



# Acute exposure to a commercial formulation of Azoxystrobin alters antioxidant enzymes and elicit damage in the aquatic macrophyte *Myriophyllum quitense*

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**Abstract** Azoxystrobin is a strobilurin of growing concern in aquatic environments because it is the most sold fungicide worldwide, however, the information available about its effect on aquatic non-target organisms is scarce. The objective of the present study was to evaluate potential physiological, biochemical, and genetic effects at environmentally relevant (1–10 µg/L) and elevated (100–500 µg/L) concentrations in the aquatic macrophyte *Myriophyllum quitense* exposed to the commercial formulation AMISTAR<sup>®</sup>. Following an acute 24-h exposure, there were no effects of AMISTAR<sup>®</sup> on photosynthetic pigments at any of the concentrations evaluated. Glutathione-S-transferase activity was significantly elevated at 1 and 10 µg/L AZX. Significant decrease of catalase and guaiacol peroxidase activities in plants exposed to 500 µg/L, and to 100 and 500 µg/L, respectively, and an increase in glycolate oxidase activity at 500 µg/L was observed. DNA damage at 100 and 500 µg/L was observed. These data indicate that although environmentally relevant levels of AMISTAR<sup>®</sup> did not result cytotoxic, this fungicide was genotoxic, affecting the physiological process of photorespiration and caused oxidative damage at high

concentrations. In this sense, it is necessary to explore sub-lethal responses in non-target organisms because some effects could promote further potential long-term biological consequences in a context of repeated pulses of exposure.

**Keywords** Azoxystrobin · *Myriophyllum quitense* · Antioxidant enzymes · “Comet” assay · Chlorophyll

## Introduction

Globally, agricultural producers apply around USD 40 billion worth of pesticides per annum (Popp et al. 2013). Consequently, their presence in the environment has grown in the past few years and has become an intensive and burning issue of discussion (Ghosh and Singh 2009). Particularly, the consumption of agrochemicals in Argentina increased from 73 to 236 million kg per year over the last 10 years (De Gerónimo et al. 2014). Fungicides are widely applied mostly in arable crops, which are more sensitive to fungal infections (Hocinat and Boudemagh 2016). The strobilurins are an important class of agricultural fungicides (Bartlett et al. 2001). Azoxystrobin (AZX) was the first strobilurin compound launched in the market in 1996 under the commercial name of AMISTAR<sup>®</sup> (Bartlett et al. 2002). The fungicidal activity is generated by binding Q<sub>0</sub> site of cytochrome b blocking the electron transfer (Huang et al. 2007). Nowadays AZX is marketed in 100 countries and applied for approximately 120 crops (SYNGENTA 2017). According to international authorities, it is the principal fungicide sold in the world (Royal Society of Chemistry 2017). The information available about AZX concentration in water bodies is scarce. Concentrations from 0.06 µg/L in streams and ponds from the US (Reilly

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et al. 2012) to 29.7 µg/L in streams from Germany (Berenzen et al. 2005) have been reported. In addition, Edwards et al. (2016) showed the horizontal movement with agricultural runoff as vector for transport of AZX, which increases the likelihood of exposure to non-target aquatic species. Therefore, the increase in the volume applied due to the resistance of fungi and its massive use worldwide suggest that the detections of AZX in waterbodies near agricultural fields will be more frequent in the future. In 2010, the European Food Safety Authority (EFSA) determined a value of 3.3 µg/L as the regulatory acceptable concentration (RAC) (Rodrigues et al. 2013). However, recent data suggest a Maximum Acceptable Concentration-Environmental Quality Standard (MAC-EQS) of 1.8 µg/L as a limit value protective in acute exposure (Rodrigues et al. 2017). It is well recognized that there is a gap in the ecotoxicological information concerning fungicides in the aquatic environment and these fact limits result interpretations (Battaglin et al. 2011). In this sense, the impact on non-target organisms can be toxic, but it is not fully known and can lead to ecosystem alterations (Deb et al. 2010). Fungicides, like most of the pesticides, are always used with emulsifiers, carrying solvents, and other additives. Accordingly, other authors highlighted about the higher potency of commercial formulations than their corresponding active ingredients for several pesticides (Pereira et al. 2009).

