



Effects on Ultrastructure, Composition and Specific Surface Area of the Gills of *Odontesthes bonariensis* Under Subchronic Glyphosate-Based Herbicide Exposure

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Abstract

Gills represent one of the major sites of gas exchange of fish, consequently they are in continuous close contact with the aquatic environment and its pollutants. In the present study the effects on gills of pejerrey fish, *Odontesthes bonariensis*, under glyphosate-based herbicide subchronic exposure were analyzed. Adult animals were exposed to sublethal concentrations of a glyphosate-based commercial formulation (1 and 10 PMG mg L⁻¹, PMG: glyphosate active ingredient) for 15 days, while control group was maintained in rearing water. Ultrastructural changes in gills were observed by scanning electron microscopy (SEM). The composition of the surface epithelium and specific surface area were determined by energy dispersive spectroscopy (EDS) and N₂ (g) adsorption–desorption isotherms, respectively. The herbicide exposure induced severe alterations in gill ultrastructure, as shown in the SEM micrographs. Accordingly, an increase in surface area of the gills of exposed animals was determined. These results support that gills parameters of freshwater fish are sensitive morphological biomarkers for glyphosate exposure.

Keywords Glyphosate-based herbicide · Gill ultrastructure · SEM · Specific surface area · Pejerrey fish *Odontesthes bonariensis*

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Water pollution by agrochemicals is one of the most serious problems for the conservation of aquatic ecosystems. In Argentina, transgenic soybean cultivation has continued growing over the last decade. The cropped area increased from approximately 9 million ha at the end of the 1990 decade up to 18 million ha in 2016/17 (MAGyP 2018). Consequently, the pesticide application rates (herbicides plus insecticides, fungicides, acaricides, among others) increased from 38,000 tons of active ingredients in 1995 (before the introduction of a GM glyphosate-tolerant crops) to more than 190,000 tons in 2017 (Food and Agriculture Organization, FAO 2018). Glyphosate [*N*-(phosphonomethyl) glycine; PMG] is the main broad-spectrum herbicide used for the control of a wide range of weeds in soybean crops. The application of glyphosate formulations commonly used in Argentine agriculture increased from 70,000 tons to more than 180,000 tons between 1999 and 2014 (CASAFE 2015). Once applied to the crop, part of the herbicide may remain adsorbed to soil particles until the degradation by microorganisms. Also, the herbicide can be mobilized by the influence of several factors such as wind, rain or irrigation that increase infiltration and surface run-off (Pérez et al. 2011;

World Health Organization, WHO 1994). Thus, the organic compound can reach nearby aquatic environments and biota can be adversely affected. In this regard, some authors have reported high levels of PMG and its metabolite aminomethylphosphonic acid (AMPA) in sediment, soil and water, particularly in the surroundings of cultivation areas (Peruzzo et al. 2008; Ronco 2011). Moreover, other authors reported lower PMG levels but a high frequency of detection (Primo et al. 2017; Mac Loughlin et al. 2020), suggesting that PMG could be considered as “pseudo-persistent” pollutant because it is continuously introduced to the environment under real-world agricultural management practices.

PMG does not exhibit a significant acute lethal toxicity to freshwater fish ($LC_{50-96\text{ h}} > 100\text{ mg L}^{-1}$) (Carriquiriborde 2010; Folmar et al. 1979; Menéndez-Helman et al. 2013; Nešković et al. 1996). However, several authors have noted that formulations containing the herbicide active ingredient together with co-adjuvants (glyphosate-based herbicides, GBH) exhibit a significant increase in toxicity to diverse aquatic organisms (algae, bacteria, protozoans, invertebrates and fish) (Menéndez-Helman et al. 2013; Pérez et al. 2011). Moreover, several sublethal effects of the PMG and GBH have also been reported in fish including histopathological disorders, metabolic changes associated with oxidative stress, energy disturbances, neurotoxicity, genotoxicity, among others (Menéndez-Helman et al. 2012, 2013, 2015, and references cited therein).

