 200SL  6CNA  Average  positive control (mg L <sup>-1</sup> )  Mean	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72	39 29 39 530 620 488 445 438 387 418 393 358 318 41 40	255.6 35.6667
Fluorescence measurments(FU) mg L 1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L 1) Mean Survival %	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100	108 131 83 647 587 543 435 438 433 541 486 435 1391 962 805 7.6 351 63.266	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776	39 29 39 530 620 488 445 438 387 418 393 358 41 40 127.8 74.6667 13.4583	35.6667 6.42874
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD imidacloprid	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025	39 29 39 530 620 488 445 438 387 418 393 358 41 40 <i>127.8</i> 74.6667 13.4583 3.23606	35.6667 6.42874 1.04065
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L -1 ) Mean Survival % SD imidacloprid Mean	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025	39 29 39 530 620 488 445 438 387 418 393 358 41 40 127.8 74.6667 13.4583 3.23606 473.667	35.6667 6.42874 1.04065 423.333
Fluorescence measurments(FU) mg L <sup>-1</sup> positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD imidacloprid Mean Survival % SD imidacloprid Mean Survival % SD	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025	39 29 39 530 620 488 445 438 387 418 393 358 41 40 <i>127.8</i> 74.6667 13.4583 3.23606	35.6667 6.42874 1.04065
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L -1 ) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100 0	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936 476 85.7967 1.82929	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713 448.667 80.87 4.58239	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025 441.333 79.5482 2.81744	39 29 39 530 620 488 445 438 387 418 393 358 41 40 127.8 74.6667 13.4583 3.23606 473.667 85.3761 12.1239	35.6667 6.42874 1.04065 423.333 76.3038 5.7065
Fluorescence measurments(FU) mg L 1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival %	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100 0	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936 476 85.7967 1.82929 563 101.478	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713 448.667 80.87 4.58239	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025 441.333 79.5482	39 29 39 530 620 488 445 438 387 418 393 358 38 41 40 127.8 74.6667 13.4583 3.23606 473.667 85.3761 12.1239 401.333 72.3384	35.6667 6.42874 1.04065 423.333 76.3038
Fluorescence measurments(FU) mg L -1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L -1 ) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100 0	108 131 83 647 587 543 435 435 438 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936 476 85.7967 1.82929	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713 448.667 80.87 4.58239	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025 441.333 79.5482 2.81744	39 29 39 530 620 488 445 438 387 418 393 358 41 40 127.8 74.6667 13.4583 3.23606 473.667 85.3761 12.1239	35.6667 6.42874 1.04065 423.333 76.3038 5.7065 389.667
Fluorescence measurments(FU) mg L 1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD 6CNA Mean	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100 0 554.8 100 0	108 131 83 647 587 543 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936 476 85.7967 1.82929 563 101.478 13.1912	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713 448.667 80.87 4.58239 487.333 87.8395 9.55526	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025 441.333 79.5482 2.81744 405.333 73.0594 1.10132 1458.33	39 29 39 530 620 488 445 438 387 418 393 358 38 41 40 127.8 74.6667 13.4583 3.23606 473.667 85.3761 12.1239 401.333 72.3384 4.11154	35.6667 6.42874 1.04065 423.333 76.3038 5.7065 389.667 70.2355 5.43233 39.6667
Fluorescence measurments(FU) mg L 1 positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD 6CNA	328 521 204 579 761 567 465 485 478 647 529 513 1060 896 794 control 554.8 100 0 554.8 100 0	108 131 83 647 587 543 435 478 433 541 486 435 1391 962 805 7.6 351 63.266 28.7936 476 85.7967 1.82929 563 101.478 13.1912	72 71 73 546 649 551 427 458 439 400 404 412 2055 1396 924 25.6 107.333 19.3463 4.32713 448.667 80.87 4.58239 487.333 87.8395 9.55526	68 95 61 593 557 104 548 417 456 415 414 375 2596 1800 998 51.1 72 12.9776 0.18025 441.333 79.5482 2.81744 405.333 73.0594 1.10132	39 29 39 530 620 488 445 438 387 418 393 358 41 40 127.8 74.6667 13.4583 3.23606 473.667 85.3761 12.1239 401.333 72.3384 4.11154	35.6667 6.42874 1.04065 423.333 76.3038 5.7065 389.667 70.2355 5.43233

Table 28: Values of fluorescence measurement and % of algal growth in D. subspicatus exposed for 0, 24, 48, 72 and 96 h to IMI, Confidor 200SL and 6CNA (continued).

#### D. subspicatus

48 h						
Fluorescence measurments(FU)	7.6	25.6	51.1	127.8	255.6	
mg/L positive control	173	25.0	24	127.8	233.0	
positive control	892	52	30	21	9	
	109	35	23	18	8	
	1024	1226	1217	1202	1004	
control	1234 1557	1326 1401	1317 1676	1283 1320	1084 1286	
	1273	1583	1413	468	1403	
imidacloprid	1665	1441	1408	1589	1481	
	1731	1797	1623	1495	1522	
	1710	1676	1661	1565	1652	
confidor 200SL	1919	1540	1244	1165	1137	
	1730	1550	1114	1090	1097	
	1836	1440	1145	959	981	
6CNA	3747	3659	5030	4487	20	
	4053	4669	5111	4173	15	
	4894	4118	5320	4365	20	
Average						
positive control (mg L <sup>-1</sup> )	0	7.6	25.6	51.1	127.8	255.6
Mean Survival %	981.185 100.000	93.980 9.578	30.485 3.107	19.235 1.960	14.485 1.476	7.485 0.763
SD	66.671	8.918	2.230	1.348	0.996	0.558
imidacloprid						
Mean	981.185	1,276.485	1,228.485	1,172.985	1,162.235	1,163.700
Survival % SD	100.000 66.671	130.096 3.437	125.204 18.449	119.548 13.905	118.452 4.978	118.605 9.099
confidor 200SL	00.071	3.137	10,	15.505	, 70	2.022
Mean	981.185	1,371.235	1,132.485	875.735	803.485	803.735
Survival % SD	100.000	139.753	115.420	89.253	81.889	81.915
6CNA	66.671	9.655	6.199	6.920	10.626	8.258
Mean	2,920.133	4,231.333	4,148.667	5,153.667	4,341.667	18.333
Survival %	100.000	144.902	142.071	176.487	148.680	0.628
SD	0.000	20.339	17.318	5.124	5.421	0.099
72 h						
72 h						
72 h  Fluorescence measurments(FU)						
Fluorescence measurments(FU) mg/L	7.6	25.6	51.1	127.8	255.6	
Fluorescence measurments(FU)	153	24	17	14	13	
Fluorescence measurments(FU) mg/L						
Fluorescence measurments(FU) mg/L	153 1586	24 28	17 21	14 15	13 13 7	
Fluorescence measurments(FU) mg/L	153 1586 72 4087	24 28 29 4524	17 21 15 4081	14 15 12 3834	13 13 7 3535	
Fluorescence measurments(FU) mg/L positive control	153 1586 72 4087 4255	24 28 29 4524 3708	17 21 15 4081 6175	14 15 12 3834 3267	13 13 7 3535 3307	
Fluorescence measurments(FU) mg/L positive control	153 1586 72 4087	24 28 29 4524	17 21 15 4081	14 15 12 3834	13 13 7 3535	
Fluorescence measurments(FU) mg/L positive control	153 1586 72 4087 4255 3417 5338	24 28 29 4524 3708 3702	17 21 15 4081 6175 3596 4148	14 15 12 3834 3267 2120 4757	13 13 7 3535 3307 3575 4277	
Fluorescence measurments(FU) mg/L positive control  control	153 1586 72 4087 4255 3417 5338 5047	24 28 29 4524 3708 3702 4761 5329	17 21 15 4081 6175 3596 4148 5368	14 15 12 3834 3267 2120 4757 4503	13 13 7 3535 3307 3575 4277 4779	
Fluorescence measurments(FU) mg/L positive control  control	153 1586 72 4087 4255 3417 5338	24 28 29 4524 3708 3702	17 21 15 4081 6175 3596 4148	14 15 12 3834 3267 2120 4757	13 13 7 3535 3307 3575 4277	
Fluorescence measurments(FU) mg/L positive control  control	153 1586 72 4087 4255 3417 5338 5047	24 28 29 4524 3708 3702 4761 5329	17 21 15 4081 6175 3596 4148 5368	14 15 12 3834 3267 2120 4757 4503	13 13 7 3535 3307 3575 4277 4779	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid	153 1586 72 4087 4255 3417 5338 5047 6404	24 28 29 4524 3708 3702 4761 5329 7413	17 21 15 4081 6175 3596 4148 5368 6117 2621	14 15 12 3834 3267 2120 4757 4503 5556 2356	13 13 7 3535 3307 3575 4277 4779 5624 2018	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844	
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14	255.6
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1 )	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18	255.6 11.000
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14	255.6 11.000 0.289
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000	14 15 12 3834 3267 2120 4757 4503 5556 2198 2219 17 11 26	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18	11.000
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD imidacloprid	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18 127.8 13.667 0.358 0.040	11.000 0.289 0.091
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1 ) Mean Survival % SD imidacloprid Mean	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5,596.333	17 21 15 4081 6175 3596 4148 5368 6117 2621 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5,211.000	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18 127.8 13.667 0.358 0.040	11.000 0.289 0.091 4,893.333
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD imidacloprid	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18 127.8 13.667 0.358 0.040	11.000 0.289 0.091
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L <sup>-1</sup> ) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5,596.333 146.801 18.741	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5,834.333 153.044 36.628	14 15 12 3834 3267 2120 4757 4503 5556 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5.211.000 136.693 26.070	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18 127.8 13.667 0.358 0.040 4,938.667 129.549 14.414	11.000 0.289 0.091 4,893.333 128.360 17.857
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5.596.333 146.801 18.741 5,164.667	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5,834.333 153.044 36.628	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5,211.000 136.693 26.070 2,728.667	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14 18 127.8 13.667 0.358 0.040 4,938.667 129.549 14.414	11.000 0.289 0.091 4,893.333 128.360 17.857
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival %	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5.596.333 146.801 18.741 5,164.667 135.477	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5.834.333 153.044 36.628	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5,211.000 136.693 26.070 2,728.667 71.577	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 18 127.8 13.667 0.358 0.040 4,938.667 129.549 14.414	11.000 0.289 0.091 4,893.333 128.360 17.857 1,934.667 50.749
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5.596.333 146.801 18.741 5,164.667	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5,834.333 153.044 36.628	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5,211.000 136.693 26.070 2,728.667	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14 18 127.8 13.667 0.358 0.040 4,938.667 129.549 14.414	11.000 0.289 0.091 4,893.333 128.360 17.857
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L - 1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD 6CNA Mean	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5,596.333 146.801 18.741 5,164.667 135.477 7,552	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5,834.333 153.044 36.628 4,215.667 110.584 7.752	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 11 26 51.1 17.667 0.463 0.080 5,211.000 136.693 26.070 2,728.667 71.577 4.080	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14 18 127.8 13.667 0.358 0.040 4,938.667 129.549 14.414 2,257.667 59.222 2.251	11.000 0.289 0.091 4,893.333 128.360 17.857 1,934.667 50.749 2.288
Fluorescence measurments(FU) mg/L positive control  control  imidacloprid  confidor 200SL  6CNA  Average positive control (mg L-1) Mean Survival % SD imidacloprid Mean Survival % SD confidor 200SL Mean Survival % SD confidor 200SL Mean Survival % SD 6CNA	153 1586 72 4087 4255 3417 5338 5047 6404 4991 5006 5497 5039 4055 6057 control 3,812.200 100.000 0.000 3,812.200 100.000 0.000	24 28 29 4524 3708 3702 4761 5329 7413 3921 4214 4512 145 357 645 7.6 112.500 2.951 1.502 5.596.333 146.801 18.741 5,164.667 135.477 7,552	17 21 15 4081 6175 3596 4148 5368 6117 2621 2658 2907 29 39 182 25.6 27.000 0.708 0.069 5.834.333 153.044 36.628 4,215.667 110.584 7.752	14 15 12 3834 3267 2120 4757 4503 5556 2356 2198 2219 17 111 26 51.1 17.667 0.463 0.080 5,211.000 136.693 26.070 2,728.667 71.577 4.080	13 13 7 3535 3307 3575 4277 4779 5624 2018 1942 1844 14 14 18 127.8 13.667 0.358 0.040 4.938.667 129.549 14.414 2,257.667 59,222 2,251	11.000 0.289 0.091 4,893.333 128.360 17.857 1,934.667 50.749 2.288

Table 28: Values of fluorescence measurement and % of algal growth in D. subspicatus exposed for 0, 24, 48, 72 and 96 h to IMI, Confidor 200SL and 6CNA (continued)

## D. subspicatus

## 96 h

Fluorescence measurments(F	TU)					
mg/L	7.6	25.6	51.1	127.8	255.6	
positive control	462	21	20	12	12	
	3866	21	21	18	15	
	203	27	20	10	9	
control	6471	6929	7019	7014	6887	
	7933	7510	9929	7681	7720	
	7864	8701	8095	6062	8140	
imidacloprid	9901	10912	10546	10565	8296	
	12089	12528	10834	9157	8494	
	11920	10684	10770	10443	8183	
confidor 200SL	10026	8694	6207	4321	3388	
	9711	9875	6780	5642	3576	
	9623	9501	7800	6471	4497	
6CNA	10620	10781	12295	10421	17	
	10452	12392	12239	9493	12	
	12070	11402	12322	10662	16	
Average						
<b>positive control</b> $(mg L^{-1})$	0	7.6	25.6	51.1	127.8	255.6
Mean	7597	332.5	23	20.3333	13.3333	12
Survival %	100	4.37673	0.30275	0.26765	0.17551	0.15796
SD	0	2.4107	0.0456	0.0076	0.0548	0.03949
imidacloprid						
Mean	7597	11303.3	11374.7	10716.7	10055	8324.33
Survival %	100	148.787	149.726	141.064	132.355	109.574
SD	0	16.0246	13.2329	1.9906	10.2683	2.07219
confidor 200SL						
Mean	7597	9786.67	9356.67	6929	5478	3820.33
Survival %	100	128.823	123.163	91.2071	72.1074	50.2874
SD	0	2.7891	7.94504	10.6211	14.2733	7.81232
6CNA						
Mean	9526.73	11047.3	11525	12285.3	10192	15
Survival %	100	115.961	120.975	128.956	106.983	0.15745
SD	0	9.33825	8.52877	0.44439	6.47891	0.02777
DMSO + NMP						
% v/v	0	0.003	0.01	0.02	0.06	0.12
Mean	7597	6176.31	6054.8	4277.11	1344.66	774.8
Survival %	100	81.3	79.7	56.3	17.7	10.3
SD	0	4.56	6.23	7.32	3.21	6.02

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C).

### G. fossarum

$\mathbf{A}$								
control			102.2 μg L <sup>-1</sup>			153.3 μg L <sup>-1</sup>		
Wet weight		Length	Wet weight		Length	Wet weight		Length
0.017	†	7	0.016	†	5	0.038	†	10
0.026	†	10	0.034	†	11	0.031		12
0.017		9	0.036	†	12	0.054	†	12
0.038		12	0.037	†	12	0.054		13
0.052		14	0.028	†	12	0.029		8
0.052		15	0.342	†	13	0.032		10
0.054		15	0.037	†	12	0.017		10
0.036		19	0.023	†	14	0.043		16
0.018		16	0.018	†	10	0.032		14
0.044		14	0.068	†	14	0.027		13
0.019		10	0.026	†	9	0.017		11
0.032		12	0.018		9	0.023		11
0.032		10 7	0.053		16	0.054		12
0.038 0.043		14	0.053 0.024		15 12	0.03 0.061		11 16
0.043		10	0.024		12	0.049		12
0.051		14	0.027		14	0.049		12
0.052		15	0.034		13	0.054		13
0.046		17	0.047		14	0.029		8
0.024		10	0.052		10	0.045		17
0.047		15	0.08		16	0.043		16
0.023		11	0.023		10	0.032		15
0.026		11	0.018		10	0.038		16
0.035		13	0.056		17	0.037		13
0.041		15	0.029		11	0.038		11
0.023		11	0.029		11	0.023		12
0.03		14	0.023		11	0.026		12
0.044		15	0.017		11	0.024		12
0.031		13	0.069		16	0.031		13
0.029		12	0.058		16	0.028		12
0.022		10	0.034		14	0.022		10
0.032		15	0.037		13	0.036		15
0.047		16	0.037		14	0.021		12
0.039		14	0.013		10	0.024		12
0.023		11	0.045		15	0.022		12
0.03		14	0.034		11	0.025		12
0.019		11	0.036		13	0.04		15
0.023		12	0.033		14	0.038		12
0.043		16	0.018		12	0.038		16
0.038 0.04		15	0.026 0.028		13 14	0.032		12 14
0.027		17 12	0.028		14 16	0.028 0.03		
0.027		15	0.043		14	0.03		14 15
0.024		11	0.022		14	0.028		10
0.028		13	0.027		15	0.04		15
0.032		13	0.027		16	0.038		11
0.036		13	0.037		16	0.027		13
0.032		17	0.037		16	0.043		14
0.043		17	0.022		10	0.032		12
0.042		15	0.043		17	0.042		15
0.032		14	0.041		15	0.034		11
0.036		11	0.032		13	0.024		13
0.023		12	0.021		12	0.016		13

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C) (continued).

## G. fossarum

<b>A</b>							
A			1				
$204.5~\mu \mathrm{g~L^{-1}}$			255.6 μg L <sup>-1</sup>			511.3 μg L <sup>-1</sup>	
Wet weight		Length	Wet weight		Length	Wet weight	Length
0.05	<i>†</i>	13	0.023	†	6	0.038 †	15
0.034	<i>†</i>	11	0.025	†	6	0.03 †	11
0.048	<i>†</i>	15	0.033	†	11	0.032 †	15
0.06	†	17	0.07	<i>†</i>	17	0.038 †	14
0.025	<i>†</i>	12	0.041	†	15	0.032 †	15
0.018	<i>†</i>	11	0.033	<i>†</i>	16	0.032 †	15
0.041	<i>†</i>	15	0.025		13	0.038 †	15
0.038	<i>†</i>	15	0.04		14	0.245	11
0.07		17	0.022		12	0.038	14
0.026		14	0.053		15	0.034	15
0.061		15	0.039		13	0.057	16
0.056		14	0.027		10	0.022	11
0.05		12	0.052		13	0.043	17
0.032		9	0.056		13	0.036	10
0.038		12	0.041		15	0.027	10
0.055		15	0.049		18	0.08	18
0.048		15	0.041		15	0.036	10
0.039		14	0.045		16	0.025	8
0.024		13	0.039		16	0.03	15
0.049		16	0.037		13	0.043	15
0.019		12	0.039		13	0.044	15
0.049		18	0.349		15	0.041	14
0.034		12	0.048		10	0.027	14
0.425		14	0.048		14	0.036	15
0.035		13	0.04		15	0.036	10
0.041		13	0.022		12	0.019	12
0.029		13	0.034		15	0.026	14
0.042		17	0.02		10	0.021	12
0.045		15	0.034		15	0.044	15
0.033		15	0.042		17	0.041	14
0.025		14	0.043		15	0.027	14
0.021		12	0.04		15	0.045	16
0.032		14	0.044		18	0.043	17
0.044		15	0.021		12	0.021	12
0.029		13	0.048		16	0.019	12
0.042		17	0.041		16	0.026	14
0.025		12	0.02		10	0.042	15
0.031		12	0.046		15	0.043	16
0.026		12	0.032		13	0.044	14
0.026		12	0.024		10	0.042	15
0.028		12	0.032		16	0.039	13
0.045		12	0.036		16	0.036	11
0.034		13	0.023		14	0.041	13
0.036		15	0.032		13	0.023	13
0.033		17	0.028		14	0.026	13
0.03		12	0.023		17	0.024	17
0.019		10	0.048		15	0.022	15
0.023		10	0.042		16	0.032	15
0.043		17	0.028		12	0.014	5
0.022		11	0.027		12	0.036	10
0.036		11	0.037		12	0.025	8
0.039		15	0.019		11	0.022	7

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C) (continued).