Effects of AZX have been detected in terrestrial plants. In long-term exposures, a delay in senescence in winter wheat (*Triticum aestivum*) have been observed (Wu and von Tiedemann 2001; Zhang et al. 2010). This fact is an advantage from the agronomic point of view because it enhances the grain yield of wheat. Although it could be considered a good characteristic to producers, non-target organisms that occur in natural water bodies could suffer negative effects. Indeed, perturbation of ascorbate synthesis, fat acid metabolism, and RNA translation have been recently demonstrated in the green algae *Chlorella pyrenoidosa* exposed to 0.5–10 mg/L AZX for 10 days, suggesting that AZX inhibits algal cell growth through multiple pathways (Lu et al. 2018). On the other hand, AZX also affects mitochondrial respiration and mechanisms controlling cell growth and proliferation in fish (Olsvik et al. 2010). Adult zebrafish (*Danio rerio*) showed reduced egg production and fertilization rates, as well as other reproductive effects during exposures at 200 µg/L AZX for 21 days (Cao et al. 2016). Toxic effects of AZX on brackish water copepods has also been described, and these effect could trigger a cascade effect affecting ecosystem functioning e.g. indirect effects on phytoplankton community composition (Gustafsson et al. 2010).

Aquatic macrophytes play a key role in aquatic ecosystems as primary producers, shelter, oxygen

contributor, sediment stabilizer, and potential detoxifier. In order to maintain ecosystem structure and functions, the aquatic macrophytes, as other aquatic organisms, must be protected from adverse chemical effects (Arts et al. 2010). Macrophytes could be employed as biomonitor organisms, because they are easy to see and identify in the field, sedentary and, depending on the pollutant characteristics, they concentrate it in their tissues reflecting the environmental contamination (Nimptsch et al. 2005). In 2008, a panel of experts from the Society of Environmental Toxicology and Chemistry (SETAC)—The Aquatic Macrophyte Ecotoxicology Group (AMEG)—published guidance and recommendations to evaluate different substances in aquatic macrophyte risk assessment. They proposed *Myriophyllum* as a recommended genus because of the considerable knowledge and experience that researchers have about it. Indeed, the Organization for Economic Cooperation and Development (OECD) adopted *M. spicatum* in the guideline TG 239 (2014) as new test system for rooted aquatic plants (OECD 2014). This genus has been used to evaluate the toxicity of different substances such as chlorodifluoroacetic acid (Popp et al. 2013), Cd, Cu (Ngayila et al. 2009), Ni, Pb, Zn (Nimptsch et al. 2005), ammonia (Nimptsch and Pflugmacher 2007) and the insecticide endosulfan (Menone et al. 2008).

The xenobiotic stress in plants is manifested firstly at the biochemical level, before its manifestation at organism level, therefore this biochemical effects can work as early warning systems (Brain and Cedergreen 2009). For this reason, these type of biomarkers were chosen in the present study to evaluate the potential sub-lethal effects of AMISTAR<sup>®</sup> in acute bioassays. Particularly, in ecotoxicological studies, the oxidative stress caused by oxiradicals and the potential cellular oxidative damage is widely applied to determine effects of several xenobiotics (Valavanidis et al. 2006). Exposure to some xenobiotics may result in decrease of antioxidant defenses, leading to tissue damage, and different processes that ultimately cause cellular aging and diseases (Sohal et al. 2002), moreover the nuclear DNA is sensitive to oxidative damage by ROS. Genetic alterations can have a number of immediate effects upon the cells involved, including cell death or transformation into malignancy (Bickham et al. 2000). The most advantage method to measure DNA strand breaks is the “Comet assay”. The DNA damage is measured quantitatively by single-cell gel electrophoresis technique which is widely applied because of its rapidity and sensitivity in aquatic organisms, although scarcely applied in plant tissues (Valavanidis et al. 2006).

Considering that there is not enough available data on the effects of AZX commercial formulations on non-target aquatic organisms, its increasing application worldwide and the sensitivity of *Myriophyllum*, the objective of the

present study was to evaluate potential physiological, biochemical and genetic effects at environmentally relevant (1–10 µg/L) and elevated (100–500 µg/L) concentrations in the aquatic macrophyte *M. quitense* exposed to the commercial formulation AMISTAR®.

## Materials and methods

### Plant material and exposure conditions

*Myriophyllum quitense* plants were obtained from a natural reservoir, placed in the Estación Experimental Agropecuaria Balcarce (INTA), Argentina. Species identification was done according to Orchard (1981). Plants were acclimated for 15 days in 30 L tanks, the light was provided by fluorescence lamps with an irradiance of 8000 lux at a light/dark cycle of 12:12 h. Room temperature was maintained at 20–22 °C, and the plants were submerged in Hoagland's medium (pH 5) (Arts et al. 2010).