The pejerrey (*Odontesthes bonariensis*; Cuvier and Valenciennes, 1835) belongs to the Atherinopsidae family and is considered native to shallow lakes of Buenos Aires Province, Argentina, and probably in southern Brazil (Dyer 2006). This teleost presents socioeconomic relevance for commercial and sport fishing (Berasain et al. 2015), and additionally it is a promising species for aquaculture (Somoza et al. 2008). In several lethal toxicity studies using pejerrey as test organism, relatively low $LC_{50-96\text{ h}}$ values for different heavy metals and organic compounds have been estimated (Carriquiriborde and Ronco 2006; Carriquiriborde 2010). This fact confirms the sensitivity of *O. bonariensis* to environmental pollutants.

The fish gill is a very complex and multifunctional organ, involved in vital functions, such as gas exchange, ion exchange, acid-base balance, and nitrogenous waste excretion. This organ is in contact with the surrounding water, and its large surface area makes it an important target for waterborne pollutants (Cavalcante et al. 2008; Ferrari et al. 2009; Jiraungkoorskul et al. 2003). Thus, the ultrastructural responses are sensitive enough to indicate early effects on the organism exposed to low concentrations of chemicals, and they become potentially useful tools as morphological biomarkers to assess environmental pollution.

In this context, the aim of this study was to evaluate the exposure of pejerrey to a glyphosate-based herbicide,

specifically the effects on the ultrastructure, the composition of the surface epithelium and specific surface area of gills.

Materials and Methods

The bioassay was conducted at the aquaculture facilities of the Instituto Tecnológico de Chascomús (INTECH, Chascomús, Argentina) during the spring season. Three tanks, containing 120 L of groundwater (salinity 15 g L^{-1}), continuously aerated, were located outdoors. Adult fish (2 years old) were selected from the stock reared at the INTECH and four males and two females were placed in each tank (to prevent the disruption of the normal reproductive cycle). After acclimatization for 48 h, fish were transferred to another container while the commercial formulation of glyphosate was added to the tanks, after which fish were placed back into the tanks. Glifosato II® (manufactured by Atanor, Argentina) was the tested GBH formulation that contained 43.8% w/v mono potassium salt of PMG simulating a single contamination event. The doses used were 0 (control), 1 or $10\text{ mg PMG}_{\text{GBH}}\text{ L}^{-1}$ (PMG_{GBH} : PMG expressed as glyphosate acid contained in GBH solutions). During the assays, fish were fed commercial fish food pellets (Shulet, Buenos Aires, Argentina). The static trials were carried out for 15 days. Temperature and dissolved oxygen (DO) were measured in each tank every 2 days at 2 p.m. (the mean values \pm standard error were $19 \pm 1^\circ\text{C}$ and $6.2 \pm 0.3\text{ mg L}^{-1}$, respectively) and samples of the assay media were taken at the initial and final time to quantify the analytical concentration of the herbicide. At the end of the exposure period, fish were anesthetized with benzocaine (100 mg L^{-1}), weighed (W), and the total length (L_T) was measured. These parameters were used to calculate the Fulton's condition factor index ($K = [(W \times 100) \times (L_T)^{-3}]$). After this, the dissections were performed and gills were removed intact. Gills were washed with phosphate buffer (0.1 M, pH 7.2), fixed for 3 h in 2.5% glutaraldehyde solution in phosphate buffer and 4 additional washes with phosphate buffer were carried out during the next 48 h. Later, dehydration was carried out rinsing during 30 min with a graded series of 10%, 30% and 50% ethanol. Finally, samples were stored in 70% ethanol until analysis.

Gills were transferred from 70% ethanol to absolute ethanol, and samples were dried by Critical Point technique (Anderson 1951). The method is based on the displacement of ethanol by liquid CO_2 under high pressure up to critical pressure, and then heating until a temperature higher than critical temperature. This process allows CO_2 to pass from liquid to gaseous phase, at pressure and temperature conditions above the critical point of CO_2 , without forming a liquid/gas interface. Therefore, tissue samples were dried without collapsing or deforming their original structure. Then samples were fixed

on stubs by carbon conductive tape and coated with gold/palladium using a sputter coater (Thermo VG Scientific Sc7620). The surface morphology of the samples, an advanced form of high spatial resolution imaging, was visualized by a scanning electron microscope (SEM; Philips XL30 TMP New Look) in high vacuum conditions.

The elemental composition of the surface after metal binding was determined by energy dispersive spectroscopy (EDS, Oxford Instruments detector with INCA software) coupled to scanning electron microscope with field emission gun (FEG-SEM Zeiss Supra 40). In this case samples were coated with gold. One holobranch of each specimen ($n=2$ specimens per treatment) was bombarded by the SEM's electron beam in three different surface places for EDS analysis.