## G. fossarum

B

control / DMSO+ NMP			$102.2~\mu { m g~L}^{-1}$			$153.3~\mu g~L^{-1}$
Wet weight		Length	Wet weight		Length	Wet weight
0.017	†	7	0.019	†	7	0.015
0.026	†	10	0.026	†	10	0.024
0.017		9	0.017		9	0.038
0.038		12	0.039		12	0.037
0.052		14	0.052		14	0.029
0.052		15	0.042		15	0.342
0.054		15	0.054		15	0.035
0.036		19	0.026		19	0.023
0.018		16	0.018		16	0.018
0.044		14	0.054		14	0.067
0.019		10	0.019		10	0.026
0.032		12	0.032		12	0.018
0.032		10	0.032		10	0.053
0.038		7	0.038		7	0.053
0.043		14	0.033		14	0.031
0.031		10	0.035		11	0.0542
0.052		14	0.04		14	0.0292
0.052		15	0.05		14	0.0318
0.046		17	0.036		17	0.0503
0.024		10	0.028		12	0.0345
0.047		15	0.047		15	0.08
0.023		11	0.013		10	0.023
0.026		11	0.026		11	0.018
0.035		13	0.035		12	0.056
0.041		15	0.041		15	0.029
0.023		11	0.033		11	0.029
0.023		14	0.033		13	0.029
0.044		15	0.044		13	0.03
0.031		13	0.021		13	0.032
0.029		12	0.021		12	0.032
0.022		10	0.022		10	0.032
0.032		15	0.022		15	0.032
0.032		16	0.032		16	0.032
0.039		14	0.039		14	0.038
0.023		11	0.023		11	0.015
0.023		14	0.023		14	0.034
0.019		11	0.027		10	0.034
0.023		12	0.052		14	0.034
0.023		16	0.056		13	0.033
0.043		15	0.030		15	0.028
0.038		17	0.039		18	0.028
0.027		12	0.039		15	0.028
0.027		15	0.041		13	0.042
0.024		11	0.024		11	0.038
		13	0.024		13	
0.028						0.027
0.032 0.036		13 13	0.035 0.036		13 13	0.039 0.037
0.032		17	0.032		15	0.033
0.043		17	0.023		13	0.042
0.042		15	0.042		15	0.043
0.032		14	0.042		14	0.031
0.036		11	0.036		11	0.032
0.023		12	0.023		12	0.021

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C) (continued).

## G. fossarum B

Wet weight	В								
Wet weight	204.5 $μg L^{-1}$			255.6 μg L <sup>-1</sup>			511.3 μg L <sup>-1</sup>		
0.037			Length	Wet weight		Length	Wet weight		Length
0.044	0.038	†	10	0.023	<i>†</i>	6	0.023	†	6
0.054   12	0.037	†	12	0.025	<i>†</i>	6	0.025	†	6
0.022         †         8         0.019         †         7         0.044         †         15           0.022         †         10         0.026         †         11         0.033         †         16           0.019         †         10         0.039         †         12         0.04         †         15           0.036         †         14         0.016         †         5         0.023         †         12           0.017         †         13         0.034         †         11         0.053         †         15           0.017         †         11         0.034         †         11         0.053         †         15           0.017         †         11         0.037         †         12         0.019         †         11           0.023         111         0.037         †         12         0.019         †         11           0.03         11         0.038         †         10         0.028         †         12           0.04         12         0.054         †         12         0.043         †         15           0.045         12	0.044	†	12	0.033	†	11	0.033	†	12
0.022         †         10         0.026         †         10         0.033         †         16           0.013         †         16         0.037         †         9         0.026         †         14           0.036         †         14         0.016         †         5         0.023         †         12           0.027         †         13         0.034         †         11         0.053         †         15           0.017         †         11         0.036         †         12         0.036         †         13           0.023         11         0.037         †         12         0.019         †         11           0.054         12         0.038         †         10         0.028         †         12           0.03         11         0.031         †         12         0.043         †         15           0.04         12         0.054         †         13         0.043         †         15           0.049         12         0.054         †         13         0.044         †         17           0.031         12         0.054         †	0.054	†	13	0.07				<i>†</i>	15
0.019         †         10         0.017         †         9         0.026         †         14           0.036         †         16         0.039         †         12         0.04         †         15           0.026         †         14         0.016         †         5         0.023         †         12           0.017         †         13         0.034         †         11         0.053         †         15           0.017         †         11         0.036         †         12         0.036         †         13           0.023         11         0.037         †         12         0.049         †         11           0.034         12         0.043         †         12         0.043         †         16           0.040         16         0.054         †         12         0.043         †         16           0.049         12         0.054         †         13         0.047         †         17           0.031         12         0.0654         †         13         0.027         †         12           0.041         †         13         0.027 <td>0.039</td> <td>†</td> <td>8</td> <td>0.019</td> <td></td> <td></td> <td></td> <td><i>†</i></td> <td>15</td>	0.039	†	8	0.019				<i>†</i>	15
0.043         †         16         0.039         †         12         0.04         †         15           0.027         †         13         0.034         †         11         0.053         †         12           0.017         †         11         0.036         †         12         0.036         †         13           0.023         111         0.037         †         12         0.019         †         11           0.054         12         0.038         †         10         0.028         †         12           0.03         11         0.031         †         12         0.043         †         15           0.049         12         0.054         †         13         0.04         †         17           0.031         12         0.064         †         13         0.027         †         12           0.043         13         0.026         †         13         0.027         †         12           0.029         8         0.0173         †         15         0.014         †         15           0.029         8         0.0173         †         15         0.0	0.022		10	0.026			0.033		16
0.036         †         14         0.016         †         5         0.023         †         12           0.017         †         13         0.034         †         11         0.053         †         15           0.017         †         11         0.036         †         12         0.019         †         11           0.054         12         0.038         †         12         0.049         †         11           0.061         16         0.054         †         12         0.043         †         16           0.061         16         0.054         †         12         0.038         †         15           0.049         12         0.064         †         13         0.04         †         17           0.031         12         0.0169         †         13         0.027         †         12           0.044         13         0.026         †         13         0.027         †         12           0.029         8         0.0173         †         15         0.014         †         15           0.043         16         0.023         †         17         0.									
0.027         †         13         0.036         †         11         0.036         †         12         0.036         †         13           0.023         111         0.037         †         12         0.019         †         11           0.054         12         0.038         †         10         0.028         †         12           0.03         11         0.031         †         12         0.043         †         16           0.061         16         0.054         †         12         0.043         †         16           0.049         12         0.054         †         13         0.04         †         17           0.031         12         0.054         †         13         0.027         †         12           0.054         13         0.026         †         11         0.041         †         15           0.029         8         0.0173         †         15         0.014         †         5           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †									
0.017         †         11         0.036         †         12         0.019         †         13           0.023         11         0.037         †         12         0.019         †         11           0.054         12         0.038         †         10         0.028         †         12           0.03         11         0.031         †         12         0.043         †         16           0.040         12         0.054         †         13         0.04         †         17           0.031         12         0.0169         †         13         0.027         †         12           0.049         12         0.054         †         13         0.027         †         12           0.054         13         0.026         †         11         0.041         †         5           0.029         8         0.0173         †         15         0.014         †         5           0.045         17         0.038         †         17         0.038         †         17         0.036         †         10           0.042         1         7         0.038         †									
0.023									
0.054         12         0.038         † 10         0.028         † 12           0.061         16         0.054         † 12         0.043         † 15           0.049         12         0.054         † 13         0.04         † 17           0.031         12         0.0169         † 13         0.04         † 17           0.054         13         0.026         † 11         0.041         † 15           0.029         8         0.0173         † 15         0.014         † 5           0.045         17         0.038         † 17         0.036         † 10           0.0443         16         0.037         † 12         0.025         † 8           0.032         15         0.023         † 14         0.022         † 7           0.038         16         0.023         † 14         0.022         † 7           0.037         13         0.068         † 14         0.022         † 7           0.038         16         0.018         † 9         0.041         † 13           0.023         12         0.017         10         0.034         † 15           0.024         12         0.017		†							
0.03         11         0.031         †         12         0.043         †         16           0.061         16         0.054         †         12         0.038         †         15           0.049         12         0.054         †         13         0.027         †         12           0.054         13         0.026         †         11         0.041         †         15           0.029         8         0.0173         †         15         0.041         †         5           0.043         16         0.038         †         17         0.036         †         10           0.043         16         0.023         †         14         0.025         †         8           0.032         15         0.023         †         14         0.039         †         15           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.026         †         9         0.036         †         11           0.023         12         0.018         †         10         0.039         †         13									
0.061         16         0.054         †         12         0.038         †         15           0.049         12         0.054         †         13         0.04         †         17           0.031         12         0.0169         †         13         0.027         †         12           0.054         13         0.026         †         11         0.041         †         15           0.029         8         0.0173         †         15         0.014         †         5           0.045         17         0.038         †         17         0.036         †         10         0.014         †         5           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0									
0.049         12         0.054         †         13         0.04         †         17           0.054         13         0.026         †         11         0.041         †         15           0.029         8         0.0173         †         15         0.014         †         5           0.045         17         0.038         †         17         0.036         †         10           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.041         †         15           0.037         13         0.068         †         14         0.039         †         13           0.023         12         0.018         9         0.041         †         13           0.									
0.031         12         0.0169         †         13         0.027         †         12           0.029         8         0.0173         †         15         0.014         †         15           0.045         17         0.038         †         17         0.036         †         10           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.034         †         13           0.022         12         0.017         10         0.034         †         15 <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></td<>									
0.054         13         0.026         †         11         0.041         †         15           0.029         8         0.0173         †         15         0.014         †         5           0.045         17         0.038         †         17         0.036         †         10           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.066         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13									
0.029         8         0.0173         †         15         0.014         †         5           0.045         17         0.038         †         17         0.036         †         10           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13									
0.045         17         0.038         †         17         0.036         †         10           0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.041         †         13           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.029         10         0.017         11         0.043         15 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.043         16         0.037         †         12         0.025         †         8           0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.022         10         0.017         11         0.043         15           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021									
0.032         15         0.023         †         14         0.022         †         7           0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.041         †         13           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.023         11         0.04         15         0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022									
0.038         16         0.018         †         10         0.042         †         15           0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.028         12         0.027         13         0.042         17           0.029         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15									
0.037         13         0.068         †         14         0.039         †         13           0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.025         12         0.0362         14         0.041         16           0.025         12         0.0362         14         0.041         16 </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.038         11         0.026         †         9         0.036         †         11           0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.022         12           0.024         12         0.0159         15         0.022         12           0.025         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038<									
0.023         12         0.018         9         0.041         †         13           0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         16         0.026         12         0.034         13           0.032         12									
0.026         12         0.017         10         0.034         †         15           0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.026         12         0.034         13           0.038         14         0.028					Ŧ				
0.024         12         0.043         16         0.026         †         10           0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         12         0.026         12         0.034         13           0.032         12         0.026         12         0.034         13           0.032         12         0.026         12									
0.031         13         0.032         14         0.034         15           0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.028         12         0.032         16           0.03         14         0.045         12         0.036									
0.028         12         0.027         13         0.042         17           0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.026         12         0.024         10           0.028         14         0.028         12         0.032         16           0.047         15         0.034         13         0.028         14           0.028         10         0.036         15         0.032								1	
0.022         10         0.017         11         0.043         15           0.036         15         0.023         11         0.04         15           0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.026         12         0.034         13           0.032         14         0.028         12         0.032         16           0.03         14         0.045         12         0.036         16           0.047         15         0.034         13         0.028         14           0.028         10         0.036         15         0.032									
0.036       15       0.023       11       0.04       15         0.021       12       0.0541       15       0.044       18         0.024       12       0.0159       15       0.022       12         0.022       12       0.034       17       0.048       16         0.025       12       0.0362       14       0.041       16         0.04       15       0.018       15       0.02       10         0.038       12       0.0676       14       0.046       15         0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.034       13         0.038       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17 </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.021         12         0.0541         15         0.044         18           0.024         12         0.0159         15         0.022         12           0.022         12         0.034         17         0.048         16           0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.026         12         0.024         10           0.028         14         0.028         12         0.032         16           0.03         14         0.045         12         0.036         16           0.047         15         0.034         13         0.028         14           0.028         10         0.036         15         0.032         13           0.04         15         0.033         17         0.028         14           0.038         11         0.03         12         0.023									
0.024       12       0.0159       15       0.022       12         0.022       12       0.034       17       0.048       16         0.025       12       0.0362       14       0.041       16         0.04       15       0.018       15       0.02       10         0.038       12       0.0676       14       0.046       15         0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16 </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.022       12       0.034       17       0.048       16         0.025       12       0.0362       14       0.041       16         0.04       15       0.018       15       0.02       10         0.038       12       0.0676       14       0.046       15         0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.042       13       0.043       17       0.028       12 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.025         12         0.0362         14         0.041         16           0.04         15         0.018         15         0.02         10           0.038         12         0.0676         14         0.046         15           0.038         16         0.026         12         0.034         13           0.032         12         0.026         12         0.024         10           0.028         14         0.028         12         0.032         16           0.03         14         0.045         12         0.036         16           0.047         15         0.034         13         0.028         14           0.028         10         0.036         15         0.032         13           0.04         15         0.033         17         0.028         14           0.038         11         0.03         12         0.023         17           0.027         13         0.019         10         0.048         15           0.043         14         0.023         10         0.042         16           0.032         12         0.043         17         0.028									
0.04       15       0.018       15       0.02       10         0.038       12       0.0676       14       0.046       15         0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.038       12       0.0676       14       0.046       15         0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.042       15       0.022       11       0.037       12 </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.038       16       0.026       12       0.034       13         0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>									
0.032       12       0.026       12       0.024       10         0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11									
0.028       14       0.028       12       0.032       16         0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11									
0.03       14       0.045       12       0.036       16         0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11									
0.047       15       0.034       13       0.028       14         0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11									
0.028       10       0.036       15       0.032       13         0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11									14
0.04       15       0.033       17       0.028       14         0.038       11       0.03       12       0.023       17         0.027       13       0.019       10       0.048       15         0.043       14       0.023       10       0.042       16         0.032       12       0.043       17       0.028       12         0.042       15       0.022       11       0.028       12         0.034       11       0.036       11       0.037       12         0.024       13       0.039       15       0.019       11			10	0.036		15	0.032		13
0.027     13     0.019     10     0.048     15       0.043     14     0.023     10     0.042     16       0.032     12     0.043     17     0.028     12       0.042     15     0.022     11     0.028     12       0.034     11     0.036     11     0.037     12       0.024     13     0.039     15     0.019     11	0.04		15	0.033		17			14
0.043     14     0.023     10     0.042     16       0.032     12     0.043     17     0.028     12       0.042     15     0.022     11     0.028     12       0.034     11     0.036     11     0.037     12       0.024     13     0.039     15     0.019     11	0.038		11	0.03		12	0.023		17
0.032     12     0.043     17     0.028     12       0.042     15     0.022     11     0.028     12       0.034     11     0.036     11     0.037     12       0.024     13     0.039     15     0.019     11	0.027		13	0.019		10	0.048		15
0.042     15     0.022     11     0.028     12       0.034     11     0.036     11     0.037     12       0.024     13     0.039     15     0.019     11	0.043		14	0.023		10	0.042		16
0.034     11     0.036     11     0.037     12       0.024     13     0.039     15     0.019     11	0.032		12	0.043		17	0.028		12
0.024 13 0.039 15 0.019 11	0.042			0.022		11	0.028		
	0.034		11	0.036		11	0.037		12
0.016 13 0.032 12 0.036 12				0.039					
0.010 15 0.032 12 0.030 12	0.016		13	0.032		12	0.036		12

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C) (continued).

# G. fossarum C

C								
control			102.2 μg $L^{-1}$			153.3 μg L <sup>-1</sup>		
Wet weight		Length	Wet weight		Length	Wet weight		Length
0.017	†	7	0.08	<del>†</del>	18	0.017	†	8
0.026	<i>†</i>	10	0.036	<del>†</del>	10	0.023	†	12
0.017		9	0.025	<del>†</del>	8	0.038	†	13
0.038		12	0.03	†	15	0.037	†	12
0.052		14	0.043		15	0.032		11
0.052		15	0.044		15	0.42		13
0.054		15	0.245		11	0.025		12
0.036		19	0.038		14	0.036		14
0.018		16	0.034		15	0.018		12
0.044		14	0.057		16	0.067		12
0.019		10	0.022		11	0.046		9
0.032		12	0.043		17	0.018		10
0.032		10	0.037		10	0.043		15
0.038		7	0.027		10	0.053		15
0.043		14	0.039		16	0.021		14
0.031		10	0.037		13	0.042		15
0.052		14	0.039		13	0.029		14
0.052		15	0.349		15	0.038		12
0.046		17	0.048		10	0.03		16
0.024		10	0.048		14	0.035		11
0.047		15	0.04		15	0.08		16
0.023		11	0.041		14	0.02		10
0.026		11	0.027		14	0.018		10
0.035		13	0.036		15	0.056		17
0.041		15	0.036		10	0.039		11
0.023		11	0.019		12	0.029		12
0.03		14	0.026		14	0.018		15
0.044		15	0.021		12	0.03		11
0.031		13	0.044		15	0.032		13
0.029		12	0.041		14	0.028		12
0.022		10	0.027		14	0.032		15
0.032		15	0.045		16	0.032		15
0.047		16	0.043		17	0.018		14
0.039		14	0.021		12	0.023		10
0.023		11	0.01		12	0.045		15
0.03		14	0.026		14	0.034		11
0.019		11	0.042		15	0.034		13
0.023		12	0.043		16	0.033		14
0.043		16	0.044		14	0.028		12
0.038		15	0.042		15	0.024		13
0.04		17	0.039		13	0.028		14
0.027		12	0.036		11	0.042		16
0.041		15	0.041		13	0.038		14
0.024		11	0.023		13	0.025		14
0.028		13	0.026		13	0.017		13
0.032		13	0.024		17	0.039		14
0.036		13	0.022		15	0.037		16
0.032		17	0.032		15	0.033		16
0.043		17	0.032		17	0.042		17
0.042		15	0.036		15	0.053		17
0.032		14	0.046		16	0.031		15
0.036		11	0.032		12	0.022		13
0.023		12	0.041		16	0.021		11

Table 29: G. fossarum wet weight (g) and total body length (mm) measured during whole experimental trial. Dead individuals are signed by  $\dagger$  and grey colour. n=50 for every exposure concentration and tested compound IMI (A), Confidor 200SL (B) and 6CNA (C) (continued).