Two experiments were carried out to evaluate oxidative biomarkers and genotoxicity, respectively. Four concentrations of AZX were evaluated, two environmental concentrations, and two concentrations that exceed the actual environmental values, but would be expected if the consumption of AZX continues increasing. Plants of *M. quitense* (six independent replicates per treatment, n = 6) were exposed to AMISTAR® Syngenta® equivalent to 0 (Co-), 1, 10, 100 and 500 µg/L of AZX, in a volume of 350 mL each, for 24 h, under constant light and temperature conditions. AMISTAR® was dissolved in bidestillated water in a stock solution of 2000 mg/L. Each exposure concentration was prepared by diluting the appropriate amount of AMISTAR® in Hoagland medium to a final volume of 350 mL. In the negative control treatment (Co-), AMISTAR® was omitted from the medium. One positive mutagenic control (Co +) consisting of 7.14 g/L of hydrogen peroxide was also included. After exposures, plants were frozen with liquid N<sub>2</sub> before being stored at – 80 °C until processing for analyzing enzymes activities, H<sub>2</sub>O<sub>2</sub> and MDA contents, or directly submerged in extraction buffer for the “Comet” assay.

### Azoxystrobin quantification

Water samples (250 ml) were extracted with 20 ml chloroform. 50 µl of 5 ppm 1-bromo-2-nitrobenzene in toluene was added as internal standard. The method was based on extraction with chloroform, followed by a clean-up with QuEChErs (AOAC Methods 2003, 2007). Sample preparation, based on QuEChErs method, combined a liquid–liquid extraction with acetonitrile and a dispersive-SPE clean-up. The organic phase was transferred into a beaker

(250 mL) and evaporated under nitrogen stream at 40 °C ± 2 °C in a water bath to a volume of about 0.3–0.5 ml. The residue was taken up with toluene to a volume of about 1 ml and transferred to a chromatography vial refrigerated until use.

Chromatographic conditions: An Agilent 7890 chromatograph was used for the chromatographic run, associated with a 5975 Series Mass Selective Detector (MSD). Oven temperatures were 70 °C (2 min), 25 °C up to 150 °C (0 min), 3 °C/min up to 200 (0 min), and 8 °C/min up to 280 °C (10 min). Injection was in pulsed splitless mode with an automatic injector. Helium was used as carrier gas. The mass detector was used in SIM Mode, Source, Quadrupole, Temperature of Transfer Line mode: 230, 150, and 280, respectively, Solvent delay: 5.00 min, Multiplier voltage: Gain = 25. After the run was completed, AZX was quantified with the chromatograph (Agilent G1701 EA GC/MSD ChemStation) software. An analyte recovery percentage between 82 and 110% was obtained with a coefficient of variation ≤ 14%. The detection limits were generally 10 ng/L. To evaluate the method performance, calibration curves at concentrations of 10, 20, 50, 100, 200 ng/L were constructed. Injections of the standards were repeated at concentrations of 50 ng/L in matrix to assure system stability. The limits of detection and quantification were 1 µg/L (signal-to-noise ratio > 3).

### Battery of biomarkers

#### Chlorophyll content

Chlorophyll content was measured according to Inskeep and Bloom (1985). Leaf samples of 0.2 g each were put into vials containing 2 ml of N, N-dimethylformamide (DMF). The vials were kept closed for 72 h, at 4 °C in darkness until the complete loss of green color. The absorbance values of the supernatant were recorded at 647 and 664 nm using a spectrophotometer. For calculation of quantitative values of pigments, the following equations were applied:

$$\begin{aligned} \text{Chl } a &= (12.70A_{664.5} - 2.79A_{647})/\text{g Fw}^{-1} \\ \text{Chl } b &= (20.70A_{647} - 4.62A_{664.5})/\text{g Fw}^{-1} \\ \text{Total Chl} &= (17.90A_{647} + 8.08A_{664.5})/\text{g Fw}^{-1} \end{aligned}$$