Specific surface area (SSA) is a property defined as the total surface area of a material per mass unit expressed in m^2g^{-1} . SSA was determined using a Micrometrics AccuSorb 2100 by N_2 (g) adsorption–desorption isotherms recorded at 77 K of temperature. Gill samples were degassed for 12 h at 150°C prior to measurement. The SSA was calculated using Langmuir model. Also, the pore volume (micropore and mesopore volume) define as the total volume of very small openings in a bed of adsorbent particles was calculated using the Gurvich rule from the plateau-like region of the nitrogen adsorption isotherm at a relative pressure of $p/p^0=0.99$, which is in the plateau-region of the isotherm (Lowell et al. 2004). For SSA analysis, holobranch from 5 to 6 specimens of each treatment were analyzed.

PMG concentrations in the exposure media (PMG_{GBH}) were analyzed by ion chromatography (Zhu et al. 1999), using a Dionex DX-100 chromatograph with a conductivity detector and a $25\ \mu\text{L}$ sample loop. Dionex AG-4 and AS-4 were used as analytical columns. A mixture of $\text{NaOH}/\text{CO}_3^{2-}$ (4 mM/9 mM) was used as eluent with a flow rate of $1\ \text{mL min}^{-1}$. Samples from three tanks and standards (0, 0.5, 1, 5, 10, 25 mg PMGL^{-1}) freshly prepared were injected. PMG standard was purchased from Sigma (St. Louis, MO, USA). The detection limit of the method was $0.1\ \text{mg L}^{-1}$. Data acquisition was performed using the Clarity Lite software.

The statistical comparisons for morphometric (L_T , W and K), SSA and elemental composition parameters between exposed groups and the control were performed using Kruskal–Wallis non-parametric ANOVA followed by Dunn's Multiple Comparison Test (GraphPad Prism Software). The level of significance for all statistical tests was set at $\alpha=0.05$.

Results and Discussion

The analytical concentrations of PMG_{GBH} (mean \pm standard error) at the initial time were: not detectable for control group, 0.9 ± 0.1 and $9.2 \pm 0.4\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$ for

the groups exposed to nominal concentrations of 1 and $10\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$, respectively. A significant reduction in PMG_{GBH} occurred at the end of the exposure time, being PMG_{GBH} no detectable ($<0.1\ \text{mg L}^{-1}$) in the control and the lower concentration tanks; and $1.1 \pm 0.2\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$ in the higher concentration tank. Moreover, condition factor (K) values at the final time were: 0.78 ± 0.05 for the control group; 0.83 ± 0.05 and 0.83 ± 0.04 for 1 and $10\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$, respectively. No statistically significant changes in fish morphometric parameters (L_T , W and K) were found among treatments.

Scanning electron micrographs of *O. bonariensis* gills at the end of the exposure period, are shown in Fig. 1.

Gill structure of control fish included filaments and lamellae showing normal morphological features (Fig. 1a). These characteristics are in accordance to those previously reported by other authors for this species (Tano de la Hoz et al. 2014; Vigliano et al. 2006). On the other hand, fish exposed to 1 and $10\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$ exhibited gill impairment. Gill of fish exposed to $1\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$ (Fig. 1b) showed signs of disorganization, while these alterations in fish exposed to $10\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$ (Fig. 1c) were severe, in both the lamellae as well as in the interlamellar epithelium spaces. The gill damages induced by GBH include mucous globules, both in epithelium of the filaments as well as in the mucosal cells of the interlamellar space and primary lamellae. The mucus production may be interpreted as a nonspecific stress response as well as the evidence of a protective role. These effects have been previously reported by several authors in gills of fish exposed to other environmental pollutants (Ferrari et al. 2009).

The fish gill is a multipurpose organ that plays a central role in the responses to environmental changes. In this sense, gill ultrastructural biomarkers can be useful tools to characterize the health status of fish in polluted environment. Several studies have shown the adverse histopathological impacts of commercial formulations of glyphosate after acute or chronic exposure (Hued et al. 2012; Jiraungkoorskul et al. 2003; Nešković et al. 1996). Filament cell proliferation, lamellar cell hyperplasia, lamellar fusion, epithelial lifting, and aneurysm were also observed in the gills. Cavalcante et al. (2008) determined genotoxic effects by the comet assay in gill cells of *Prochilodus lineatus* after sublethal acute exposure to $10\ \text{mg L}^{-1}$ of Roundup.