# G. fossarum C

C								
$204.5 \ \mu g \ L^{-1}$			255.6 μg L <sup>-1</sup>			511.3 μg L <sup>-1</sup>		
Wet weight		Length	Wet weight		Length	Wet weight		Length
0.032	<i>†</i>	10	0.038	<i>†</i>	15	0.0142	†	10
0.038	<i>†</i>	7	0.036	; <del>'</del>	10	0.0193	<i>†</i>	11
0.043	; ;	14	0.027	; ;	10	0.201	<i>†</i>	18
0.031	<i>†</i>	10	0.080	; †	18	0.072	; ;	14
0.052	; ;	14	0.036	; ;	10	0.072	<i>†</i>	18
0.052	1	15	0.025	† †	8	0.033	; ;	12
0.046		17	0.030	1	15	0.026	<i>†</i>	17
0.024		10	0.043		15	0.04	,	12
0.047		15	0.044		15	0.023		12
0.023		11	0.041		14	0.053		12
0.026		11	0.027		14	0.036		16
0.020		14	0.027		12	0.019		8
0.052		10	0.0208		10	0.028		o
0.023		16	0.0423		12	0.043		19
0.023		10	0.0236		12	0.038		13
0.018		10	0.0343		13	0.04		15
0.056		17	0.0198		13	0.027		12
0.029		11	0.0383		11	0.041		10
0.029		11	0.0486		15	0.014		10
0.023		11	0.0213		17	0.036		15
0.017		11	0.0423		12	0.025		15
0.054		12	0.0394		14	0.022		12
0.061		11	0.032		10	0.042		11
0.061		16	0.0423		14	0.039		18
0.049		12	0.0432		9	0.036		14
0.031		12	0.0283		9	0.041		17
0.054		13	0.0301		10	0.034		13
0.029		8	0.0238		16	0.026		10
0.045		17	0.0324		14	0.034		14
0.043		16	0.0183		13	0.042		11
0.032		15	0.0239		11	0.043		11
0.038		16	0.0313		11	0.04		11
0.037		13	0.0193		15	0.044		11
0.031		12	0.0182		15	0.022		14
0.032		9	0.0473		17	0.048		16
0.038		12	0.0306		14	0.041		17
0.055		15	0.0412		15	0.02		18
0.048		15	0.0273		14	0.046		16
0.039		14	0.0201		12	0.034		11
0.024		13	0.0423		12	0.024		11
0.049		16	0.02		12	0.032		17
0.019		12	0.0197		12	0.036		14
0.049		18	0.0132		13	0.028		11
0.039		13	0.0214		15	0.032		13
0.027		10	0.0227		17	0.028		14
0.052		13	0.0412		12	0.023		17
0.056		13	0.0432		10	0.048		15
0.041		15	0.0327		10	0.042		16
0.049		18	0.0402		17	0.028		12
0.041		15	0.0187		11	0.028		12
0.045		16	0.0156		11	0.037		12
0.039		16	0.0428		15	0.019		11
0.037		13	0.0231		12	0.036		12

Table 30: CAT activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 2 min measurement (selected for every 30 s) and data normalised on total protein content (µmol/min/mg protein). (n=10).

CAT										
24 h - A										
Control										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.249984	0.236015	0.328127	0.189747	0.944999	0.526705	0.15148	0.103817	0.147454	0.26248
30 s 60 s	0.1434 0.1278	0.144931 0.107958	0.267479 0.261221	0.124783 0.110829	0.884891 0.878326	0.39884 0.449446	0.076513 0.040124	0.041044 0.003254	0.071584 0.034201	0.16636 0.13993
90 s	0.1278	0.107938	0.261221	0.110829	0.876684	0.449446	0.040124	0.003234	0.034201	0.13353
120 s	0.1319	0.092474	0.262044	0.10956	0.872544	0.461109	0.023223	0.020759	0.00724	0.13065
$102.2~\mu g~L^{-1}$										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.622296	0.378397	0.248703	0.244712	0.562342	0.303023	0.134211	0.29079	0.148624	0.43806
30 s	0.510951	0.309595	0.192585	0.17747	0.51609	0.205498	0.081636	0.211563	0.073619	0.09428
60 s	0.473377	0.300228	0.186722	0.156488	0.513651	0.194052	0.044354	0.186296	0.044421	0.08821
90 s 120 s	0.4616 0.459534	0.299063 0.298819	0.189618 0.193574	0.15161 0.150546	0.513624 0.512424	0.197784 0.207905	0.023377 0.015575	0.179639 0.178033	0.030781 0.019442	0.08333 0.07377
153.3 μg L <sup>-1</sup>	0.437334	0.270017	0.175574	0.130340	0.312424	0.207703	0.013373	0.170033	0.017442	0.07377
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.387351	0.293125	0.232242	0.368594	0.208142	0.21532	0.421714	0.199429	0.099711	0.3391
30 s	0.321101	0.228271	0.156464	0.331384	0.123799	0.145265	0.297398	0.136172	0.032089	0.2359
60 s	0.316213	0.223401	0.135936	0.329843	0.09466	0.102403	0.275566	0.119574	0.007261	0.23418
90 s	0.317878	0.223038	0.129975	0.331618	0.088608	0.079409	0.271171	0.115876	0.028495	0.23091
120 s 204.5 μg L <sup>-1</sup>	0.3209	0.223233	0.127897	0.332949	0.094649	0.067767	0.273193	0.115107	0.040036	0.20964
204.5 μg L time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.109651	0.568216	0.308946	0.456586	0.51346	0.384129	0.196056	0.157861	0.218183	0.39877
30 s	0.107902	0.48655	0.254737	0.415276	0.465719	0.311387	0.106084	0.087623	0.147254	0.28689
60 s	0.10576	0.447169	0.245754	0.414603	0.461604	0.30384	0.091746	0.05773	0.096521	0.30647
90 s	0.105092	0.427466	0.244844	0.414191	0.461798	0.307386	0.098287	0.04877	0.068282	0.3165
120 s	0.104304	0.417472	0.245152	0.413845	0.462283	0.312876	0.109563	0.047428	0.054617	0.32143
255.6 μg L <sup>-1</sup>	477	477	477	ATT	477	ATT	4.77	477	477	477
time 0 s	AU 0.370032	AU 0.222188	AU 0.308351	AU 0.72374	AU 0.265941	AU 0.122743	AU 0.189076	AU 0.434314	AU 0.370469	AU 0.67924
30 s	0.2753	0.181938	0.230039	0.708714	0.172579	0.044143	0.12084	0.352311	0.285773	0.60047
60 s	0.23435	0.170581	0.216697	0.722104	0.144939	0.017542	0.077435	0.321198	0.249752	0.61264
90 s	0.221666	0.17146	0.215311	0.72614	0.137805	0.011349	0.055976	0.31635	0.240864	0.60824
120 s	0.220359	0.176104	0.217619	0.734509	0.138946	0.012736	0.047281	0.31852	0.239757	0.59708
511.3 μg L <sup>-1</sup>										
time 0 s	AU 0.372761	AU 0.257039	AU 0.295672	AU 0.214039	AU 0.207367	AU 0.414632	AU 0.359254	AU 0.220962	AU 0.206582	AU 0.28352
30 s	0.372701	0.170586	0.225766	0.214039	0.152519	0.392458	0.339234	0.145373	0.123168	0.28332
60 s	0.280295	0.124015	0.214258	0.215311	0.142815	0.389164	0.244308	0.148769	0.10143	0.19975
90 s	0.265205	0.106573	0.213684	0.216409	0.142374	0.39124	0.250666	0.162347	0.094843	0.20069
120 s	0.258302	0.103872	0.212975	0.21806	0.142141	0.396799	0.260805	0.177832	0.095391	0.20278
		lues normalised o	•							
24 h	μmol/min	mg protein sample	μmol/min/mg prot	μmol/min	mg protein sample	μmol/min/mg prot	μmol/min	mg protein sample	μmol/min/mg prot	
control	3.30	0.14	23.26	3.61	0.17	21.04	2.44	0.14	17.20	
102.2 μg L <sup>-1</sup>	4.27	0.17	25.11	3.42	0.17	20.62	/	/	/	
153.3 μg L <sup>-1</sup>	2.93	0.15	20.21	3.26	0.16	21.04	4.68	0.17	28.20	
204.5 μg L <sup>-1</sup>	3.23	0.14	23.26	3.57	0.17	21.04	3.15	0.16	20.34	
255.6 μg L <sup>-1</sup>	3.92	0.13	29.92	4.06	0.12	34.78	12.01	0.13	90.31	
511.3 µg L <sup>-1</sup> control	3.00 3.34	0.12 0.14	25.64 23.58	1.71	0.15 0.14	11.56	4.54 2.77	0.17 0.17	26.38 16.70	
102.2 μg L <sup>-1</sup>				3.67		25.90			/	
102.2 μg L 153.3 μg L <sup>-1</sup>	2.08 2.99	0.13 0.13	16.64 23.58	6.75 2.30	0.13 0.17	54.06 13.56	2.01	0.12	17.20	
204.5 μg L <sup>-1</sup>	3.62	0.13	23.36	3.68	0.17	25.40	2.55	0.12	17.20	
204.5 μg L 255.6 μg L <sup>-1</sup>								/	/	
233.0 μg L 511.3 μg L <sup>-1</sup>	1.12 4.02	0.17 0.15	6.51 27.16	4.49 3.54	0.14 0.15	32.34 23.94	2.09	0.13		
control	2.34	0.15	14.30	3.54	0.15	22.30	2.09	0.15	15.58	
102.2 μg L <sup>-1</sup>	1.45	0.16	9.04	3.61	0.16	22.59				
153.3 μg L <sup>-1</sup>	3.89	0.16	24.35	4.68	0.10	22.30				
204.5 μg L <sup>-1</sup>	2.03	0.10	14.30	3.79	0.17	22.30				
255.6 μg L <sup>-1</sup>	1.21	0.14	9.22	3.79	0.17	27.55				
233.0 μg L 511.3 μg L <sup>-1</sup>	2.38	0.13	14.00	3.30	0.14	25.18				
control	4.41	0.17	27.96	4.68	0.17	28.20				
102.2 μg L <sup>-1</sup>	1.31	0.17	7.70	4.82	0.16	31.10				
153.3 μg L <sup>-1</sup>										
	4.61	0.17	27.96	2.22	0.13	16.70				
204.5 μg L <sup>-1</sup>	4.61 2.96	0.17 0.17	27.96 17.89	2.22	0.13 0.14	16.70 16.70				
$204.5~\mu\mathrm{g~L}^{\text{-}1}$	2.96	0.17	17.89	2.34	0.14	16.70				

Table 30: CAT activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 2 min measurement (selected for every 30 s) and data normalised on total protein content (µmol/min/mg protein). (n=10), (continued).

CAT										
24 h - B										
Control										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.249984	0.236015	0.328127	0.189747	0.944999	0.526705	0.15148	0.103817	0.147454	0.26248
30 s 60 s	0.1434	0.144931	0.267479	0.124783	0.884891	0.39884	0.076513	0.041044	0.071584	0.16636
90 s	0.1278 0.1225	0.107958 0.095192	0.261221 0.261815	0.110829 0.109345	0.878326 0.876684	0.449446 0.466036	0.040124 0.025223	0.003254 0.014785	0.034201 0.016996	0.13993 0.13353
120 s	0.1319	0.092474	0.262044	0.10956	0.872544	0.461109	0.023223	0.020759	0.00724	0.13065
102.2 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.522296	0.38397	0.248703	0.23414	0.5232	0.289814197	0.29285006	0.26924486	0.278629182	0.25714
30 s	0.510951	0.38595	0.198534	0.18726	0.52341	0.222244718	0.25269492	0.193485563	0.261264094	0.23475
60 s	0.473377	0.300228	0.185623	0.16285	0.51321	0.156715849	0.24618021	0.173053472	0.240855059	0.19025
90 s	0.4616	0.29963	0.18431	0.15064	0.51201	0.106401488	0.23923729	0.161487643	0.198077666	0.17877
120 s	0.359534	0.29819	0.183574	0.150741	0.512	0.100619267	0.21348412	0.132165169	0.182304731	0.17474
153.3 μg L <sup>-1</sup>	477	477	477	4.77	477	4.77	4 77	477	477	4.77
time 0 s	AU 0.159624	AU 0.42762	AU 0.205672	AU 0.214039	AU 0.106056	AU 0.21532	AU 0.421714	AU 0.100420	AU 0.000711	AU 0.2201
30 s	0.158624 0.093619	0.43763 0.2341	0.295672 0.225766	0.214039	0.196056 0.106084	0.145265	0.421714	0.199429 0.136172	0.099711 0.032089	0.3391 0.2359
60 s	0.054431	0.08549	0.214258	0.215311	0.091746	0.102403	0.275566	0.119574	0.007261	0.23418
90 s	0.023783	0.08235	0.213684	0.216409	0.098287	0.079409	0.271171	0.115876	0.028495	0.23091
120 s	0.01946	0.07323	0.212975	0.21806	0.109563	0.067767	0.273193	0.115107	0.040036	0.20964
204.5 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.109651	0.568216	0.308946	0.456586	0.51346	0.385123	0.232121	0.157861	0.218183	0.39877
30 s	0.107902	0.48655	0.254737	0.415276	0.465719	0.311341	0.146464	0.087421	0.147254	0.28689
60 s	0.10576	0.447169	0.245754	0.414603	0.461604	0.31352	0.135923	0.04772	0.096521	0.30647
90 s 120 s	0.105092 0.104304	0.427466 0.417472	0.244844 0.245152	0.414191 0.413845	0.461798 0.462283	0.30723 0.30235	0.129295 0.127897	0.041237 0.037428	0.068282 0.054617	0.3165 0.32143
255.6 μg L <sup>-1</sup>	0.104304	0.417472	0.243132	0.413643	0.402283	0.30253	0.12/89/	0.037428	0.034017	0.32143
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.37123	0.22215	0.307653	0.72374	0.206582	0.189076	0.265941	0.434314	0.122743	0.67924
30 s	0.2681	0.2231	0.220456	0.708714	0.124566	0.12084	0.172579	0.352311	0.044143	0.60142
60 s	0.22319	0.222	0.21035	0.722104	0.100233	0.077435	0.15679	0.321198	0.017542	0.60344
90 s	0.2241	0.20345	0.20458	0.72614	0.090345	0.055976	0.137805	0.31635	0.011349	0.63454
120 s	0.22012	0.1934	0.200134	0.734509	0.08391	0.047281	0.124546	0.31852	0.012736	0.59756
511.3 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.372761	0.257039	0.368594	0.208142	0.207367	0.414632	0.359254	0.220962	0.290469	0.28352
30 s	0.35451	0.170586	0.331384	0.123799	0.152519	0.392458	0.255978	0.175372	0.284573	0.21291
60 s 90 s	0.28195 0.27532	0.124015 0.11173	0.329843 0.321517	0.09466 0.088608	0.142815 0.142374	0.389164 0.39124	0.24138 0.25661	0.148769 0.152347	0.26527 0.24459	0.19975 0.20069
120 s	0.27332	0.11173	0.302149	0.088608	0.142141	0.396799	0.23001	0.13256	0.23977	0.20009
1200		ues normalised on p		0.07.017	0.1.21.1	0.570777	0.2 1505	0.13230	0.23777	0.20270
24 h	μmol/min	nes normansea on p	orem comem							
control		ma protoin comple	umal/min/ma prot	umol/min	ma protoin comple				umal/min/ma prot	
102.2 μg L <sup>-1</sup>	3 30	mg protein sample	μmol/min/mg prot	µmol/min	mg protein sample	μmol/min/mg prot 21.04	μmol/min 2 44	mg protein sample	μmol/min/mg prot 17.20	
	3.30	0.14	23.26	3.61	0.17	21.04	2.44	0.14	17.20	
	1.54	0.14 0.07	23.26 21.12	3.61 3.59	0.17 0.26	21.04 13.71	2.44 2.64	0.14 0.23	17.20 11.42	
$153.3~\mu g~L^{\text{-}1}$	1.54 1.41	0.14 0.07 0.07	23.26 21.12 20.21	3.61 3.59 5.28	0.17 0.26 0.25	21.04 13.71 21.04	2.44 2.64 7.08	0.14 0.23 0.25	17.20 11.42 28.20	
153.3 $\mu g L^{-1}$ 204.5 $\mu g L^{-1}$	1.54 1.41 1.61	0.14 0.07 0.07 0.08	23.26 21.12 20.21 20.21	3.61 3.59 5.28 3.33	0.17 0.26 0.25 0.24	21.04 13.71 21.04 14.00	2.44 2.64 7.08 7.51	0.14 0.23 0.25 0.28	17.20 11.42 28.20 27.00	
153.3 $\mu g L^{-1}$ 204.5 $\mu g L^{-1}$ 255.6 $\mu g L^{-1}$	1.54 1.41 1.61 3.00	0.14 0.07 0.07 0.08 0.08	23.26 21.12 20.21 20.21 39.73	3.61 3.59 5.28 3.33 2.86	0.17 0.26 0.25 0.24 0.26	21.04 13.71 21.04 14.00 10.95	2.44 2.64 7.08 7.51 6.83	0.14 0.23 0.25 0.28 0.23	17.20 11.42 28.20 27.00 29.18	
$153.3 \ \mu g \ L^{-1}$ $204.5 \ \mu g \ L^{-1}$ $255.6 \ \mu g \ L^{-1}$ $511.3 \ \mu g \ L^{-1}$	1.54 1.41 1.61 3.00 0.42	0.14 0.07 0.07 0.08 0.08 0.08	23.26 21.12 20.21 20.21 39.73 78.08	3.61 3.59 5.28 3.33 2.86 2.56	0.17 0.26 0.25 0.24 0.26 0.19	21.04 13.71 21.04 14.00 10.95 13.33	2.44 2.64 7.08 7.51 6.83 12.19	0.14 0.23 0.25 0.28 0.23 0.17	17.20 11.42 28.20 27.00 29.18 72.14	
153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 511.3 μg L <sup>-1</sup> control	1.54 1.41 1.61 3.00 0.42 3.34	0.14 0.07 0.07 0.08 0.08 0.05 0.14	23.26 21.12 20.21 20.21 39.73 78.08 23.58	3.61 3.59 5.28 3.33 2.86 2.56 3.67	0.17 0.26 0.25 0.24 0.26 0.19 0.14	21.04 13.71 21.04 14.00 10.95 13.33 25.90	2.44 2.64 7.08 7.51 6.83 12.19 2.77	0.14 0.23 0.25 0.28 0.23 0.17 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70	
153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> control 102.2 µg L <sup>-1</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16	0.17 0.26 0.25 0.24 0.26 0.19 0.14	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55	2.44 2.64 7.08 7.51 6.83 12.19 2.77	0.14 0.23 0.25 0.28 0.23 0.17 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70	
153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56	2.44 2.64 7.08 7.51 6.83 12.19 2.77 /	0.14 0.23 0.25 0.28 0.23 0.17 0.17 /	17.20 11.42 28.20 27.00 29.18 72.14 16.70 /	
153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23	
153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$ control 102.2 $\mbox{ µg L}^1$ 153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$ control 102.2 $\mbox{ µg L}^1$ 153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23	
153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$ control 102.2 $\mbox{ µg L}^1$ 153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$ control	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.16	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ 511.3 $\mbox{ µg L}^1$ control 102.2 $\mbox{ µg L}^1$ 153.3 $\mbox{ µg L}^1$ 204.5 $\mbox{ µg L}^1$ 255.6 $\mbox{ µg L}^1$ control 102.2 $\mbox{ µg L}^1$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.26	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 511.3 $\mbox{ ig } L^4$ 102.2 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 251.3 $\mbox{ ig } L^4$ 1511.3 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.16 0.29 0.25	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 511.3 $\mbox{ ig L}^{-1}$ control 102.2 $\mbox{ ig L}^{-1}$ 153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 151.3 $\mbox{ ig L}^{-1}$ control 102.2 $\mbox{ ig L}^{-1}$ 153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.26	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 511.3 $\mbox{ ig } L^4$ 102.2 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 251.3 $\mbox{ ig } L^4$ 1511.3 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.16 0.29 0.25	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 511.3 $\mbox{ ig L}^{-1}$ control 102.2 $\mbox{ ig L}^{-1}$ 153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 255.6 $\mbox{ ig L}^{-1}$ 151.3 $\mbox{ ig L}^{-1}$ control 102.2 $\mbox{ ig L}^{-1}$ 153.3 $\mbox{ ig L}^{-1}$ 204.5 $\mbox{ ig L}^{-1}$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.16 0.29 0.25	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 µg L <sup>4</sup> 204.5 µg L <sup>4</sup> 255.6 µg L <sup>4</sup> 511.3 µg L <sup>4</sup> control 102.2 µg L <sup>4</sup> 153.3 µg L <sup>4</sup> 204.5 µg L <sup>4</sup> 511.3 µg L <sup>4</sup> control 102.2 µg L <sup>4</sup> 511.3 µg L <sup>4</sup> control 102.2 µg L <sup>4</sup> 153.3 µg L <sup>4</sup> 204.5 µg L <sup>4</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.05 0.05 0.01 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.05	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.16 0.29 0.25 0.25	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40 2.89	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.03 0.03 0.16 0.16 0.17 0.17 0.17	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61 10.58	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11 5.48	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.16 0.29 0.25 0.25 0.12	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70 45.26	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 211.3 µg L <sup>-1</sup> control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> control	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40 2.89 4.41	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.03 0.16 0.16 0.17 0.17 0.14 0.27 0.16 0.16	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61 10.58 27.96 4.65	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11 5.48 4.68	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.27 0.16 0.29 0.25 0.12 0.12 0.15	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70 45.26 28.20 20.89	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 511.3 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.3 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40 2.89 4.41 0.75 4.08	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.03 0.16 0.16 0.17 0.17 0.17 0.14 0.27 0.16 0.16 0.16 0.16	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61 10.58 27.96 4.65 17.45	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11 5.48 4.68 3.03 3.91	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.16 0.29 0.25 0.12 0.12 0.15 0.15	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70 45.26 28.20 20.89 26.81	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup>	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40 2.89 4.41 0.75 4.08 5.98	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.03 0.16 0.16 0.17 0.17 0.14 0.27 0.16 0.16 0.23 0.21	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61 10.58 27.96 4.65 17.45 27.96	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11 5.48 4.68 3.03 3.91 2.47	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.16 0.29 0.25 0.12 0.12 0.15 0.15	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70 45.26 28.20 20.89 26.81 16.70	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	
153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 511.3 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 153.3 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 255.6 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 204.5 $\mbox{ ig } L^4$ 255.3 $\mbox{ ig } L^4$ 255.3 $\mbox{ ig } L^4$	1.54 1.41 1.61 3.00 0.42 3.34 1.83 0.69 1.06 3.67 2.53 2.34 1.51 4.01 4.09 3.40 2.89 4.41 0.75 4.08	0.14 0.07 0.07 0.08 0.08 0.05 0.14 0.05 0.05 0.05 0.05 0.03 0.16 0.16 0.17 0.17 0.17 0.14 0.27 0.16 0.16 0.16 0.16	23.26 21.12 20.21 20.21 39.73 78.08 23.58 35.26 13.58 23.58 123.56 78.08 14.30 9.20 24.35 24.35 23.61 10.58 27.96 4.65 17.45	3.61 3.59 5.28 3.33 2.86 2.56 3.67 3.16 3.70 2.28 2.86 19.09 3.65 3.47 5.46 4.94 3.11 5.48 4.68 3.03 3.91	0.17 0.26 0.25 0.24 0.26 0.19 0.14 0.12 0.16 0.17 0.27 0.16 0.29 0.25 0.12 0.12 0.15 0.15	21.04 13.71 21.04 14.00 10.95 13.33 25.90 26.55 23.56 13.56 10.00 71.23 22.30 12.13 22.30 20.00 25.70 45.26 28.20 20.89 26.81	2.44 2.64 7.08 7.51 6.83 12.19 2.77 / 3.66 3.54 4.57	0.14 0.23 0.25 0.28 0.23 0.17 0.17 / 0.21 0.18 0.17	17.20 11.42 28.20 27.00 29.18 72.14 16.70 / 17.20 19.23 27.23	