### Oxidative stress biomarkers

Preparation of guaiacol peroxidase (POD), glutathione-S-transferase (GST) and catalase (CAT) enzymes was performed according to Pflugmacher (2004). Approximately 10 g fresh weight of plant leaves were homogenized with sodium-phosphate buffer 0.1 M (pH 6.5), containing glycerol 20%, dithioerythritol (DTE) 14 mM and

ethylenediaminetetraacetic acid (EDTA) 1 mM. The homogenate was centrifuged at 10,000 g for 10 min. The enzyme glycolate oxidase (GOX) was extracted with a buffer containing sodium-phosphate buffer 100 mM (pH 6.8), with EDTA 1 mM, BSA 0.1% and Triton 0.1%. Protein determination was done according to Bradford (1976) using bovine serum albumin (BSA) as the standard. Enzyme activity was calculated in nanokatals per milligram of protein from percent substrate conversion. Measurement of the POD activity using guajacol as the substrate was performed as described by Drotar et al. (1985). Determination of GST activity with the model substrate 1-chloro-2, 4-dinitrobenzene (CDNB) was done according to Habig et al. (1974). CAT activity was assayed according to Claiborne (1985). GOX activity was measured following the method of Archer and Ting (1996). All measurements were made in duplicate.

#### *H<sub>2</sub>O<sub>2</sub> content*

The level of cell internal H<sub>2</sub>O<sub>2</sub> was determined calorimetrically by monitoring the formation of titanium peroxide according to Jana and Choudhuri (1982). H<sub>2</sub>O<sub>2</sub> was extracted by homogenizing 0.4 g of leaf tissue with 3 ml of 50 mM sodium phosphate buffer (pH 7.0). The homogenate was spun down at 10,000 g for 2 min at 4 °C and the transparent supernatant was collected. 750 µl of supernatant was mixed with 250 µl of 0.1% titanium sulphate (TiSO<sub>4</sub>) in 20% (v/v) sulfuric acid (H<sub>2</sub>SO<sub>4</sub>). Then, the intensity of red–orange colour of the mixture was measured at 410 nm after 1 min. All measurements were made in triplicate with each extract from 6 independent replicates. The content of H<sub>2</sub>O<sub>2</sub> was calculated using the extinction coefficient 0.28 l/mmol cm and expressed as µmol g/FW.

#### **Lipid peroxidation**

Malondialdehyde (MDA), the principal product of polyunsaturated fatty acid peroxidation, was evaluated according to Shi et al. (2006). A leaf sample of 0.3 g was homogenized in 3 mL of trichloroacetic acid (TCA) 0.1% (w/v), centrifuged at 10,000 g for 10 min, and 3 mL of TCA 20% containing thiobarbituric acid (TBA) 0.5% (w/v) was added to 1 mL of supernatant. The mixture was heated at 95° C for 30 min, placed into an ice-bath and centrifuged at 10,000g for 10 min, and the absorbance of the supernatant was read at 532 and 600 nm. After subtracting the non-specific absorbance at 600 nm, the TBARS concentration was calculated by its extinction coefficient of 155 mM/cm.

#### **DNA fragmentation**

The isolation of nucleoids and the alkaline “Comet” assay was made according to Garanzini and Menone (2015). Briefly, the isolation was made through a sucrose cushion and the electrophoresis was run in denaturing solution at 0.72 V/cm for 30 min. Gels were silver stained according to Nadin et al. (2001) for optic microscope observation using a microscope OLYMPUS CX31. The quantification of the level of DNA damage was made by measuring the relative length of the tail of the “Comet” in 200 nuclei per sample. Where each nucleoid was classified into five classes according to tail size (from undamaged, class 0, to maximally damaged, class 4), resulting in a single DNA damage score (Damage Index, DI) for each slide. The DI was calculated as follows:  $DI = n_1 + 2n_2 + 3n_3 + 4n_4$ ; where  $n_1$ ,  $n_2$ ,  $n_3$  and  $n_4$  are the number of cells in class 1, 2, 3 and 4 of damage respectively according to Poletta et al. (2009).

#### **Statistical analysis**

Normality and homogeneity of variances were verified by Shapiro–Wilk and Levene tests, respectively. Photosynthetic pigment data showed normality, therefore a one-way ANOVA test was applied followed by the post hoc Dunnett’s test. For the other parameters, the median was used as the measure of central tendency, and non-parametric tests were applied, because data did not fit with a normal distribution and variance homogeneity. Kruskal–Wallis and, a posteriori, Dunn test for testing differences among treatments were applied (Zar 1999). In all cases, statistical analyses were determined at a 0.05% significance level. A canonical discriminant analysis of principal coordinates (CAP) using 9999 permutations (Anderson and Robinson 2003, Anderson and Willis 2003) were performed in order to determine to what extent the set of 11 variables can actually explain treatment differences. For this, Euclidean distances was applied (Anderson and Willis 2003).