Other authors have also described changes in gill ultrastructure in fish exposed to glyphosate formulations (Samanta et al. 2016; 2018a, b; Senapati et al. 2009).

In order to perform a quantitative analysis of glyphosate exposure impact, the gills SSA was measured (Fig. 2). In the exposed groups an SSA increase was observed that was statistically significant for the group exposed to $1\ \text{mg PMG}_{\text{GBH}}\ \text{L}^{-1}$.

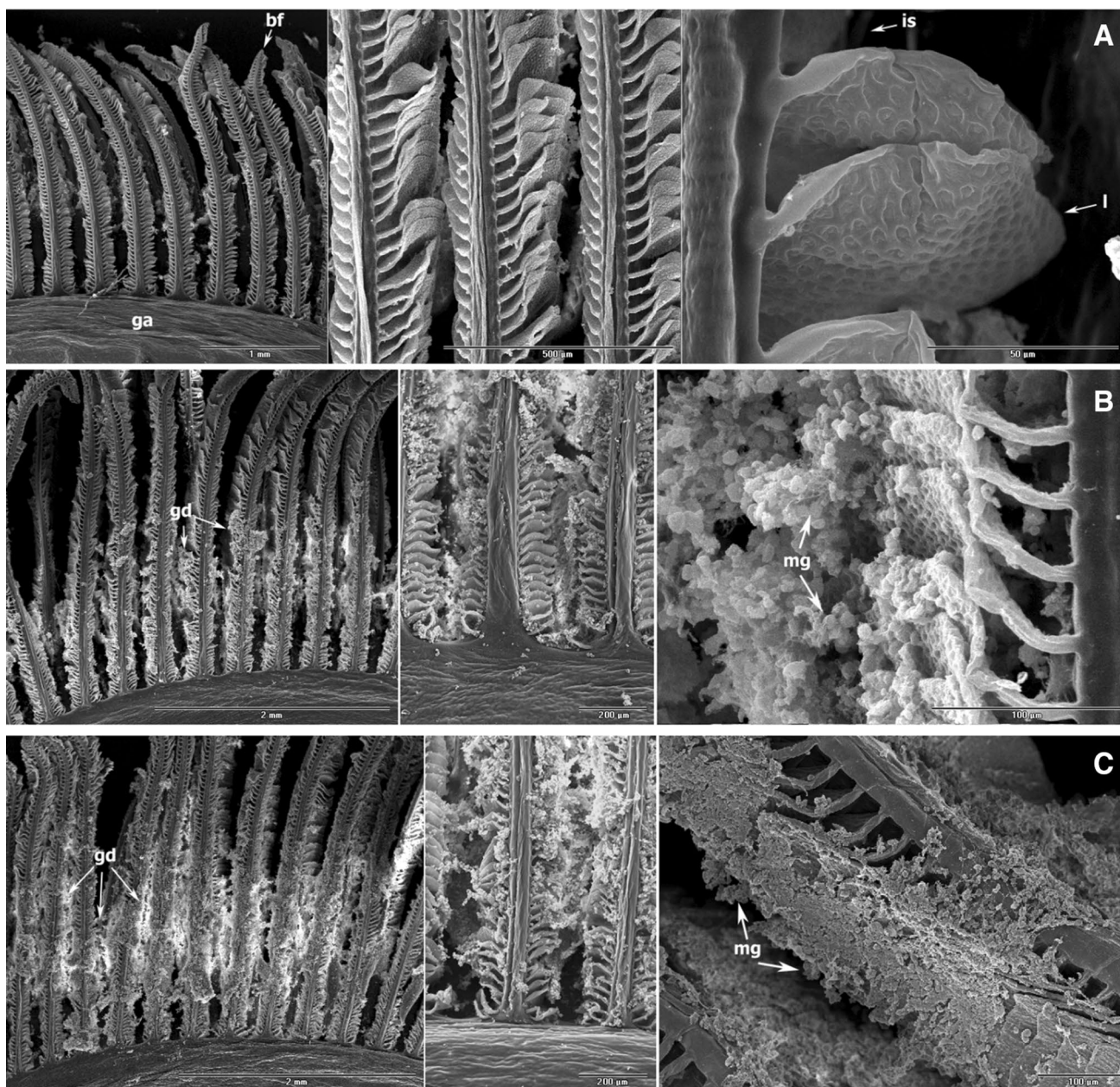


Fig. 1 Scanning electron micrographs of a holobranch of the freshwater fish *Odontesthes bonariensis* showing gill arch (ga), branchial filament (bf), the lamellae (l) and interlamellar spaces (is). Signs of gill disorganization (gd) including a large number of mucous glob-