Table 30: CAT activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 2 min measurement (selected for every 30 s) and data normalised on total protein content (µmol/min/mg protein). (n=10), (continued).

CAT										
24 h - C										
Control										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.417296	0.223984	0.447792	0.280212	0.282202	0.308197	0.15148	0.103817	0.156924	0.26248
30 s 60 s	0.289438 0.301048	0.113222 0.084728	0.295567 0.382694	0.16861 0.11518	0.153354 0.151725	0.206356 0.18544	0.08513 0.04124	0.041044 0.003654	0.091584 0.0458201	0.18635 0.13993
90 s	0.307308	0.072444	0.370074	0.11318	0.131723	0.154564	0.035223	0.014995	0.026396	0.13993
120 s	0.304934	0.071731	0.379914	0.100404	0.132029	0.137677	0.024539	0.020759	0.00921	0.1361
102.2 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s 30 s	0.280964 0.168339	0.469795 0.351609	0.300103 0.150794	0.406851 0.243353	0.478454 0.318117	0.51487 0.381618	0.2928501 0.2578167	0.26243884 0.19411345	0.278684764 0.26004986	0.25894 0.2322
60 s	0.164014	0.378154	0.113745	0.260187	0.339945	0.382504	0.2474515	0.17630525	0.24079709	0.19238
90 s	0.175256	0.371781	0.124816	0.254015	0.344779	0.400398	0.2186981	0.16448088	0.1927135	0.17532
120 s	0.177516	0.371946	0.131744	0.243844	0.333275	0.391553	0.2170096	0.13275557	0.18059799	0.17627
153.3 μg L <sup>-1</sup>	477	477	477	477	477	477	477	477	477	477
time 0 s	AU 0.15435	AU 0.43763	AU 0.295472	AU 0.214039	AU 0.195969	AU 0.568216	AU 0.308946	AU 0.199429	AU 0.385123	AU 0.23537
30 s	0.09619	0.233	0.24591	0.21662	0.16943	0.48655	0.254737	0.147011	0.323971	0.14493
60 s	0.056793	0.08549	0.214958	0.215311	0.09946	0.447169	0.245754	0.12395	0.314891	0.14393
90 s	0.023569	0.08435	0.219714	0.216409	0.09596	0.427466	0.244844	0.115921	0.30491	0.14885
120 s 204.5 μg L <sup>-1</sup>	0.019502	0.07458	0.210365	0.215936	0.0235	0.417472	0.245152	0.115592	0.30855	0.1279
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s	0.104551	0.21532	0.421714	0.456586	0.51346	0.099711	0.339102	0.157861	0.218183	0.39877
30 s	0.104402	0.1456765	0.2934498	0.415276	0.465719	0.032089	0.33593	0.087326	0.147254	0.29989
60 s 90 s	0.103 0.103092	0.10456 0.0759	0.211166 0.2111	0.414603 0.414191	0.461604 0.461798	0.02261 0.021495	0.3341 0.280905	0.04767 0.0414555	0.106521 0.098282	0.30047 0.3165
120 s	0.103092	0.0719356	0.20333	0.413845	0.462283	0.020035	0.59687	0.038923	0.084617	0.31114
$255.6~\mu g~L^{-1}$										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
0 s 30 s	0.695786	0.190596	0.482663	0.231408	0.170252	0.1896986 0.167924	0.265941	0.434314	0.122743 0.044143	0.35025 0.27892
50 s 60 s	0.578729 0.582328	0.05812 0.015624	0.352929 0.355879	0.073957 0.030101	0.05696 0.013514	0.156435	0.252579 0.25679	0.352311 0.321198	0.044143	0.27892
90 s	0.588393	0.016179	0.357232	0.008665	0.016508	0.1545976	0.237815	0.31635	0.011349	0.23166
120 s	0.595026	0.034017	0.347654	0.016339	0.032874	0.147281	0.244546	0.31852	0.012736	0.22231
511.3 μg L <sup>-1</sup>										
time 0 s	AU 0.292449	AU 0.202655	AU 0.294565	AU 0.175453	AU 0.500334	AU 0.190094	AU 0.679242	AU 0.220962	AU 0.290469	AU 0.28352
30 s	0.158771	0.088393	0.17603	0.074383	0.353917	0.093607	0.60142	0.175372	0.284563	0.27603
60 s	0.120821	0.040104	0.141868	0.028976	0.394518	0.060739	0.60344	0.148769	0.278902	0.24507
90 s 120 s	0.10851 0.093683	0.00626 0.020868	0.118905 0.112824	0.005186 0.029738	0.397787 0.397524	0.041749 0.028976	0.634543 0.59756	0.152347 0.13256	0.265459 0.2567977	0.20694 0.20659
		s normalised on pr		0.029738	0.397324	0.028976	0.39730	0.13236	0.2367977	0.20039
24 h	μmol/min	•	μmol/min/mg prot	μmol/min	mg protein sample	μmol/min/mg prot	μmol/min	mg protein sample	μmol/min/mg prot	
control	3.30	0.14	23.26	3.61	0.17	21.04	2.44	0.14	17.20	
$102.2~\mu\mathrm{g~L}^{\text{-}1}$	2.71	0.12	22.17	3.81	0.13	29.17	4.86	0.15	31.56	
153.3 μg L <sup>-1</sup>	4.53	0.15	30.22	3.71	0.17	22.50	3.62	0.12	29.45	
204.5 μg L <sup>-1</sup>	2.65	0.26	10.23	3.37	0.16	20.79	4.21	0.14	29.65	
255.6 μg L <sup>-1</sup>	2.64	0.13	20.98	8.22	0.10	85.09	5.03	0.12	41.20	
511.3 μg L <sup>-1</sup> control	5.21 3.34	0.11 0.14	46.10 23.58	2.69 3.67	0.10 0.14	26.40 25.90	8.72 2.77	0.14 0.17	62.30 16.70	
102.2 μg L <sup>-1</sup>	2.56	0.14	23.44	3.23	0.14	23.56	4.22	0.17	23.20	
153.3 μg L <sup>-1</sup>	2.64	0.11	23.58	4.87	0.15	33.56	2.87	0.15	19.23	
204.5 μg L <sup>-1</sup>	1.61	0.12	13.45	1.51	0.10	14.56	3.56	0.15	24.56	
255.6 μg L <sup>-1</sup>	9.19	0.14	67.30	10.01	0.15	69.03	4.80	0.12	40.00	
$511.3~\mu g~L^{\text{-}1}$	6.83	0.11	61.81	4.22	0.13	33.71	4.55	0.12	37.89	
control	2.34	0.16	14.30	3.65	0.16	22.30				
102.2 μg L <sup>-1</sup>	4.41	0.11	39.17	3.68	0.17	22.31				
153.3 μg L <sup>-1</sup>	1.38	0.11	12.57	3.26	0.17	19.78				
204.5 μg L <sup>-1</sup>	3.04	0.12	24.53	3.76	0.17	22.13				
255.6 μg L <sup>-1</sup>	3.54	0.10	35.79	4.13	0.16	25.34				
511.3 μg L <sup>-1</sup> control	4.77 4.41	0.11 0.16	44.33 27.96	7.49 4.68	0.16 0.17	45.67 28.20				
102.2 μg L <sup>-1</sup>	4.27	0.13	32.57	4.07	0.17	23.40				
153.3 μg L <sup>-1</sup>	4.05	0.15	27.56	2.76	0.17	16.70				
204.5 μg L <sup>-1</sup>	4.71	0.15	32.45	5.87	0.17	35.60				
255.6 μg L <sup>-1</sup>										
	7.79	0.20	38.97	10.32	0.16	65.34				
511.3 μg L <sup>-1</sup>	7.79 7.69	0.20 0.10	38.97 74.44	10.32 7.27	0.16 0.16	65.34 44.30				

Table 31: GST activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 5 min measurement (selected for every min) and data normalised on total protein content (nmol/min/mg protein). (n=10).