## **Results**

#### **Azoxystrobin quantification**

Analytical measurements of AZX were carried out to corroborate the expected concentration according to the nominal concentrations in AMISTAR®. Measured concentrations of 1.1, 9.8, 99.7 and 503.2 µg/L AZX corresponding to nominal concentrations of 1, 10, 100 and 500 µg/L AZX were obtained.

## Battery of biomarkers

### Chlorophyll content

None of the concentrations evaluated exerted effects in total chlorophyll, chlorophyll a, chlorophyll b and chlorophyll a/b ratio as none differed significantly from the control ( $p > 0.05$ ) (Table 1).

### Biochemical biomarkers

The activity of GST increased in plants exposed at 1 and 10  $\mu\text{g/L}$  ( $p < 0.05$ ) (Fig. 1 D). The antioxidant enzymes CAT and POD showed a decrease in comparison to controls at 500  $\mu\text{g/L}$ , and at 100 and 500  $\mu\text{g/L}$ , respectively ( $p < 0.05$ ) (Fig. 1a, b). The activity of GOX was increased compared to controls at 500  $\mu\text{g/L}$  ( $p < 0.05$ ) (Fig. 1c). Levels of  $\text{H}_2\text{O}_2$  as well as lipid peroxidation measured as TBARS were not different in plants exposed to AMISTAR<sup>®</sup> with respect to control plants ( $p > 0.05$ ) (Fig. 1e, f).

### DNA fragmentation

Figure 2 shows a significant increment of DNA fragmentation in plants exposed to the positive control with respect to the negative control. Increased DNA damage was observed at 100 and 500  $\mu\text{g/L}$  compared to the negative controls ( $p < 0.05$ ).

### Canonical discriminant analysis of principal coordinates (CAP)

The CAP analysis showed the same patterns of response observed in the individual statistical analysis. The plot produced by visualizing the values obtained on the new canonical axes (Fig. 3) graphically separated the different treatments, mainly controls from plants exposed to AMISTAR<sup>®</sup>. The cross validation of allocation of observations to groups showed a value of 80% ( $p < 0.0001$ ).

## Discussion

### Azoxystrobin quantification

The AZX measured concentrations were highly concordant with the nominal concentrations. These results are in agreement with Singh et al. (2010) as they found that this fungicide is hydrolytically fairly stable between pH 4 and 9.

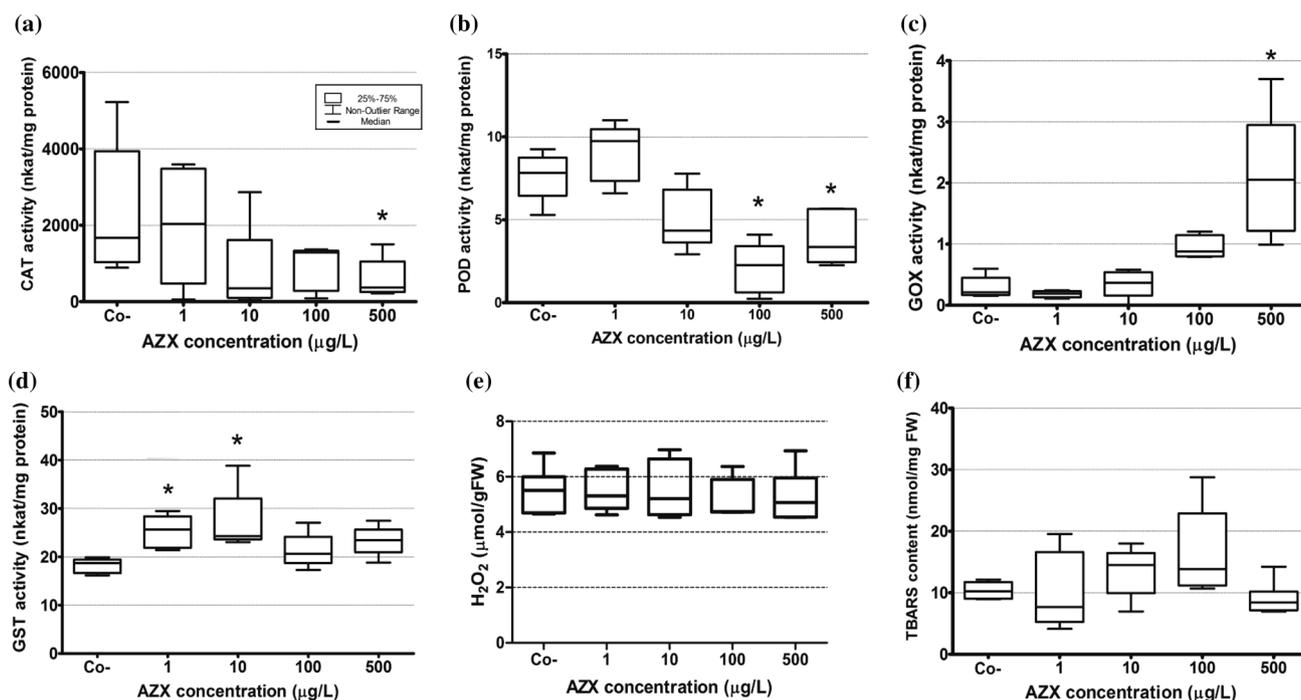
### Battery of biomarkers

The results indicated that the fungicide AMISTAR<sup>®</sup> did not disturb the photosynthetic pigments at any of the concentrations evaluated, from environmentally low to high. Kozłowski et al. (2009) also found no differences in chlorophyll content in the terrestrial plant *Phaseolus vulgaris* exposed to AMISTAR<sup>®</sup> during 21 days. Indeed, previous studies, for other fungicides like tebuconazole have shown the same results, in the green algae *Pseudokirchneriella subcapitata* and *Nannochloris oculata* (Martinez et al. 2015). In all cases, this lack of response was interpreted as a non-cytotoxic effect. On the other hand, a decrease in the net photosynthetic rate, measured through the gas exchange and intercellular carbon dioxide concentration, in *Triticum aestivum* L., *Hordeum vulgare* L. and *Glycine max* exposed to the individual active ingredient AZX for at least 24 h have been detected (Nason et al. 2007). This disagreement with our data could be due to excipients of the commercial formulation or to differences among species.

GST activity is sensitive to acute exposures to xenobiotics (< 24 hs), and this response allows its application as an early warning biomarker (Brain and Cedergreen 2009). This enzyme system could be important in a protective role catalyzing the conjugation and detoxification of many types of pesticides. For example, conjugation of herbicides with reduced glutathione (7-glutamyl cysteinyl glycine, GSH) is an important and irreversible mode of detoxification (Cole 1994). However, in addition to its detoxifying function in plants and animals (conjugation with

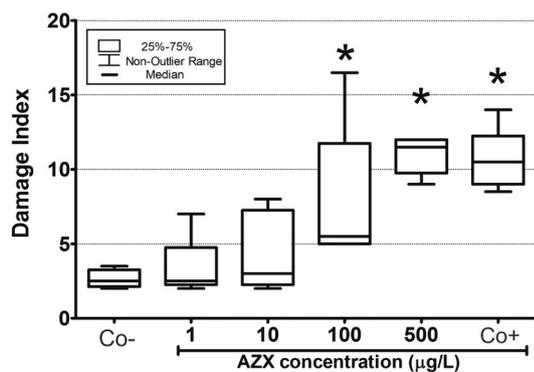
**Table 1** Photosynthetic parameters (mean  $\pm$  SD) in the aquatic macrophyte *Myriophyllum quitense* exposed to AMISTAR<sup>®</sup>

	AZX Concentrations ( $\mu\text{g/L}$ )				
	Co-	1	10	100	500
Chl a ( $\text{mg/g Fw}^{-1}$ )	430.1 $\pm$ 376.0	468.7 $\pm$ 118.3	696.1 $\pm$ 103.2	595.0 $\pm$ 122.5	526.1 $\pm$ 130.6
Chl b ( $\text{mg/g Fw}^{-1}$ )	2226 $\pm$ 873.2	1812 $\pm$ 687.8	1575 $\pm$ 231.2	1499 $\pm$ 134.3	1443 $\pm$ 604.2
Total Chl ( $\text{mg/g Fw}^{-1}$ )	2691 $\pm$ 637.8	2291 $\pm$ 608.6	1978 $\pm$ 638.2	2071 $\pm$ 133.6	1979 $\pm$ 519.6
Ratio Chl a/b	0.16 $\pm$ 0.13	0.31 $\pm$ 0.19	0.45 $\pm$ 0.02	0.36 $\pm$ 0.01	0.46 $\pm$ 0.31