GST										
24 h - A										
Control										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.146236	0.042268	0.118228	-0.025388	0.238809	0.225794	0.268316	0.37774	0.256141	0.13408
2 min 3 min	0.215348 0.277997	0.092 0.13941	0.180029 0.234261	0.015482 0.054798	0.32088 0.348072	0.265945 0.387399	0.313015 0.42122	0.462245 0.518991	0.24998 0.249274	0.21085 0.33364
3 min 4 min	0.333979	0.179978	0.279339	0.102651	0.348072	0.409343	0.44122	0.563051	0.362658	0.38016
5 min	0.384914	0.21773	0.322515	0.129369	0.457125	0.444866	0.424734	0.593668	0.377037	0.40418
$102.2~\mu\mathrm{g~L^{\text{-}1}}$										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.909622	0.802723	1.176711	1.074273	0.222303	0.460296	0.39944	0.44331	0.220164	0.14542
2 min	1.545629	0.933118	1.428991	1.243378	0.388799	0.79716	0.656945	0.709702	0.381393	0.28287
3 min	2.144172	1.053127	1.653984	1.39861	0.541888	1.093051	0.888451	0.948591	0.529834	0.41317
4 min 5 min	2.7892 3.45683	1.16699 1.275071	1.859892 2.048848	1.546903 1.687243	0.687614 0.825741	1.357828 1.597588	1.099222 1.289486	1.166439 1.363909	0.669636 0.801064	0.53381 0.64913
153.3 μg L <sup>-1</sup>	3.43003	1.273071	2.040040	1.007243	0.025741	1.577566	1.207400	1.505707	0.001004	0.04713
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.590015	1.021672	1.132618	0.469923	0.366457	0.124729	0.186352	0.056928	0.102605	0.17689
2 min	0.999548	1.188764	1.289452	0.769374	0.615384	0.229973	0.350374	0.220762	0.221323	0.19082
3 min	1.357925	1.34323	1.436688	1.039295	0.838152	0.32992	0.500128	0.373464	0.332224	0.20525
4 min	1.696015	1.480263	1.573813	1.282897	1.001446	0.423767	0.641188	0.516052	0.437991	0.21928
5 min	2.02275	1.612571	1.703852	1.503956	1.170568	0.514482	0.770873	0.650197	0.538277	0.23116
204.5 μg L <sup>-1</sup> time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.817308	1.056696	1.131106	0.280531	0.348402	0.19448	0.054815	0.115823	0.359723	0.98658
2 min	1.355639	1.277275	1.313312	0.496001	0.566677	0.35748	0.217042	0.207232	0.496969	1.07743
3 min	1.829354	1.481491	1.48226	0.693072	0.766985	0.506428	0.365746	0.293624	0.623925	1.16675
4 min	2.273806	1.666513	1.638484	0.876962	0.953984	0.647927	0.503935	0.376172	0.743778	1.25782
5 min	2.895171	1.832829	1.785761	1.049342	1.129494	0.779441	0.632618	0.454382	0.856411	1.34941
255.6 μg L <sup>-1</sup>										
time 1 min	AU 0.973248	AU 1.046696	AU 0.870266	AU 0.140116	AU 0.376066	AU 0.399981	AU -0.058228	AU -0.002271	AU 0.064088	AU 0.37052
2 min	1.487858	1.191279	0.888636	0.284721	0.651402	0.659692	0.064438	0.130072	0.132022	0.54618
3 min	1.850204	1.327096	0.906303	0.419976	0.901567	0.897886	0.179641	0.257808	0.197218	0.71663
4 min	2.104055	1.452359	0.92442	0.546363	1.128708	1.132115	0.28668	0.381754	0.257507	0.87727
5 min	2.368362	1.574003	0.942526	0.666827	1.342134	1.346819	0.388663	0.501742	0.31682	1.04344
511.3 μg L <sup>-1</sup>										
time	AU	AU	AU 1.107798	AU	AU	AU	AU	AU	AU	AU
				1.004165	0.115112	0.398807	0.798974	0.274961	0.366106	0.04136
1 min	1.122983	0.272581							0.600006	0.00187
2 min	1.746232	0.310429	1.286498	1.130215	0.243017	0.678667	1.262047	0.444148	0.609996 0.835636	0.09187 0.14179
									0.609996 0.835636 1.044832	0.09187 0.14179 0.18935
2 min 3 min	1.746232 2.158413	0.310429 0.346949	1.286498 1.45279	1.130215 1.249372	0.243017 0.36618	0.678667 0.926952	1.262047 1.662634	0.444148 0.601651	0.835636	0.14179
2 min 3 min 4 min	1.746232 2.158413 2.35678 2.459443	0.310429 0.346949 0.385358	1.286498 1.45279 1.608845 1.75555	1.130215 1.249372 1.363838	0.243017 0.36618 0.48214	0.678667 0.926952 1.152651	1.262047 1.662634 2.005436	0.444148 0.601651 0.746919	0.835636 1.044832	0.14179 0.18935
2 min 3 min 4 min	1.746232 2.158413 2.35678 2.459443 Calculated va μmol/min	0.310429 0.346949 0.385358 0.420955	1.286498 1.45279 1.608845 1.75555 on protein content nmol/min/mg prot	1.130215 1.249372 1.363838 1.478732	0.243017 0.36618 0.48214	0.678667 0.926952 1.152651	1.262047 1.662634 2.005436	0.444148 0.601651 0.746919	0.835636 1.044832	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control	1.746232 2.158413 2.35678 2.459443 Calculated vo µmol/min 0.07	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06	0.678667 0.926952 1.152651 1.361604	1.262047 1.662634 2.005436 2.332274	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06	0.835636 1.044832 1.240717	0.14179 0.18935
2 min 3 min 4 min 5 min  24 h control 102.2 µg L <sup>-1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated va μmol/min	0.310429 0.346949 0.385358 0.420955 ulues normalised o mg protein sample	1.286498 1.45279 1.608845 1.75555 on protein content nmol/min/mg prot	1.130215 1.249372 1.363838 1.478732	0.243017 0.36618 0.48214 0.591267 mg protein sample	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot	1.262047 1.662634 2.005436 2.332274 μmol/min	0.444148 0.601651 0.746919 0.883645 mg protein sample	0.835636 1.044832 1.240717 nmol/min/mg prot	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated vo µmol/min 0.07 0.12 0.05	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09	1.286498 1.45279 1.608845 1.75555 on protein content nmol/min/mg prot 833 1410 833	1.130215 1.249372 1.363838 1.478732 μmol/min 0.01 0.05 0.04	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 /	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated vo pmol/min 0.07 0.12 0.05 0.02	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320	1.130215 1.249372 1.363838 1.478732 <b>µmol/min</b> 0.01 0.05 0.04 0.06	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07	0.444148 0.601651 0.746919 0.883645 <b>mg protein sample</b> 0.06 0.09 0.09	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>4</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve µmol/min 0.07 0.12 0.05 0.02	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08	1.286498 1.45279 1.608845 1.75555 on protein content nmol/min/mg prot 833 1410 833 320 650	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01 0.05 0.04 0.06 0.07	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457	1.262047 1.662634 2.005436 2.332274 µmol/min 0.01 / 0.02 0.07 0.14	0.444148 0.601651 0.746919 0.883645 <b>mg protein sample</b> 0.06 0.09 0.09	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621 978	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>4</sup> 204.5 µg L <sup>4</sup> 255.6 µg L <sup>1</sup> 511.3 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated vo pmol/min 0.07 0.12 0.05 0.02 0.05 0.06	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01 0.05 0.04 0.06 0.07 0.05	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474	1.262047 1.662634 2.005436 2.332274 µmol/min 0.01 / 0.02 0.07 0.14 0.02	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621 978 133	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>3</sup> 255.6 µg L <sup>4</sup> 511.3 µg L <sup>1</sup> control	1.746232 2.158413 2.35678 2.459443 Calculated ve  µmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09	0.310429 0.346949 0.385358 0.420955  alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158	1.130215 1.249372 1.363838 1.478732 <b>µmol/min</b> 0.01 0.05 0.04 0.06 0.07 0.05 0.02	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.08 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261	1.262047 1.662634 2.005436 2.332274 µmol/min 0.01 / 0.02 0.07 0.14	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621 978 133 324	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve  µmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05	0.310429 0.346949 0.385358 0.420955  alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369	1.262047 1.662634 2.005436 2.332274 µmol/min 0.01 / 0.02 0.07 0.14 0.02 0.03 /	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621 978 133 324 /	0.14179 0.18935
2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>2</sup> 511.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve  µmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 an protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.09 0.10 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369 464	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 /	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.08	0.835636 1.044832 1.240717 nmol/min/mg prot 156 / 201 621 978 133 324 /	0.14179 0.18935
2 min 3 min 4 min 5 min 4 min 5 min 24 h control 102.2 µg L <sup>4</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve  µmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340	1.130215 1.249372 1.363838 1.478732 µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.10 0.07	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359	1.262047 1.662634 2.005436 2.332274 pmol/min 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.08	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234	0.14179 0.18935
2 min 3 min 4 min 5 min  24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 211.3 µg L <sup>1</sup> 102.2 µg L <sup>4</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480	1.130215 1.249372 1.363838 1.478732  µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.10 0.07 0.08 0.10 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min  24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>1</sup> 211.3 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 153.3 µg L <sup>2</sup> 153.3 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 153.3 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06	0.310429 0.346949 0.385358 0.420955  alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 an protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03	0.243017 0.36618 0.48214 0.591267 mg protein sample 0.06 0.07 0.08 0.08 0.10 0.07 0.08 0.10 0.07 0.08	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506	1.262047 1.662634 2.005436 2.332274 pmol/min 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01	0.444148 0.601651 0.746919 0.883645 mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.08	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 255.6 µg L <sup>3</sup> 511.3 µg L <sup>1</sup> control 102.2 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 255.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> control	1.746232 2.158413 2.35678 2.459443 Calculated ve  µmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.06 0.07	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380	1.130215 1.249372 1.363838 1.478732  µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.07 0.008 0.005 0.03 0.05 0.03 0.10	0.678667 0.926952 1.152651 1.361604 nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 225.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> control 102.2 µg L <sup>3</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 251.3 µg L <sup>1</sup> 204.2 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 202.2 µg L <sup>2</sup> 151.2 µg L <sup>1</sup> 151.2 µg L <sup>1</sup> 151.3 µ	1.746232 2.158413 2.35678 2.459443 Calculated ve	0.310429 0.346949 0.385358 0.420955  alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435	1.130215 1.249372 1.363838 1.478732  µmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.04 0.02	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.008 0.10 0.07	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 225.6 µg L <sup>2</sup> 151.3 µg L <sup>1</sup> control 102.2 µg L <sup>2</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 255.6 µg L <sup>2</sup> 511.3 µg L <sup>2</sup> 151.3 µg L <sup>2</sup> 151.3 µg L <sup>3</sup> 151.3 µg L <sup>3</sup> 151.3 µg L <sup>3</sup> 151.3 µg L <sup>3</sup> 151.3 µg L <sup>4</sup> 151.3 µg L <sup>3</sup> 151.3 µg L <sup>4</sup> 151.3 µg L <sup>4</sup> 153.3 µg L <sup>4</sup> 153.3 µg L <sup>4</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.04 0.06 0.03 0.02 0.03	0.310429 0.346949 0.385358 0.420955  **alues normalised a**  **mg protein sample* 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.10 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.07 0.08 0.07 0.07 0.08	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 5 min 5 min 24 h control 10.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 153.3 µg L <sup>4</sup> 204.5 µg L <sup>2</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 151.3 µg L <sup>1</sup> 255.6 µg L <sup>2</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 151.3 µg L <sup>2</sup> 204.5 µg L <sup>2</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>2</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>2</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02	0.310429 0.346949 0.385358 0.420955  alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.07 0.008 0.006 0.07 0.010 0.07 0.07 0.03 0.10 0.07 0.07 0.07 0.07	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 10.22 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2255.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2255.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.008 0.10 0.07 0.008 0.006 0.05 0.03 0.10 0.07 0.07 0.013 0.09	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 4 min 5 min 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2255.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 2255.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 154.3 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02	0.310429 0.346949 0.385358 0.420955 alues normalised a mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.009 0.07 0.06 0.08 0.09 0.07 0.06 0.08 0.08	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.04 0.02 0.03 0.01 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.04 0.07 0.06 0.112	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.05 0.03 0.10 0.07 0.01 0.07 0.07 0.09 0.10	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 511.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 255.6 µg L <sup>1</sup> 255.9 µg L <sup>1</sup> 265.9 µg L <sup>1</sup> 255.9 µg L <sup>1</sup> 2	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518 466 351	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.04 0.07 0.05 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.04 0.07 0.06 0.12 0.03	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.07 0.08 0.010 0.07 0.08 0.006 0.05 0.03 0.10 0.07 0.07 0.07 0.07 0.07 0.13 0.09 0.15 0.11	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827 258	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 205.2 µg L <sup>1</sup> 153.2 µg L <sup>1</sup> 205.2 µg L <sup>1</sup> 11.3 µg L <sup>1</sup> 205.2 µg L	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02	0.310429 0.346949 0.385358 0.420955 alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518 466 351 564	1.130215 1.249372 1.363838 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.04 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.07 0.05 0.02 0.03 0.04 0.07 0.06 0.01 0.07 0.06 0.12 0.03 0.04	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.006 0.07 0.01 0.07 0.08 0.06 0.05 0.03 0.10 0.07 0.07 0.13 0.09 0.15 0.11 0.07	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827 258	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 5 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 2153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.04 0.06 0.03 0.02 0.03 0.02	0.310429 0.346949 0.385358 0.420955  alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518 466 351 564	1.130215 1.249372 1.36383 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.04 0.07	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.08 0.10 0.07 0.03 0.10 0.07 0.01 0.07 0.13 0.09 0.13 0.09 0.15 0.11 0.07 0.06	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827 258 597	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 5 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 255.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>3</sup> 151.3 µg L <sup>1</sup> 204.5 µg L <sup>3</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>3</sup> 204.5 µg L <sup></sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.03	0.310429 0.346949 0.385358 0.420955 nlues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.008 0.009 0.07 0.06 0.08 0.08 0.09 0.07	1.286498 1.45279 1.608845 1.75555 m protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518 466 351 564 551	1.130215 1.249372 1.36383 1.478732  pumol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.04 0.07 0.06 0.10 0.07 0.06 0.10 0.03 0.04 0.02 0.03 0.04 0.07 0.06 0.12 0.03 0.04 0.07	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.008 0.10 0.07 0.03 0.10 0.07 0.03 0.10 0.07 0.13 0.09 0.15 0.11 0.07 0.06 0.12	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827 258 597 258	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935
2 min 3 min 4 min 5 min 5 min 5 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> 205.6 µg L <sup>1</sup> 151.3 µg L <sup>1</sup> control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 2153.3 µg L <sup>1</sup> 204.5 µg L <sup>1</sup> 153.3 µg L <sup>1</sup>	1.746232 2.158413 2.35678 2.459443 Calculated ve pmol/min 0.07 0.12 0.05 0.02 0.05 0.06 0.09 0.05 0.03 0.03 0.04 0.06 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.04 0.06 0.03 0.02 0.03 0.02	0.310429 0.346949 0.385358 0.420955  alues normalised of mg protein sample 0.08 0.09 0.06 0.07 0.08 0.08 0.08 0.08 0.08 0.08 0.08	1.286498 1.45279 1.608845 1.75555 in protein content nmol/min/mg prot 833 1410 833 320 650 789 1158 600 458 340 480 635 380 435 380 245 518 466 351 564	1.130215 1.249372 1.36383 1.478732  pmol/min 0.01 0.05 0.04 0.06 0.07 0.05 0.02 0.03 0.04 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.02 0.03 0.04 0.07	0.243017 0.36618 0.48214 0.591267  mg protein sample 0.06 0.07 0.08 0.08 0.08 0.10 0.07 0.08 0.08 0.08 0.10 0.07 0.03 0.10 0.07 0.01 0.07 0.13 0.09 0.13 0.09 0.15 0.11 0.07 0.06	0.678667 0.926952 1.152651 1.361604  nmol/min/mg prot 157 696 457 734 878 474 261 369 464 359 632 506 313 269 513 504 623 827 258 597	1.262047 1.662634 2.005436 2.332274 <b>µmol/min</b> 0.01 / 0.02 0.07 0.14 0.02 0.03 / 0.01 0.03	0.444148 0.601651 0.746919 0.883645  mg protein sample 0.06 0.09 0.09 0.12 0.14 0.12 0.08 0.08 0.08 0.06 0.11 0.11	0.835636 1.044832 1.240717  nmol/min/mg prot 156 / 201 621 978 133 324 / 224 234 326	0.14179 0.18935

Table 31: GST activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 5 min measurement (selected for every min) and data normalised on total protein content (nmol/min/mg protein). (n=10), (continued).

GST										
24 h - B										
Control										
time 1 min	AU 0.146236	AU 0.042268	AU 0.118228	AU -0.025388	AU 0.238809	AU 0.225794	AU 0.268316	AU 0.37774	AU 0.256141	AU 0.13408
2 min	0.215348	0.092	0.110220	0.015482	0.32088	0.265945	0.313015	0.462245	0.24998	0.13408
3 min	0.277997	0.13941	0.234261	0.054798	0.348072	0.387399	0.42122	0.518991	0.249274	0.33364
4 min	0.333979	0.179978	0.279339	0.102651	0.3573	0.409343	0.4467295	0.563051	0.362658	0.38016
5 min	0.384914	0.21773	0.322515	0.129369	0.457125	0.444866	0.424734	0.593668	0.377037	0.40418
102.2 μg L <sup>-1</sup>	477	4.7.7	4.77	4.77	4.77	477	4.77	477	4.77	4.77
time 1 min	AU 0.246487	AU 0.214286	AU 0.165125	AU 0.1922	AU 0.231688	AU 0.099655	AU 0.205051	AU 0.20567	AU 0.16857	AU 0.34794
2 min	0.463121	0.412305	0.322555	0.32956	0.40615	0.209559	0.373268	0.3568	0.356932	0.47834
3 min	0.65974	0.595308	0.467042	0.478132	0.569532	0.314147	0.531424	0.49535	0.46854	0.7792
4 min	0.842501	0.764909	0.594008	0.576049	0.725613	0.416258	0.6795	0.793	0.57891	0.98673
5 min	1.011514	0.926191	0.707406	0.723529	0.876813	0.512416	0.836608	0.98214	0.61234	1.05743
153.3 μg L <sup>-1</sup> time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.278601	0.342678	0.411578	0.977062	1.045793	0.209089	0.213854	0.39062	0.164019	0.20859
2 min	0.498475	0.593665	0.624214	1.109753	1.184476	0.38398	0.399634	0.661876	0.297811	0.33613
3 min	0.700568	0.818906	0.824877	1.235106	1.312686	0.546653	0.571913	0.908192	0.418247	0.4597
4 min	0.88758	1.026637	1.008798	1.351356	1.434061	0.698393	0.73199	1.130932	0.528877	0.57721
5 min 204.5 μg L <sup>-1</sup>	1.06179	1.212562	1.181936	1.460631	1.548873	0.837494	0.881334	1.33405	0.62801	0.68869
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.411578	0.534649	0.409561	1.042248	0.977062	1.132618	0.870266	0.34278	0.4178	0.45982
2 min	0.624214	0.828774	0.657076	1.72172	1.109753	1.289452	0.888636	0.593725	0.624568	0.75602
3 min	0.824877	1.095027	0.841081	2.405085	1.235106	1.436688	0.906303	0.928906	0.83987	0.90587
4 min 5 min	1.008798 1.181936	1.338607 1.556883	1.007319 1.162938	3.2345 3.65467	1.351356 1.460631	1.573813 1.703852	0.92442 0.942526	1.16537 1.214362	1.0088 1.18128	1.05743 1.25733
255.6 μg L <sup>-1</sup>	1.101730	1.550005	1.102936	3.03407	1.400031	1.703632	0.742320	1.214302	1.10120	1.23733
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.163465	0.304731	0.399482	0.072382	0.507572	0.649887	0.539649	0.478956	0.327894	0.2439
2 min	0.301388	0.44971	0.579965	0.182122	0.79617	0.986566	0.923537	0.67845	0.468921	0.45689
3 min 4 min	0.431836 0.553647	0.576334 0.700386	0.7453 0.897622	0.287766 0.387947	1.057329 1.291838	1.291838 1.567477	1.272385 1.589471	0.894821 1.26733	0.68392 0.789032	0.78943 0.90458
5 min	0.673951	0.845805	1.043112	0.387947	1.512174	1.836475	1.871736	1.56201	0.81239	1.0583
511.3 μg L <sup>-1</sup>									*******	
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.219327	0.266947	0.327678	0.350998	0.02081	0.31938	0.459346	0.668968	0.432442	0.12378
1 min 2 min	0.219327 0.433335	0.266947 0.473167	0.327678 0.458036	0.350998 0.515273	0.02081 0.028369	0.31938 0.5383	0.459346 0.643325	0.668968 0.96649	0.432442 0.618388	0.12378 0.14482
1 min 2 min 3 min	0.219327 0.433335 0.631108	0.266947 0.473167 0.664827	0.327678 0.458036 0.581985	0.350998 0.515273 0.670777	0.02081 0.028369 0.037408	0.31938 0.5383 0.743934	0.459346 0.643325 0.807949	0.668968 0.96649 1.248589	0.432442 0.618388 0.79022	0.12378 0.14482 0.1651
1 min 2 min	0.219327 0.433335	0.266947 0.473167	0.327678 0.458036	0.350998 0.515273	0.02081 0.028369	0.31938 0.5383	0.459346 0.643325	0.668968 0.96649	0.432442 0.618388	0.12378 0.14482
1 min 2 min 3 min 4 min 5 min	0.219327 0.433335 0.631108 0.814218 0.986821	0.266947 0.473167 0.664827 0.844545	0.327678 0.458036 0.581985 0.700259 0.814241	0.350998 0.515273 0.670777 0.815064	0.02081 0.028369 0.037408 0.045741	0.31938 0.5383 0.743934 0.937396	0.459346 0.643325 0.807949 0.965685	0.668968 0.96649 1.248589 1.547566	0.432442 0.618388 0.79022 0.955609	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min	0.219327 0.433335 0.631108 0.814218 0.986821	0.266947 0.473167 0.664827 0.844545 1.014197	0.327678 0.458036 0.581985 0.700259 0.814241	0.350998 0.515273 0.670777 0.815064	0.02081 0.028369 0.037408 0.045741	0.31938 0.5383 0.743934 0.937396	0.459346 0.643325 0.807949 0.965685	0.668968 0.96649 1.248589 1.547566	0.432442 0.618388 0.79022 0.955609	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 24 h control	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content	0.350998 0.515273 0.670777 0.815064 0.953058	0.02081 0.028369 0.037408 0.045741 0.056071	0.31938 0.5383 0.743934 0.937396 1.122665	0.459346 0.643325 0.807949 0.965685 1.11235	0.668968 0.96649 1.248589 1.547566 1.783996	0.432442 0.618388 0.79022 0.955609 1.129218	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min  24 h control 102.2 µg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised o	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot	0.350998 0.515273 0.670777 0.815064 0.953058 μmol/min	0.02081 0.028369 0.037408 0.045741 0.056071 mg protein sample	0.31938 0.5383 0.743934 0.937396 1.122665	0.459346 0.643325 0.807949 0.965685 1.11235	0.668968 0.96649 1.248589 1.547566 1.783996 mg protein sample	0.432442 0.618388 0.79022 0.955609 1.129218	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min  24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689	0.350998 0.515273 0.670777 0.815064 0.953058 μmol/min 0.01 0.03 0.05	0.02081 0.028369 0.037408 0.045741 0.056071 <b>mg protein sample</b> 0.06 0.11	0.31938 0.5383 0.743934 0.937396 1.122665 <b>nmol/min/mg prot</b> 157 221 432	0.459346 0.643325 0.807949 0.965685 1.11235 μmol/min 0.01 0.01	0.668968 0.96649 1.248589 1.547566 1.783996 mg protein sample 0.06 0.08	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 156 222 501	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489	0.350998 0.515273 0.670777 0.815064 0.953058 μmol/min 0.01 0.03 0.05 0.05	0.02081 0.028369 0.037408 0.045741 0.056071 mg protein sample 0.06 0.11 0.11	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304	0.459346 0.643325 0.807949 0.965685 1.11235 μmol/min 0.01 0.01 0.04 0.05	0.668968 0.96649 1.248589 1.547566 1.783996 mg protein sample 0.06 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 156 222 501 587	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated μmol/min 0.07 0.04 0.06 0.06	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943	0.350998 0.515273 0.670777 0.815064 0.953058 μmol/min 0.01 0.03 0.05 0.05	0.02081 0.028369 0.037408 0.045741 0.056071 <b>mg protein sample</b> 0.06 0.11 0.11 0.15	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132	0.459346 0.643325 0.807949 0.965685 1.11235 µmol/min 0.01 0.01 0.04 0.05 0.05	0.668968 0.96649 1.248589 1.547566 1.783996 mg protein sample 0.06 0.08 0.08 0.09	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 1.56 222 501 587 542	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 24 h control 102.2 µg L <sup>-1</sup> 153.3 µg L <sup>-1</sup> 204.5 µg L <sup>-1</sup> 255.6 µg L <sup>-1</sup> 511.3 µg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295	0.350998 0.515273 0.670777 0.815064 0.953058 <b>µmol/min</b> 0.01 0.03 0.05 0.05 0.05	0.02081 0.028369 0.037408 0.045741 0.056071 mg protein sample 0.06 0.11 0.11 0.15 0.12	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132 1205	0.459346 0.643325 0.807949 0.965685 1.11235 <b>µmol/min</b> 0.01 0.01 0.04 0.05 0.05	0.668968 0.96649 1.248589 1.547566 1.783996 <b>mg protein sample</b> 0.06 0.08 0.08 0.09 0.09	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 156 222 501 587 542 1023	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 24 h control 102.2 µg L <sup>1</sup> 153.3 µg L <sup>1</sup> 204.5 µg L <sup>2</sup> 255.6 µg L <sup>2</sup> 113.3 µg L <sup>1</sup> control	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158	0.350998 0.515273 0.670777 0.815064 0.953058 <b>µmol/min</b> 0.01 0.03 0.05 0.05 0.02 0.13	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132 1205 261	0.459346 0.643325 0.807949 0.965685 1.11235 μmol/min 0.01 0.04 0.05 0.05 0.09 0.03	0.668968 0.96649 1.248589 1.547566 1.783996 mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 156 222 501 587 542 1023 324	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 511.3 μg L <sup>1</sup> τ control 102.2 μg L <sup>1</sup> 1512 μg L <sup>1</sup> 102.2 μg L <sup>1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431	0.350998 0.515273 0.670777 0.815064 0.953058 <b>µmol/min</b> 0.01 0.03 0.05 0.05 0.02 0.13 0.02	0.02081 0.028369 0.037408 0.045741 0.056071 mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132 1205 261 164	0.459346 0.643325 0.80794 0.965685 1.11235 µmol/min 0.01 0.04 0.05 0.05 0.05 0.09	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08	0.432442 0.618388 0.79022 0.955609 1.129218 nmol/min/mg prot 156 222 501 587 542 1023 324 346	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>2</sup> 155.6 μg L <sup>3</sup> 11.3 μg L <sup>3</sup> 11.3 μg L <sup>3</sup> 11.3 μg L <sup>3</sup> 153.3 μg L	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231	0.350998 0.515273 0.6707777 0.815064 0.953058 µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132 1205 261 164 218	0.459346 0.643325 0.807949 0.965685 1.11235 punol/min 0.01 0.01 0.04 0.05 0.05 0.09 0.03 0.02	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403	0.350998 0.515273 0.670777 0.815064 0.953058  µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08	0.31938 0.5383 0.743934 0.937396 1.122665 mmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> control 102.2 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 205.6 μg L <sup>-1</sup> 155.6	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.03 0.09 0.17 /	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403	0.350998 0.515273 0.670777 0.815064 0.953058 µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08	0.31938 0.5383 0.743934 0.937396 1.122665 mmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403	0.350998 0.515273 0.670777 0.815064 0.953058  µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08	0.31938 0.5383 0.743934 0.937396 1.122665 mmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> control 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 213.3 μg L <sup>-1</sup> 113.3 μg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.06 0.06 0.02 0.04 0.09 0.03 0.07	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.08 0.02 0.06 0.07 / 0.03	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038	0.350998 0.515273 0.670777 0.815064 0.953058  pmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.02 0.03 0.01 0.08	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08	0.31938 0.5383 0.743934 0.937396 1.122665 nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 511.3 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 255.6 μg L <sup>1</sup> 2151.3 μg L <sup>1</sup> 200.0000000000000000000000000000000000	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.08 0.02 0.06 0.07	0.327678 0.458036 0.581985 0.700259 0.814241 nm protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380	0.350998 0.515273 0.670777 0.815064 0.953058  µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03 0.01 0.08	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.10	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1511.3 μg L <sup>1</sup> 201.5 μg L <sup>1</sup> 102.2 μg L <sup>1</sup> 102.2 μg L <sup>1</sup> 102.2 μg L <sup>1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 /	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.17 0.03 0.09 0.07 /	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230	0.350998 0.515273 0.670777 0.815064 0.953058  pmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03 0.01 0.08 0.03 0.01	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.08 0.10 0.12	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 151.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 211.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 /	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.03 0.09 0.17 0.03 0.00 0.00 0.07 / 0.03 0.07 0.13 0.12	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248	0.350998 0.515273 0.670777 0.815064 0.953058  µmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03 0.01 0.08 0.03 0.01	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.08 0.10 0.12 0.012	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 2055.6 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 2055.6 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 2055.6 μg L <sup>1</sup> 151.3 μg L <sup>1</sup> 2055.6 μg L <sup>1</sup> 151.3 μg L <sup>1</sup> 2055.6 μg L <sup>1</sup> 151.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 255.6 μg L <sup>1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.04 0.03 0.03 0.03 0.05 0.05	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.01 0.03 0.07 0.03 0.07 0.13 0.12 0.10 0.13 0.110	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / / 1038 380 230 248 509 192 242	0.350998 0.515273 0.670777 0.815064 0.953058   pmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.03 0.01 0.08 0.03 0.03 0.04 0.02 0.03 0.04 0.02	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.10 0.12 0.06 0.16 0.16	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 205.6 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 205.6 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 201.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 201.5 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 201.5 μg L <sup>1</sup> 255.6 μg L <sup>1</sup> 1513.3 μg L <sup>1</sup> 201.5 μg L <sup>1</sup> 255.6 μg L <sup>1</sup> 1511.3 μg L <sup>1</sup> 200.1 μg L <sup>1</sup> 201.5 μg	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated pmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.04 0.03 0.05 0.03 0.05 0.03 0.03 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.06 0.07 / 0.03 0.07 0.13 0.12 0.10 0.13 0.10 0.08	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248 509 192 242 351	0.350998 0.515273 0.670777 0.815064 0.953058  pmol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.02 0.02 0.03 0.01 0.08 0.03 0.04 0.02 0.03 0.04 0.02 0.03	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.10 0.12 0.08 0.08 0.10 0.12 0.08 0.06 0.11 0.12 0.08	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459 258	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 2 min 3 min 4 min 5 min 4 min 5 min 24 h control 102.2 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 255.6 μg L <sup>1</sup> 511.3 μg L <sup>1</sup> 200.0 μg L <sup>1</sup> 153.3 μg L <sup>1</sup> 204.5 μg L <sup>1</sup> 151.3 μg L <sup>1</sup> 205.6 μg L <sup>1</sup> 1511.3 μg L <sup>1</sup> 200.0 μg L <sup>1</sup> 1511.3 μg L <sup>1</sup> 200.0 μg L <sup>1</sup> 1511.3 μg L <sup>1</sup> 102.2 μg L <sup>1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.03 0.03 0.03 0.03 0.03 0.03 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.01 0.03 0.09 0.01 0.07 / 0.03 0.07 0.13 0.12 0.10 0.13 0.11 0.10 0.08	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248 509 192 242 351 400	0.350998 0.515273 0.670777 0.815064 0.953058  punol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.03 0.01 0.08 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.008 0.008 0.008 0.10 0.12 0.08 0.06 0.16 0.16 0.16 0.11 0.08	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459 258 306	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>3</sup> 113.3 µg L <sup>3</sup> 204.5 µg L <sup>3</sup> 153.3 µg L	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated punol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.04 0.03 0.05 0.03 0.05 0.03 0.05 0.03 0.05	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.01 0.03 0.09 0.12 0.01 0.01 0.01 0.01 0.01 0.01 0.01	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248 509 192 242 351 400 456	0.350998 0.515273 0.670777 0.815064 0.953058  punol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.03 0.01 0.08 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.06	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.08 0.10 0.12 0.08 0.06 0.16 0.12 0.11 0.07	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459 258 306 340	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 2255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 102.2 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup> 211.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 2153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 153.3 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup>	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated µmol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.04 0.03 0.03 0.03 0.03 0.03 0.03 0.03	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.03 0.03 0.09 0.12 0.13 0.10 0.13 0.11 0.11 0.13 0.10 0.08 0.05 0.05 0.12 0.10	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248 509 192 242 351 400 456 476	0.350998 0.515273 0.670777 0.815064 0.953058  punol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.03 0.01 0.08 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.06 0.03 0.06 0.03 0.02 0.02	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.08 0.08 0.10 0.12 0.01 0.12 0.08 0.06 0.16 0.12 0.11 0.08 0.06	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459 258 306 340 202	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655
1 min 2 min 2 min 3 min 4 min 5 min 4 min 5 min 102.2 µg L <sup>1</sup> 153.3 µg L <sup>2</sup> 204.5 µg L <sup>3</sup> 113.3 µg L <sup>3</sup> 204.5 µg L <sup>3</sup> 153.3 µg L	0.219327 0.433335 0.631108 0.814218 0.986821 Calculated punol/min 0.07 0.04 0.06 0.06 0.02 0.04 0.09 0.03 0.07 0.03 / 0.04 0.03 0.05 0.03 0.05 0.03 0.05 0.03 0.05	0.266947 0.473167 0.664827 0.844545 1.014197 values normalised of mg protein sample 0.08 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.03 0.09 0.12 0.01 0.03 0.09 0.12 0.01 0.01 0.01 0.01 0.01 0.01 0.01	0.327678 0.458036 0.581985 0.700259 0.814241 on protein content nmol/min/mg prot 833 1257 689 489 943 1295 1158 1431 1231 403 / 1038 380 230 248 509 192 242 351 400 456	0.350998 0.515273 0.670777 0.815064 0.953058  punol/min 0.01 0.03 0.05 0.05 0.02 0.13 0.02 0.02 0.03 0.01 0.08 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.04 0.02 0.03 0.06	0.02081 0.028369 0.037408 0.045741 0.056071  mg protein sample 0.06 0.11 0.11 0.15 0.12 0.11 0.07 0.14 0.09 0.08 0.08 0.08 0.08 0.10 0.12 0.08 0.06 0.16 0.12 0.11 0.07	0.31938 0.5383 0.743934 0.937396 1.122665  nmol/min/mg prot 157 221 432 304 132 1205 261 164 218 365 184 1034 313 217 456 309 180 459 258 306 340	0.459346 0.643325 0.807949 0.965685 1.11235  pmol/min 0.01 0.04 0.05 0.05 0.09 0.03 0.02 0.07 0.04 0.01	0.668968 0.96649 1.248589 1.547566 1.783996  mg protein sample 0.06 0.08 0.08 0.09 0.09 0.09 0.09 0.08 0.08	0.432442 0.618388 0.79022 0.955609 1.129218  nmol/min/mg prot 156 222 501 587 542 1023 324 346 543 374 133	0.12378 0.14482 0.1651 0.18655

Table 31: GST activity in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h. Representative absorbance (AU) values of whole 5 min measurement (selected for every min) and data normalised on total protein content (nmol/min/mg protein). (n=10), (continued).

GST										
24 h - C										
Control										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.146236	0.042268	0.118228	-0.025388	0.238809	0.225794	0.268316	0.37774	0.256141	0.13408
2 min	0.215348	0.092	0.180029	0.015482	0.32088	0.265945	0.313015	0.462245	0.24998	0.21085
3 min 4 min	0.277997 0.333979	0.13941 0.179978	0.234261 0.279339	0.054798 0.102651	0.348072 0.3573	0.387399 0.409343	0.42122 0.44673	0.518991 0.563051	0.249274 0.362658	0.33364 0.38016
5 min	0.384914	0.21773	0.322515	0.102031	0.457125	0.444866	0.424734	0.593668	0.377037	0.40418
102.2 μg L <sup>-1</sup>			***	*******	*****		****			
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.12955	0.128328	0.149784	0.156438	0.340086	0.646463	0.736234	0.447378	1.094399	0.48672
2 min	0.196323	0.200735	0.216454	0.227732	0.441932	0.856097	1.082817	0.805987	1.072066	0.67022
3 min	0.252649	0.260363	0.281522	0.28501	0.621114	1.02245	1.620269	1.267982	1.426711	0.83406
4 min	0.29972	0.299787	0.332891	0.340268	0.837298	1.066789	2.06054	1.509875	1.714141	1.06496
5 min	0.339826	0.363025	0.37819	0.384669	1.694785	1.086095	2.741782	2.569846	2.112663	1.15584
153.3 μg L <sup>-1</sup> time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.4604	0.4584	0.4529	0.8663	0.481	0.792	0.198045	0.1909	0.2426	0.20689
2 min	0.6432	0.552	0.6775	1.0951	0.667	0.9934	0.349859	0.2552	0.3258	0.33563
3 min	0.8153	0.6526	0.9255	1.3347	0.8494	1.1812	0.434303	0.3208	0.4047	0.45678
4 min	0.9551	0.7609	1.0706	1.5821	1.0103	1.347	0.548586	0.389	0.4888	0.57123
5 min	1.0785	0.9093	1.1767	1.818	1.1224	1.4802	0.693385	0.4573	0.5768	0.6896
204.5 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	0.8046	0.4861	0.237564	0.293934	0.1423	0.0924	0.0802	0.5305	0.4554	0.4582
2 min 3 min	1.0142 1.2352	0.7114 0.9174	0.417888 0.523421	0.487141 0.623672	0.1499 0.1538	0.0936 0.0958	0.085 0.0891	0.9098 1.0502	0.8349 1.0258	0.7232 0.9712
4 min	1.452	1.1393	0.64742	0.790631	0.1623	0.0958	0.0831	1.2966	1.1768	1.05892
5 min	1.6715	1.3467	0.800012	0.975637	0.17	0.0976	0.0981	1.6087	1.5041	1.3239
255.6 μg L <sup>-1</sup>										
time	AU	AU	AU	AU	AU	AU	AU	AU	AU	AU
1 min	1.362637	0.143184	0.133422	0.167281	0.705267	0.517212	0.584282	0.615233	0.533033	0.69791
2 min	1.431397	0.193688	0.213913	0.224842	0.971772	0.721539	0.861622	0.854004	0.744021	1.0215
3 min	1.500165	0.23797	0.282359	0.272417	1.297737	0.929441	1.133561	1.146285	0.93774	1.38988
4 min	1.589687	0.27838	0.343447	0.31585	1.509149	1.151035	1.367819	1.45848	1.161672	2.08173
5 min	1.619464	0.313766	0.408222	0.354798	1.76466	1.296845	1.800128	1.679468	1.375759	2.25647
511.3 μg L <sup>-1</sup> time	ATT	AIT	AIT	AU	ATT	477	A 77	ATT	AII	ATT
lime 1 min	AU 0.252108	AU 0.162706	AU 0.235301	0.185771	AU 0.45971	AU 0.487406	AU 0.746899	AU 0.413927	AU 0.454748	AU 0.27377
2 min	0.313766	0.204598	0.330724	0.253613	0.665818	0.708818	1.020324	0.572275	0.680329	0.29711
3 min	0.369359	0.242534	0.414906	0.310934	0.847307	0.91868	1.87416	0.722131	0.829636	0.37389
4 min	0.415519	0.275198	0.486956	0.361553	1.014473	1.147863	2.360805	0.863753	1.214336	0.44814
5 min	0.459547	0.307573	0.552477	0.419567	1.170572	1.384937	3.28585	1.446372	1.623092	0.45919
	Calculated	values normalised	on protein conten	t						
24 h	μmol/min	mg protein sample	nmol/min/mg prot	μmol/min	mg protein sample	nmol/min/mg prot	μmol/min	mg protein sample	nmol/min/mg prot	
control	0.07	0.08	833	0.01	0.06	157	0.01	0.06	156	
102.2 μg L <sup>-1</sup>	0.01	0.09	112	0.05	0.07	763	0.04	0.08	599	
153.3 μg L <sup>-1</sup>	0.04	0.08	523	0.03	0.07	382	0.01	0.12	105	
204.5 μg L <sup>-1</sup>	0.05	0.07	654	0.03	0.07	398	0.01	0.03	346	
255.6 μg L <sup>-1</sup>	0.01	0.09	155	0.04	0.06	700	0.03	0.06	588	
511.3 μg L <sup>-1</sup>	0.01	0.06	156	0.03	0.07	454	0.05	0.06	822	
control	0.09	0.08	1158	0.02	0.07	261	0.03	0.08	324	
102.2 μg L <sup>-1</sup>	0.01	0.08	144	0.01	0.08	153	0.03	0.05	561	
153.3 μg L <sup>-1</sup>	0.05	0.07	654	0.02	0.09	255	0.03	0.04	607	
204.5 μg L <sup>-1</sup>	0.04	0.06	702	0.07	0.09	765	0.04	0.08	456	
255.6 μg L <sup>-1</sup>	0.01	0.08	105	0.03	0.06	514	0.08	0.08	970	
511.3 μg L <sup>-1</sup>	0.01	0.07	95	0.04	0.06	626	0.02	0.07	322	
control	0.03	0.07	380	0.03	0.10	313				
102.2 μg L <sup>-1</sup>	0.02	0.13	190	0.09	0.12	776				
153.3 μg L <sup>-1</sup>	0.06	0.11	571	0.05	0.10	502				
204.5 μg L <sup>-1</sup>	0.05	0.09	504	0.07	0.12	601				
255.6 μg L <sup>-1</sup>	0.03	0.13	208	0.05	0.06	805				
511.3 μg L <sup>-1</sup>	0.02	0.14	113	0.12	0.08	1638				
control	0.02	0.08	351	0.03	0.11	258				
102.2 μg L <sup>-1</sup>	0.01	0.08	142	0.09	0.12	802				
153.3 μg L <sup>-1</sup>	0.03	0.09	312	0.02	0.09	201				
204.5 μg L <sup>-1</sup>	0.04	0.08	502	0.05	0.08	573				
255.6 μg L <sup>-1</sup>	0.05	0.10	495	0.05	0.08	594				
511.3 μg L <sup>-1</sup>	0.09	0.13	687	0.04	0.06	597				
511.5 μg L	0.07	0.13	007	0.04	0.00	371				

Table 32: Absorbance (AU) values for LP measurements in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h and data normalised on total protein content (AU final), n=10.