**Fig. 1** Oxidative stress parameters in *Myriophyllum quitense* exposed to AMISTAR® ( $\mu\text{g/L}$  AZX). Co-: negative control. **a** Catalase (CAT) activity; **b** guaiacol peroxidase activity (POD) activity; **c** glycolate oxidase (GOX) activity; **d** glutathione-S-transferase (GST) activity; **e**  $\text{H}_2\text{O}_2$  content ( $\mu\text{mol/gFW}$ ) and **f** lipid

peroxidation as TBARS content ( $\text{nmol/mgFW}$ ). \*Significantly different from the control (Kruskal–Wallis with post hoc Dunn’s test) ( $p < 0.05$ ). In all cases, box limits show 25–75 percentiles and whiskers represent the non-outlier range



**Fig. 2** DNA fragmentation measured through the alkaline “Comet” assay in the aquatic macrophyte *Myriophyllum quitense* exposed to AMISTAR®. Co-: negative control, CO+: positive control ( $7.14 \text{ g/L}$   $\text{H}_2\text{O}_2$ ). \*Significantly different from the control (Kruskal–Wallis with post hoc Dunn’s test) ( $p < 0.05$ ). In all cases, box limits show 25–75 percentiles and whiskers represent the non-outlier range

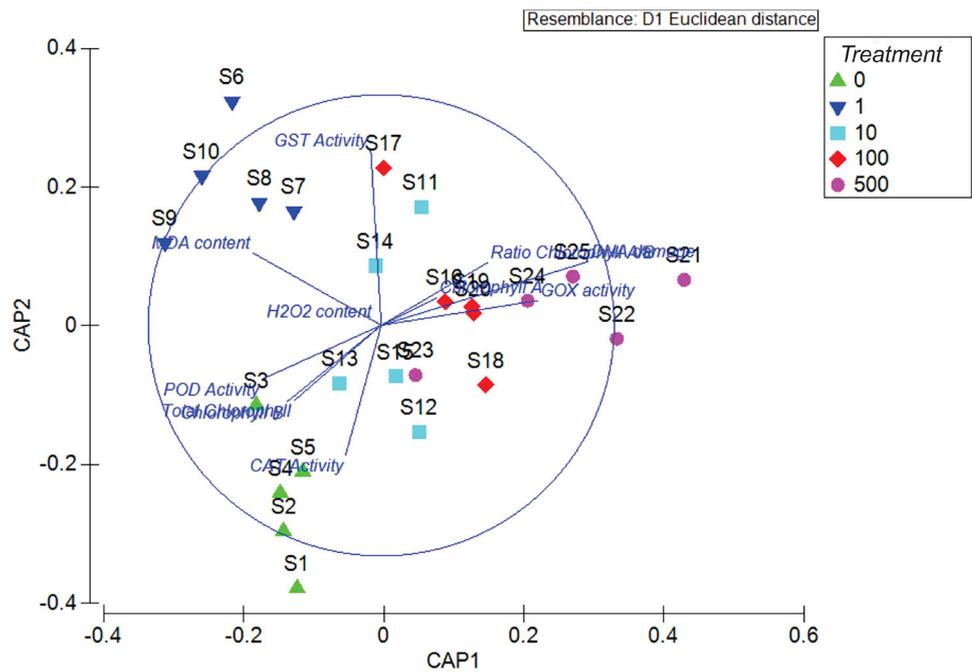
glutathion), GSTs conjugate GSH with a variety of electrophilic substrates to produce less toxic forms, which results in their detoxification, and reduce oxidative stress (Marrs 1996; Liu et al. 2015). The latter may be the process that takes place in *M. quitense* exposed to AMISTAR® where the GST activity increased in plants exposed to environmentally relevant concentrations. Taking into account that the most abundant metabolite of AZX in

plants is the product of the ether bond cleavage, cyanophenoxy-pyrimidinol, which is readily converted as N-glucoside conjugate (Balba 2007); the response in GST observed in *M. quitense* could be related to its role as an antioxidant enzyme by conjugating membrane intermediate products of lipid peroxidation in a direct manner (Marrs 1996; Cummins et al. 1999) instead to biotransformation.

The decrease in CAT activity observed could indicate oxidative damage, as proposed by Kono and Fridovich (1982) who observed that CAT activity may decrease when there is an excess of radical superoxide anion. Our results were obtained immediately after an exposure of 24 h, simulating a runoff event; but in long-term exposures, increase of these enzymes in plants of *Triticum aestivum* after 37–80 days (Wu and von Tiedemann 2001; Zhang et al. 2010); and in the algae *Chlorella pyrenoidosa* after 10 days (Lu et al. 2018) was observed. A peak of  $\text{H}_2\text{O}_2$  would be expected when peroxidases activities were decreased, however no changes in this biomarker were found. Our results are in agreement with data reported by Liang et al. (2016) in ginseng stem (*Panax ginseng*) exposed to AZX.