LP									
24 h - A									
	AU	AU	AU	AU final		AU	AU	AU	${m AUfinal}$
control	535 nm 0.065	600 nm 0.025	280 nm 0.28	0.14	control	535 nm 0.0664	600 nm 0.04	280 nm 0.24	0.11
control	0.039	0.012	0.287	0.09	control	0.0004	0.06	0.192	0.11
control	0.07164	0.009	0.261	0.24	control	0.15072	0.12	0.192	0.16
control	0.11328	0.08	0.208	0.16	control	0.1266	0.09	0.183	0.2
control 102.2 μg L <sup>-1</sup>	0.04808	0.032	0.201	0.08	control	0.11768	0.09	0.173	0.16
102.2 μg L 102.2 μg L <sup>-1</sup>	0.03792 0.0438	0.014 0.015	0.184 0.18	0.13 0.16	102.2 μg L <sup>-1</sup> 102.2 μg L <sup>-1</sup>	0.0164 0.09496	0.006 0.07	0.08 0.096	0.13 0.26
102.2 μg L 102.2 μg L <sup>-1</sup>	0.0438	0.013	0.185	0.10	102.2 μg L <sup>-1</sup>	0.09490	0.07	0.098	0.20
102.2 μg L <sup>-1</sup>	0.23571	0.016	0.173	1.27	102.2 μg L <sup>-1</sup>	0.06452	0.05	0.121	0.12
102.2 μg L <sup>-1</sup>	0.07171	0.05	0.167	0.13	102.2 μg L <sup>-1</sup>	0.13058	0.05	0.102	0.79
153.3 μg L <sup>-1</sup>	0.04838	0.011	0.267	0.14	153.3 μg L <sup>-1</sup>	0.08936	0.01	0.248	0.32
153.3 μg L <sup>-1</sup>	0.02204	0.008	0.156	0.09	153.3 μg L <sup>-1</sup>	0.0394	0.001	0.384	0.1
153.3 $\mu g L^{-1}$	0.05309	0.008	0.167	0.27	$153.3~\mu g~L^{-1}$	0.02072	0.002	0.234	0.08
153.3 $\mu g L^{-1}$	0.07132	0.007	0.201	0.32	153.3 μg L <sup>-1</sup>	0.04972	0.001	0.203	0.24
153.3 μg L <sup>-1</sup>	0.09617	0.05	0.243	0.19	153.3 μg L <sup>-1</sup>	0.0403	0.007	0.222	0.15
204.5 μg L <sup>-1</sup>	0.06175	0.014	0.191	0.25	204.5 μg L <sup>-1</sup>	0.06192	0.012	0.312	0.16
204.5 μg L <sup>-1</sup>	0.09889	0.003	0.223	0.43	204.5 μg L <sup>-1</sup>	0.0628	0.004	0.28	0.21
204.5 μg L <sup>-1</sup>	0.03163	0.013	0.207	0.09	204.5 μg L <sup>-1</sup>	0.00968	0.005	0.234	0.02
204.5 μg L <sup>-1</sup>	0.03338	0.004	0.226	0.13	204.5 μg L <sup>-1</sup>	0.03036	0.006	0.203	0.12
204.5 μg L <sup>-1</sup>	0.03616	0.004	0.201	0.16	204.5 μg L <sup>-1</sup>	0.12314	0.1	0.178	0.13
255.6 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup>	0.046 0.035	0.017	0.195	0.15	255.6 μg L <sup>-1</sup> 255.6 μg L <sup>-1</sup>	0.06028 0.0404	0.003	0.179	0.32
255.6 μg L <sup>-1</sup>	0.033	0.011 0.07	0.195 0.198	0.12 0.43	255.6 μg L <sup>-1</sup>	0.0404	0.002 0.06	0.192 0.203	0.2 0.18
255.6 μg L <sup>-1</sup>	0.13314	0.07	0.194	0.36	255.6 μg L <sup>-1</sup>	0.0708	0.03	0.203	0.18
255.6 μg L <sup>-1</sup>	0.08328	0.06	0.194	0.12	255.6 μg L <sup>-1</sup>	0.03764	0.002	0.297	0.12
511.3 μg L <sup>-1</sup>	0.05136	0.021	0.276	0.11	511.3 μg L <sup>-1</sup>	0.0516	0.006	0.24	0.19
511.3 μg L <sup>-1</sup>	0.081	0.015	0.2	0.33	511.3 μg L <sup>-1</sup>	0.04532	0.011	0.264	0.13
511.3 μg L <sup>-1</sup>	0.0312	0.004	0.16	0.17	511.3 μg L <sup>-1</sup>	0.03342	0.005	0.203	0.14
$511.3 \mu g L^{-1}$	0.05136	0.021	0.276	0.11	511.3 μg L <sup>-1</sup>	0.03842	0.008	0.234	0.13
$511.3~\mu g~L^{-1}$	0.05053	0.002	0.211	0.23	511.3 μg L <sup>-1</sup>	0.05195	0.02	0.213	0.15
24 h - B									
	AU	AU	AU	AU final		AU	AU	AU	AU final
$102.2~\mu\mathrm{g~L^{-1}}$	535 nm	600 nm	280 nm	0.15	102.2 μg L <sup>-1</sup>	535 nm 0.01	600 nm	280 nm	0.07
102.2 μg L 102.2 μg L <sup>-1</sup>	0.021 0.0164	0.006 0.002	0.098 0.18	0.15 0.08	102.2 μg L 102.2 μg L <sup>-1</sup>	0.01	0.002 0.07	0.112 0.096	0.07 0.1
102.2 μg L <sup>-1</sup>	0.01318	0.002	0.174	0.07	102.2 μg L <sup>-1</sup>	0.017	0.002	0.209	0.07
102.2 μg L <sup>-1</sup>	0.01288	0.001	0.132	0.09	102.2 μg L <sup>-1</sup>	0.05968	0.05	0.121	0.08
102.2 μg L <sup>-1</sup>	0.06503	0.05	0.167	0.09	102.2 μg L <sup>-1</sup>	0.008	0.004	0.2	0.02
153.3 $\mu g L^{-1}$	0.02435	0.011	0.267	0.05	153.3 μg L <sup>-1</sup>	0.008	0.01	0.248	0.08
153.3 $\mu g L^{-1}$	0.0158	0.008	0.156	0.05	153.3 μg L <sup>-1</sup>	0.008	0.001	0.384	0.14
153.3 $\mu g L^{-1}$	0.0247	0.008	0.167	0.1	153.3 μg L <sup>-1</sup>	0.008	0.002	0.234	0.133
153.3 μg L <sup>-1</sup>	0.01102	0.007	0.201	0.02	153.3 μg L <sup>-1</sup>	0.008	0.001	0.203	0.012
153.3 μg L <sup>-1</sup>	0.05729	0.05	0.243	0.03	153.3 μg L <sup>-1</sup>	0.008	0.007	0.222	0.12
204.5 μg L <sup>-1</sup>	0.01782	0.014	0.191	0.02	204.5 μg L <sup>-1</sup>	0.008	0.012	0.312	0.08
204.5 μg L <sup>-1</sup>	0.01192	0.003	0.223	0.04	204.5 μg L <sup>-1</sup>	0.008	0.004	0.28	0.08
204.5 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup>	0.03163	0.013	0.207	0.09	204.5 μg L <sup>-1</sup>	0.008	0.005	0.234	0.07
204.5 μg L 204.5 μg L <sup>-1</sup>	0.00852 0.02812	0.004 0.004	0.226	0.02 0.12	204.5 μg L <sup>-1</sup> 204.5 μg L <sup>-1</sup>	0.008	0.006 0.1	0.203	0.02 0.09
255.6 μg L <sup>-1</sup>	0.02812	0.004	0.201 0.205	0.12	255.6 μg L <sup>-1</sup>	0.008	0.001	0.178 0.149	0.08
255.6 μg L <sup>-1</sup>	0.013	0.002	0.195	0.06	255.6 μg L <sup>-1</sup>	0.013	0.001	0.124	0.08
255.6 μg L <sup>-1</sup>	0.0227	0.011	0.0472	0.76	255.6 μg L <sup>-1</sup>	0.009	0.002	0.11	0.07
255.6 μg L <sup>-1</sup>	0.01	0.004	0.107	0.05	255.6 μg L <sup>-1</sup>	0.009	0.001	0.11	0.72
255.6 μg L <sup>-1</sup>	0.0988	0.06	0.194	0.2	255.6 μg L <sup>-1</sup>	0.30494	0.002	0.297	1.02
511.3 μg L <sup>-1</sup>	0.021	0.004	0.149	0.11	511.3 μg L <sup>-1</sup>	0.015	0.002	0.151	0.09
511.3 $\mu g L^{-1}$	0.025	0.004	0.115	0.18	511.3 μg L <sup>-1</sup>	0.02624	0.011	0.254	0.06
511.3 μg L <sup>-1</sup>	0.0216	0.004	0.16	0.11	511.3 μg L <sup>-1</sup>	0.0165	0.005	0.23	0.05
511.3 μg L <sup>-1</sup>	0.015	0.002	0.194	0.07	511.3 μg L <sup>-1</sup>	0.03572	0.008	0.231	0.12
511.3 μg L <sup>-1</sup>	0.01	0.001	0.147	0.06	511.3 μg L <sup>-1</sup>	0.04343	0.02	0.213	0.11

Table 32: Absorbance (AU) values for LP measurements in G. fossarum exposed to IMI (A), Confidor 200SL (B) and 6CNA (C) for 24 h and data normalised on total protein content (AU final), n=10, (continued).

## LP 24 h - C

24 h - C									
	AU	AU	AU	AU final		AU	AU	AU	AU final
	535 nm	600 nm	280 nm			535 nm	600 nm	280 nm	
control	0.065	0.025	0.28	0.14	control	0.0664	0.04	0.24	0.11
control	0.039	0.012	0.287	0.09	control	0.0792	0.06	0.192	0.1
control	0.07164	0.009	0.261	0.24	control	0.15072	0.12	0.192	0.16 0.2
control control	0.11328 0.04808	0.08 0.032	0.208 0.201	0.16 0.08	control control	0.1266 0.11768	0.09 0.09	0.183 0.173	0.2 0.16
102.2 μg L <sup>-1</sup>	0.04808	0.006	0.2382	0.06	102.2 μg L <sup>-1</sup>	0.02	0.002	0.175	0.10
102.2 μg L <sup>-1</sup>	0.05001	0.001	0.2883	0.17	102.2 μg L <sup>-1</sup>	0.0384	0	0.192	0.2
102.2 μg L <sup>-1</sup>	0.0717	0.008	0.245	0.26	102.2 μg L <sup>-1</sup>	0.014	0.001	0.13	0.1
102.2 μg L <sup>-1</sup>	0.0197	0	0.197	0.1	102.2 μg L <sup>-1</sup>	0.004	0.001	0.161	0.01
$102.2~\mu g~L^{-1}$	0.008	0	0.185	0.04	$102.2~\mu g~L^{\text{-}1}$	/	/	/	1
$153.3~\mu g~L^{\text{-}1}$	0.00418	0.002	0.218	0.01	$153.3~\mu\mathrm{g~L}^{\text{-}1}$	0.0053	0.002	0.165	0.02
$153.3~\mu g~L^{-1}$	0.03372	0.002	0.244	0.13	153.3 $\mu g L^{-1}$	0.0518	0.01	0.209	0.2
153.3 $\mu g L^{-1}$	0.001	0.001	0.212	0	153.3 $\mu g L^{-1}$	0.02082	0.003	0.198	0.09
$153.3~\mu g~L^{-1}$	0.04232	0.002	0.288	0.14	153.3 $\mu g L^{-1}$	0.05	0.05	0.184	0
$153.3~\mu g~L^{-1}$	0.03743	0	0.197	0.19	153.3 $\mu g L^{-1}$	0.03812	0.03	0.203	0.04
$204.5~\mu g~L^{-1}$	0.01611	0	0.179	0.09	$204.5 \ \mu g \ L^{-1}$	0.05656	0.05	0.164	0.04
$204.5~\mu g~L^{-1}$	0.00382	0	0.191	0.02	$204.5~\mu g~L^{\text{-}1}$	0.03107	0.02	0.123	0.09
$204.5~\mu g~L^{-1}$	0.02029	0.003	0.133	0.13	$204.5 \ \mu g \ L^{-1}$	0.09248	0.08	0.156	0.08
204.5 μg L <sup>-1</sup>	0.02004	0.001	0.136	0.14	$204.5 \ \mu g \ L^{-1}$	0.01972	0.001	0.208	0.09
$204.5 \ \mu g \ L^{-1}$	0.0157	0.002	0.137	0.1	$204.5 \ \mu g \ L^{-1}$	0.00498	0.001	0.199	0.02
255.6 $\mu g L^{-1}$	0.009	0.001	0.208	0.04	255.6 $\mu g L^{-1}$	0.02392	0	0.184	0.13
255.6 $\mu g L^{-1}$	0.007	0.002	0.265	0.02	255.6 $\mu g L^{-1}$	0.0163	0.002	0.143	0.1
255.6 $\mu g L^{-1}$	0.004	0.002	0.264	0.08	$255.6 \ \mu g \ L^{-1}$	0.005	0	0.181	0.03
255.6 $\mu g L^{-1}$	0.007	0.002	0.265	0.01	255.6 $\mu g L^{-1}$	0.03277	0.03	0.0231	0.12
$255.6~\mu\mathrm{g~L}^{-1}$	0.0434	0.02	0.26	0.09	255.6 μg L <sup>-1</sup>	0.002	0	0.05	0.04
511.3 μg $L^{-1}$	0.005	0.002	0.257	0.01	$511.3 \mu g L^{-1}$	0.009	0	0.1799	0.05
$511.3~\mu g~L^{-1}$	0.004	0.001	0.239	0.01	$511.3 \mu g L^{-1}$	0.014	0	0.21	0.07
$511.3~\mu g~L^{-1}$	0.004	0.001	0.258	0.01	$511.3~\mu g~L^{-1}$	0	0	0.217	0
$511.3~\mu g~L^{-1}$	0.00305	0.001	0.205	0.01	$511.3~\mu g~L^{-1}$	0.003	0	0.26	0.01
$511.3~\mu g~L^{-1}$	0.006	0	0.206	0.03	$511.3~\mu g~L^{-1}$	0	0	0.199	0

Table 33: Values of AChE, CAT and GST activity and LP measurements in G. fossarum exposed to lower concentrations of IMI/Confidor 200SL  $(6.3-51.1~\mu g~L^{-1})$  and 6CNA  $(3.9-31.4~\mu g~L^{-1})$  for 24 h. Data are presented as values normalised on total protein content; AChE and CAT  $(\mu mol/min/mg~protein)$ , GST (nmol/min/mg~protein) and LP (AU). (n=10).

# G. fossarum n=10

n = 10									
Mean values					Raw data				
IMI/Confidor 200SL					6CNA				
$ m \mu g~L^{-1}$	AChE	CAT	GST	LP	$\mu \mathrm{g}~\mathrm{L}^{ ext{-}1}$	AChE	CAT	GST	LP
control	71.17	19.06	427.10	0.22	3.9	71.68	19.98	413.02	0.22
control	70.82	19.52	422.03	0.18	3.9	71.82	20.74	404.56	0.19
control	70.59	21.05	416.21	0.21	3.9	71.98	22.21	403.74	0.20
control	70.96	21.81	417.68	0.21	3.9	70.73	20.36	418.50	0.23
control	71.56	22.07	417.87	0.20	3.9	71.34	22.35	411.47	0.19
control	71.12	22.59	401.71	0.21	3.9	70.73	20.12	430.45	0.21
control	70.41	20.92	407.89	0.18	3.9	70.86	22.47	418.61	0.21
control	71.02	19.14	421.68	0.22	3.9	71.78 70.75	21.25	401.11	0.22 0.23
control	71.51	19.06	421.22	0.20	3.9 3.9		19.64	429.41	
control DMSO+NMP	71.37 71.67	21.25 20.15	420.42 431.33	0.21 0.22	7.8	70.54 71.58	20.96 22.01	407.59 399.10	0.18 0.22
DMSO+NMP	71.07	19.71	426.94	0.22	7.8 7.8	71.38	21.83	420.90	0.22
DMSO+NMP	70.27	19.71	411.76	0.23	7.8	70.00	21.68	405.41	0.21
DMSO+NMP	68.74	20.65	420.03	0.22	7.8	71.34	21.44	410.68	0.19
DMSO+NMP	68.50	19.25	430.27	0.23	7.8	71.72	19.69	402.85	0.21
DMSO+NMP	70.23	20.78	401.70	0.22	7.8	71.31	22.07	416.86	0.20
DMSO+NMP	70.75	19.38	400.58	0.21	7.8	70.36	21.14	430.15	0.20
DMSO+NMP	70.49	21.63	417.07	0.21	7.8	70.47	21.78	429.14	0.21
6.3	71.90	22.10	403.48	0.18	7.8	70.36	19.33	414.66	0.22
6.3	71.92	22.33	420.24	0.18	7.8	71.14	21.14	419.08	0.21
6.3	70.09	21.72	410.12	0.19	15.7	70.92	19.60	416.28	0.23
6.3	70.06	21.77	403.10	0.19	15.7	70.28	21.30	428.16	0.23
6.3	71.71	20.17	409.34	0.19	15.7	71.17	21.48	407.42	0.21
6.3	70.70	19.93	403.79	0.19	15.7	71.32	20.26	416.09	0.23
6.3	70.43	21.23	406.35	0.20	15.7	71.24	22.05	419.91	0.22
6.3	70.10	21.88	407.96	0.23	15.7	70.78	21.23	411.91	0.22
6.3	70.22	20.39	402.38	0.19	15.7	71.59	21.45	416.39	0.20
6.3	70.79	21.62	413.96	0.18	15.7	70.80	22.21	409.72	0.22
12.7 12.7	70.61 71.03	19.22 20.36	405.72 399.98	0.21 0.21	15.7 15.7	71.64 71.52	19.86 20.76	401.17 422.52	0.21 0.19
12.7	70.15	20.30	425.08	0.21	31.4	71.32	21.65	398.82	0.19
12.7	71.90	21.78	426.87	0.10	31.4	70.64	21.99	422.53	0.23
12.7	70.91	22.87	412.12	0.18	31.4	70.90	21.36	418.10	0.22
12.7	71.21	22.06	425.85	0.20	31.4	70.79	22.83	413.89	0.20
12.7	71.74	22.45	429.09	0.20	31.4	71.05	21.15	411.08	0.18
12.7	70.95	21.73	427.49	0.20	31.4	71.40	20.77	406.40	0.22
12.7	71.00	20.76	423.08	0.18	31.4	70.94	21.48	406.65	0.23
12.7	71.03	19.11	416.61	0.23	31.4	71.19	22.91	410.36	0.19
25.5	71.38	22.50	404.97	0.18	31.4	70.01	19.89	399.44	0.23
25.5	70.13	19.16	410.94	0.18	31.4	71.50	21.58	410.95	0.22
25.5	71.45	19.44	416.23	0.22					
25.5	71.64	21.09	407.85	0.21					
25.5 25.5	70.58	21.02	399.60	0.22					
25.5 25.5	71.48 71.77	21.39 19.23	411.15	0.22 0.22					
25.5 25.5	70.67	19.23	398.98 430.80	0.22					
25.5	70.84	20.78	423.69	0.13					
25.5	71.96	19.80	412.25	0.18					
51.1	70.62	20.44	427.62	0.21					
51.1	70.26	22.42	426.68	0.18					
51.1	70.70	20.42	413.06	0.19					
51.1	70.29	21.83	414.38	0.22					
51.1	70.76	22.08	414.35	0.19					
51.1	71.19	20.49	400.77	0.23					
51.1	71.15	22.17	418.82	0.22					
51.1	71.63	19.29	415.75	0.18					
51.1	70.88	21.92	401.69	0.19					
51.1	71.09	20.90	401.68	0.21					

Table 34: Fluorescence (FU) values for MXR activity measurements in G. fossarum exposed to IMI/Confidor 200SL (0.7 and 7.6 mg  $L^{-1}$ ) or 6CNA (0.4 and 4.7 mg  $L^{-1}$ ) and copper (0.75, 3 and 10  $\mu$ g  $L^{-1}$ ) for 24 h and data normalised on total animal's weight (n=3).