The activity of GOX generates  $\text{H}_2\text{O}_2$  as a result of the transformation of glycolate to glyoxylate in the process of photorespiration (McCarthy et al. 2001). AMISTAR® would deregulate the photorespiration process evidenced

**Fig. 3** Canonical discriminant analysis of principal coordinates (CAP) indicating effects of AMISTAR® in the aquatic macrophyte *Myriophyllum quitense*. Treatments: 0 = negative control, 1 = 1 µg/L AZX, 10 = 10 µg/L AZX, 100 = 100 µg/L AZX and 500 = 500 µg/L AZX



by the increased activity observed in this enzyme, that may generate  $H_2O_2$  which could contribute to its total content (Jana and Choudhuri 1982). However, no changes in  $H_2O_2$  observed would indicate that it might be confined to glioxisomes where it was quickly detoxified.

### DNA fragmentation

It has been demonstrated that ROS induce oxidation of DNA, breakage of DNA chains, and other deleterious changes of the nucleotide bases (Cooke et al. 2003). Han et al. (2014) observed that an excess of ROS was likely a cause of DNA damage observed in the earthworm *Eisenia fetida*, especially when the animals were exposed to 0.1 mg AZX/Kg soil for 7 days. The same authors observed increased DNA damage levels in zebrafish (*Danio rerio*) livers, reporting DNA strand breaks under exposure to AZX in a concentration and exposure time dependent manner (Han et al. 2016).

### Integration of biomarkers

The significant inhibition of CAT and POD, as well as the increased DNA damage observed at high concentrations of AMISTAR® coincided with the breakdown of the GST enzyme at 100 and 500 µg/L, indicating an acute stress condition. Exposure to AMISTAR® may produce oxidative damage in proteins and DNA. The results obtained only showed GST as an early warning signal, since it was the only biomarker that responded at low concentrations of AMISTAR®.

The canonical discriminant analysis of principal coordinates allowed to correlate all the effects observed in the different biomarkers evaluated. This analysis showed the integrated responses in plants exposed to AMISTAR®, where plants unexposed to AMISTAR® were significantly different to the exposed plants in the two principal axes evaluated. In addition, the different levels of exposition generated different responses in the biomarkers, while plants exposed to low concentrations showed mainly differences in GST activity, plants exposed to high concentrations behaved similarly between them, particularly in DNA damage and GOX activity. Overall, the discriminant analysis allowed the integration of all biomarkers used, and helped us to identify the oxidative stress effect exerted by AMISTAR® on *M. quitense*, more difficult to understand when individual biomarkers were analyzed.

In the aquatic environment, AZX short-term pulses of few days would be expected in surface waters, because fungicide occurrence is related to its use in the associated drainage watersheds (Battaglin et al. 2011). AZX has a water solubility of 6 mg/L and a log Kow = 2.5 (Bartlett et al. 2002), therefore, it can remain partitioned to water in the environment. However, in this matrix, a short half-life of only 14 days in natural conditions due to the photolytic degradation has been described (Boudina et al. 2007). As it was mentioned before, there are few reports of AZX toxicological data in non-target organisms, even less in aquatic ones acutely exposed. In this sense, Liu et al. (2013) determined a  $LC_{50}$  value of 549 µg/L in early life stage of grass carp (*Ctenopharyngodon idella*) and also an increase in CAT and POD activities, and inhibited

expression of several genes related to growth, energy pathways and up-regulation of stress. In a study with tadpoles of *Rana temporaria*, the exposure to AZX resulted as the more toxic among six pesticides tested, exerting negative effect in the growth (Johansson et al. 2006). Recent studies of chronic toxicity of AZX on the amphipod *Hyalella azteca*, the midge *Chironomus dilutus*, the cladoceran *Ceriodaphnia dubia* and the unionid mussel *Lampsilis siliquoidea* exposed to environmentally relevant concentrations (4.2–12 µg/L) showed negative effects in reproduction and survival (Kunz et al. 2017).

Although environmentally relevant levels of AMISTAR® did not result cytotoxic in *M. quitense*, this fungicide was genotoxic, affected the physiological process of photorespiration and caused changes in antioxidant enzymes mostly at high concentrations. In this sense, it is necessary to explore sub-lethal responses in aquatic non-target organisms because some effects could promote further potential long-term biological consequences in a context of repeated pulses of exposure.

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