MXR						
Chemical (mg)	FU	FU	FU	Mean FU	Weight (mg)	FU/mg
control	5859	5938	5985	5927.33	26	227.974
control	9231	9077	8757	9021.67	23	392.246
control	6299	6762	6620	6560.33	26	252.321
control	4607	4846	4826	4759.67	35	135.99
control	5569	5392	5694	5551.67	27	205.617
CYC CYC	18099	17825 15206	18057 14453	17993.67 14633.33	35 23	514.105 636.232
CYC	14241 9457	10253	10524	10078.00	18	559.889
CYC	6714	6960	6846	6840.00	16	427.5
CYC	11235	12231	11266	11577.33	21	551.302
VER	7025	6429	7056	6836.67	14	488.333
VER	6916	6946	6955	6939.00	26	266.885
VER	11431	11193	12274	11632.67	20	581.633
VER VER	6384 7254	6267 7159	6304 7043	6318.33 7152.00	5 13	1263.67 550.154
IMI 0.7	4846	4775	5012	4877.67	25	195.107
IMI 0.7	4256	3896	4091	4081.00	25	163.24
IMI 0.7	5748	6579	7083	6470.00	17	380.588
IMI 0.7	8567	8041	7973	8193.67	21	390.175
IMI 0.7	6360	5969	5975	6101.33	22	277.333
IMI 7.6 IMI 7.6	4510	4148	4496	4384.67 8418.33	21 34	208.794 247.598
IMI 7.6	8431 8436	8709 8423	8115 8201	8353.33	36	232.037
IMI 7.6	7531	7461	7574	7522.00	24	313.417
IMI 7.6	5174	4638	5072	4961.33	23	215.71
CONFIDOR 0.7	4846	4775	5012	4877.67	31	157.344
CONFIDOR 0.7	4256	3896	4091	4081.00	41	99.5366
CONFIDOR 0.7	5748	6579	7083 7973	6470.00	32 23	202.188 356.246
CONFIDOR 0.7 CONFIDOR 0.7	8567 6360	8041 5969	7973 5975	8193.67 6101.33	23 32	190.667
CONFIDOR 7.6	7849	8797	8108	8251.33	17	485.373
CONFIDOR 7.6	13280	13531	12829	13213.33	42	314.603
CONFIDOR 7.6	10997	11237	11196	11143.33	38	293.246
CONFIDOR 7.6	9958	10005	10282	10081.67	41	245.894
CONFIDOR 7.6 6CNA 0.4	4333 18099	7676 17825	7415 18057	6474.67 17993.67	34 22	190.431 817.894
6CNA 0.4	14241	15206	14453	14633.33	36	406.481
6CNA 0.4	9457	10253	10524	10078.00	23	438.174
6CNA 0.4	6714	6960	6846	6840.00	28	244.286
6CNA 0.4	11235	12231	11266	11577.33	14	826.952
6CNA 4.7	7025	6429	7056	6836.67	35	195.333
6CNA 4.7 6CNA 4.7	6916 11431	6946 11193	6955 12274	6939.00 11632.67	26 36	266.885 323.13
6CNA 4.7	6384	6267	6304	6318.33	28	225.655
6CNA 4.7	7254	7159	7043	7152.00	21	340.571
Cu 3 μg L <sup>-1</sup>	7398	7606	7601	7535.00	38	198.289
Cu 3 μg L <sup>-1</sup>	10354	9455	9452	9753.67	35	278.676
Cu 3 μg L <sup>-1</sup>	9202	9305	9769	9425.33	33	285.616
Cu 3 μg L <sup>-1</sup>	8443	7696	12535	9558.00	41	233.122
Cu 3 μg L <sup>-1</sup>	6007	5404	7343	6251.33	20	312.567
Cu 10 μg L <sup>-1</sup>	4510	4148	4496	4384.67	41	106.943
Cu 10 μg L <sup>-1</sup>	8431	8709	8115	8418.33	23	366.014
Cu 10 μg L <sup>-1</sup>	8436	8423	8201	8353.33	35	238.667
Cu 10 μg L <sup>-1</sup>	7531	7461	7574	7522.00	32	235.063
Cu 10 μg L <sup>-1</sup>	5174	4638	5072	4961.33	23	215.71
IMI $0.7 + \text{Cu } 3  \mu\text{g L}^{-1}$	7503	6372	7149	7008.00	32	219
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	11172	10955	11097	11074.67	23	481.507
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	2824	10313	8460	7199.00	36	199.972
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	9635	10147	10409	10063.67	25	402.547
IMI $0.7 + \text{Cu } 3 \text{ µg L}^{-1}$	7263	7466	7707	7478.67	23	325.159
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	8773	8864	9420	9019.00	28	322.107
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	11518	11849	17280	13549.00	20	677.45
IMI $0.7 + \text{Cu } 10 \text{ µg L}^{-1}$	13028	12474	20861	15454.33	32	482.948
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	9671	10437	10533	10213.67	27	378.284
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	7975	8231	8486	8230.67	25	329.227
			00	,		/

Table 34: Fluorescence (FU) values for MXR activity measurements in G. fossarum exposed to IMI/Confidor 200SL (0.7 and 7.6 mg  $L^{-1}$ ) or 6CNA (0.4 and 4.6 mg  $L^{-1}$ ) and copper (0.75, 3 and 10  $\mu$ g  $L^{-1}$ ) for 24 h and data normalised on total animal's weight (n=3), (continued).

MXR						
II						
Chemical (mg)	FU	FU	FU	Mean FU	Weight (mg)	FU/mg
control	7542	7597	7530	7556.33	35	215.895
control	7811	7439	7413	7554.33	29	260.494
control	10523	10553	12116	11064.00	35	316.114
control	10934	10958	11261	11051.00	26	425.038
control	5955	5825	6577	6119.00	27	226.63
CYC	16693	15812	12700	15068.33	26	579.551
CYC	13912	12901	12726	13179.67	27	488.136
CYC	14576	14988	14940	14834.67	34	436.314 209.333
CYC CYC	7085 12863	6563 12024	7076 11792	6908.00 12226.33	33 18	679.241
VER	7076	7260	7659	7331.67	14	523.69
VER	6788	6610	7068	6822.00	26	262.385
VER	11669	12000	12035	11901.33	20	595.067
VER	9675	9610	9843	9709.33	14	693.524
VER	14049	14954	14574	14525.67	13	1117.36
IMI 0.7	4505	4410	4318	4411.00	15	294.067
IMI 0.7	5121	4986	5256	5121.00	27	189.667
IMI 0.7	8424	8863	8479	8588.67	34	252.608
IMI 0.7	6070	5386	6695	6050.33	18	336.13
IMI 0.7	4679	4574	4704	4652.33	20	232.617
IMI 7.6	7088	6047	4479	5871.33	33	177.919
IMI 7.6	6823	7579	5702	6701.33	35	191.467
IMI 7.6	8402	8049	6417	7622.67	25	304.907
IMI 7.6	5574	5329	4885	5262.67	20	263.133
IMI 7.6	5391	5478	5299	5389.33	20	269.467
CONFIDOR 0.7	8112	8068	7985	8055.00	23	350.217
CONFIDOR 0.7	5646	5464	5663	5591.00	17	328.882
CONFIDOR 0.7	6759	9084	9971	8604.67	24	358.528
CONFIDOR 0.7	9726	9879	9838	9814.33	32	306.698
CONFIDOR 0.7	5302	5172	5134	5202.67	37	140.613
CONFIDOR 7.6	10063	10201	10871	10378.33	30	345.944
CONFIDOR 7.6	5512	5184	5261	5319.00	17	312.882
CONFIDOR 7.6	5387	5088	5133	5202.67	20	260.133
CONFIDOR 7.6	6691	6351	6313	6451.67	35	184.333
CONFIDOR 7.6	4140	4349	4737	4408.67	14	314.905
6CNA 0.4	4810	4558	4975	4781.00	17	281.235
6CNA 0.4	6150	6382	6300	6277.33	18	348.741
6CNA 0.4	8319	7656	8049	8008.00	31	258.323
6CNA 0.4	6921	6951	7221	7031.00	17	413.588
6CNA 4.7	6104	5324	4649	5359.00	32	167.469
6CNA 4.7	7530 5004	7072	8045	7549.00	34	222.029
6CNA 4.7 6CNA 4.7	5004 4664	4707 5489	3894 6359	4535.00 5504.00	12 19	377.917 289.684
Cu 0.75 μg L <sup>-1</sup>	7398	7606	7601	7535.00	25	301.4
Cu 0.75 μg L <sup>-1</sup>	10354	9455	9452	9753.67	36	270.935
Cu 0.75 μg L <sup>-1</sup>	9202	9305	9769	9425.33	26	362.513
Cu 0.75 μg L <sup>-1</sup>	8443	7696	12535	9558.00	26	367.615
Cu 0.75 μg L <sup>-1</sup>	6007	5404	7343	6251.33	25	250.053

Table 34: Fluorescence (FU) values for MXR activity measurments in G. fossarum exposed to IMI/Confidor 200SL (0.7 and 7.6 mg  $L^{-1}$ ) or 6CNA (0.4 and 4.6 mg  $L^{-1}$ ) and copper (0.75, 3 and 10  $\mu$ g  $L^{-1}$ ) for 24 h and data normalised on total animal's weight (n=3), (continued).

#### **MXR**

11						
Chemical (mg)	FU	FU	FU	Mean FU	Weight (mg)	FU/mg
IMI 0.7 + Cu 0.75 μg L <sup>-1</sup>	10051	8614	10440	9701.667	15	646.778
IMI $0.7 + \text{Cu } 0.75  \mu\text{g } \text{L}^{-1}$	7857	8271	8206	8111.333	22	368.697
IMI $0.7 + \text{Cu } 0.75  \mu\text{g } \text{L}^{-1}$	6847	6757	6331	6645	20	332.25
IMI $0.7 + \text{Cu } 0.75  \mu\text{g } \text{L}^{-1}$	8406	7333	8252	7997	25	319.88
IMI $0.7 + \text{Cu } 0.75  \mu\text{g } \text{L}^{-1}$	3930	3979	3979	3962.667	12	330.222
IMI $0.7 + \text{Cu } 3  \mu\text{g } \text{L}^{-1}$	21065	19993	21218	20758.67	20	1037.93
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	9321	9275	9565	9387	16	586.688
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	8579	8316	8632	8509	20	425.45
IMI 0.7 + Cu 3 μg L <sup>-1</sup>	10373	10078	10170	10207	41	248.951
IMI $0.7 + \text{Cu } 3  \mu\text{g } \text{L}^{-1}$	8670	8919	8359	8649.333	24	360.389
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	9885	9791	11097	10257.67	37	277.234
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	10076	9314	10202	9864	29	340.138
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	8807	8501	9295	8867.667	30	295.589
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	7670	7291	6490	7150.333	20	357.517
IMI 0.7 + Cu 10 μg L <sup>-1</sup>	6528	6289	6511	6442.667	19	339.088
$6$ CNA $0.4 + $ Cu $0.75 \mu g L-1$	10413	10402	10590	10468.33	23	455.145
6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	11331	11737	12093	11720.33	20	586.017
$6$ CNA $0.4 + $ Cu $0.75 \mu g L-1$	13490	12953	13089	13177.33	28	470.619
$6$ CNA $0.4 + $ Cu $0.75 \mu g L-1$	9252	8759	5609	7873.333	20	393.667
$6$ CNA $0.4 + $ Cu $0.75 \mu g L-1$	8339	8645	7021	8001.667	20	400.083
$6$ CNA $0.4 + $ Cu $3 \mu g L^{-1}$	8458	8406	9406	8756.667	22	398.03
6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	8373	8723	8281	8459	18	469.944
$6$ CNA $0.4 + $ Cu $3 \mu g L^{-1}$	9574	10189	9842	9868.333	27	365.494
$6$ CNA $0.4 + $ Cu $3 \mu g L^{-1}$	11020	10484	10382	10628.67	28	379.595
6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	9554	9708	9150	9470.667	20	473.533
6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	8436	7834	7652	7974	33	241.636
6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	9205	9023	8953	9060.333	23	393.928
6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	11432	10897	10871	11066.67	13	851.282
6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	9223	9143	9369	9245	14	660.357
6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	6897	6970	7162	7009.667	16	438.104
Cu 3 μg L <sup>-1</sup>	5008	4555	4856	4806.333	18	267.019
Cu 3 µg L <sup>-1</sup>	5883	6261	6644	6262.667	22	284.667
Cu 3 µg L <sup>-1</sup>	6719	6524	6687	6643.333	18	369.074
Cu 3 μg L <sup>-1</sup>	15341	14956	15695	15330.67	30	511.022
Cu 3 µg L <sup>-1</sup>	8124	7732	4547	6801	23	295.696
Cu 10 μg L <sup>-1</sup>	7527	7397	8179	7701	28	275.036
Cu 10 μg L <sup>-1</sup>	10841	10541	11875	11085.67	36	307.935
Cu 10 μg L <sup>-1</sup>	5185	5168	5755	5369.333	15	357.956
Cu 10 μg L <sup>-1</sup>	5156	4608	4653	4805.667	13	369.667
Cu 10 μg L <sup>-1</sup>	13806	13428	16417	14550.33	14	1039.31

Table 34: Fluorescence (FU) values for MXR activity measurments in G. fossarum exposed to IMI/Confidor 200SL (0.7 and 7.6 mg  $L^{-1}$ ) or 6CNA (0.4 and 4.6 mg  $L^{-1}$ ) and copper (0.75, 3 and 10  $\mu$ g  $L^{-1}$ ) for 24 h and data normalised on total animal's weight (n=3), (continued).

MXR							
III Chemical (mg)	Moan Ell	Weight (mg)	EII/ma	Chamical (ma)	Mean FU	Waight (mg)	EII/ma
· <del>-</del> -	Mean FU	Weight (mg) 38	FU/mg	Chemical (mg) IMI 0.7 + Cu 0.75 µg L <sup>-1</sup>	6217.0	Weight (mg) 27	FU/mg 230.259
control	4875.0 6304.0	38 25	128.2894737 252.16	IMI 0.7 + Cu 0.75 µg L <sup>-1</sup>	16866.0	27	
control		22		IMI 0.7 + Cu 0.75 μg L  IMI 0.7 + Cu 0.75 μg L <sup>-1</sup>			624.667
control	5179.0	28	235.4090909	IMI 0.7 + Cu 0.75 μg L  IMI 0.7 + Cu 0.75 μg L <sup>-1</sup>	9881.0	31	318.742
control	4949.0		176.75	IMI 0.7 + Cu 0.75 µg L  IMI 0.7 + Cu 0.75 µg L <sup>-1</sup>	7666.0	30	255.533
control	5691.0	26	218.8846154	IMI 0.7 + Cu 3 μg L <sup>-1</sup>	8779.0	30	292.633
CYC	5681.0	32	177.53125	IMI 0.7 + Cu 3 μg L  IMI 0.7 + Cu 3 μg L <sup>-1</sup>	7144.0	23	310.609
CYC	4973.0	18	276.2777778	• •	7157.0	24	298.208
CYC	7842.0	28	280.0714286	IMI 0.7 + Cu 3 μg L <sup>-1</sup>	7260.0	34	213.529
CYC	6995.0	27	259.0740741	IMI 0.7 + Cu 3 μg L <sup>-1</sup> IMI 0.7 + Cu 3 μg L <sup>-1</sup>	9489.0	24	395.375
CYC	11071.0	22	503.2272727		8136.0	23	353.739
VER	5797.0	24	241.5416667	IMI 0.7 + Cu 10 μg L <sup>-1</sup>	6534.0	23	284.087
VER	5630.0	20	281.5	IMI 0.7 + Cu 10 μg L <sup>-1</sup>	8166.0	26	314.077
VER	6675.0	19	351.3157895	IMI 0.7 + Cu 10 μg L <sup>-1</sup>	8764.0	28	313
VER	6396.0	18	355.3333333	IMI 0.7 + Cu 10 μg L <sup>-1</sup>	28904.0	34	850.118
VER	7013.0	20	350.65	IMI 0.7 + Cu 10 μg L <sup>-1</sup>	20856.0	20	1042.8
IMI 0.7	6369.0	32	199.03125	6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	9329.0	24	388.708
IMI 0.7	4690.0	22	213.1818182	6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	11024.0	25	440.96
IMI 0.7	5864.0	24	244.3333333	6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	8096.0	15	539.733
IMI 0.7	5245.0	24	218.5416667	6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	7295.0	15	486.333
IMI 0.7	6156.0	23	267.6521739	6CNA 0.4 + Cu 0.75 μg L <sup>-1</sup>	7888.0	22	358.545
IMI 7.6	10862.0	27	402.2962963	6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	5441.0	17	320.059
IMI 7.6	7065.0	17	415.5882353	6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	9423.0	19	495.947
IMI 7.6	4847.0	22	220.3181818	6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	6262.0	11	569.273
IMI 7.6	7147.0	26	274.8846154	6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	8585.0	15	572.333
IMI 7.6	7047.0	24	293.625	6CNA 0.4 + Cu 3 μg L <sup>-1</sup>	8123.0	27	300.852
CONFIDOR 0.7	6446.0	24	268.5833333	6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	10058.0	34	295.824
CONFIDOR 0.7	8348.0	25	333.92	6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	15211.0	32	475.344
CONFIDOR 0.7	6560.0	29	226.2068966	6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	10811.0	30	360.367
CONFIDOR 0.7	6904.0	27	255.7037037	6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	13534.0	26	520.538
CONFIDOR 0.7	7216.0	25	288.64	6CNA 0.4 + Cu 10 μg L <sup>-1</sup>	8184.0	16	511.5
CONFIDOR 7.6	7117.0	31	229.5806452	Cu 3 μg L <sup>-1</sup>	8236.0	27	305.037
CONFIDOR 7.6	5592.0	24	233	Cu 3 μg L <sup>-1</sup>	10337.0	19	544.053
CONFIDOR 7.6	8788.0	27	325.4814815	Cu 3 μg L <sup>-1</sup>	8688.0	26	334.154
CONFIDOR 7.6	5457.0	28	194.8928571	Cu 3 µg L <sup>-1</sup>	5985.0	18	332.5
CONFIDOR 7.6	6925.0	30	230.8333333	Cu 3 μg L <sup>-1</sup>	8148.0	23	354.261
6CNA 0.4	5324.0	19	280.2105263	Cu 10 μg L <sup>-1</sup>	12155.0	27	450.185
6CNA 0.4	10703.0	10	1070.3	Cu 10 μg L <sup>-1</sup>	12386.0	27	458.741
6CNA 0.4	9669.0	20	483.45	Cu 10 μg L <sup>-1</sup>	9319.0	23	405.174
6CNA 0.4	5027.0	12	418.9166667	Cu 10 μg L <sup>-1</sup>	9104.0	26	350.154
6CNA 0.4	6201.0	11	563.7272727	Cu 10 μg L <sup>-1</sup>	6193.0	30	206.433
6CNA 4.7	9332.0	18	518.444444				
6CNA 4.7	6644.0	25	265.76				
6CNA 4.7 6CNA 4.7	8014.0 7262.0	21 24	381.6190476 302.5833333				
6CNA 4.7	6903.0	23	300.1304348				
Cu 0.75 μg L <sup>-1</sup>	12718.0	28	454.2142857				
Cu 0.75 μg L <sup>-1</sup>	8523.0	31	274.9354839				
Cu 0.75 μg L <sup>-1</sup>	12161.0	29	419.3448276				
Cu 0.75 μg L <sup>-1</sup>	7835.0	24	326.4583333				
Cu 0.75 μg L <sup>-1</sup>	7032.0	28	251.1428571				

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Giordano Bruno

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