ules (mg), are observed in the exposed groups. Representative micrographs at different magnification are shown in each panel. **a** Gills of control group specimens. **b** Gills of specimens exposed to 1 mg $\text{PMG}_{\text{GBH}} \text{L}^{-1}$. **c** Gills of specimens exposed to 10 mg $\text{PMG}_{\text{GBH}} \text{L}^{-1}$

The pore volume was $0.38 \pm 0.01 \text{ cm}^3 \text{ g}^{-1}$ for the control group. An increasing trend was observed when the treatment concentration increased, around 10% and 18% for the groups exposed to 1 and 10 mg $\text{PMG}_{\text{GBH}} \text{L}^{-1}$, respectively. This increase is indicative of the ultrastructure damage.

In addition, the results were grouped by sex and significant differences due to this factor were ruled out.

Finally, the elemental composition of gills was determined as shown in Table 1. No difference was observed in

the percentage composition of any of the elements in the exposed groups with respect to the control group. These results indicate that GBH effects involve a tissue disintegrative process without elemental composition changes. The adverse effects could be due to the active ingredient, co-adjuvants and/or its metabolites.

The present results are complementary to those provided by studies performed in other species. The lowest concentration tested is environmentally relevant, as there

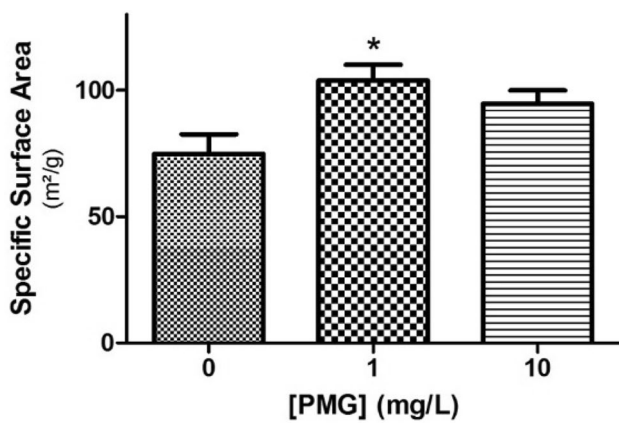


Fig. 2 Gill specific surface area of the freshwater teleost *Odontesthes bonariensis* (control group or exposed to 1 and 10 mg PMG_{GBH} L⁻¹) determined by N₂ (g) adsorption–desorption isotherms. Data as means ± standard errors (n=5–6). Asterisks indicate significant differences from control (*p<0.05)

Table 1 Gill elemental composition (%) of the freshwater teleost *Odontesthes bonariensis* determined by EDS for control group or exposed to 1 and 10 mg PMG_{GBH} L⁻¹

| Element | 0 | 1 | 10 |
|---------|-------------|-------------|-------------|
| C | 59 ± 1 | 60 ± 1 | 61 ± 2 |
| N | 17.7 ± 0.2 | 16.3 ± 0.03 | 15.1 ± 2.3 |
| O | 22.5 ± 1.3 | 23.0 ± 1.4 | 23.1 ± 0.1 |
| P | 0.46 ± 0.04 | 0.42 ± 0.12 | 0.49 ± 0.07 |
| S | 0.49 ± 0.01 | 0.47 ± 0.12 | 0.54 ± 0.17 |
| Ca | 0.07 ± 0.05 | 0.08 ± 0.02 | 0.12 ± 0.03 |

Data as means ± standard errors

have been reports of background concentrations of PMG close to 1 mg L⁻¹ in water bodies of agricultural areas of the Pampas region (Peruzzo et al. 2008). In the present work, that concentration lead to changes in gill ultrastructure of pejerrey fish. These effects were supported by SEM and SSA analyses.

For future research, it would be interesting to investigate the impact of lower PMG concentrations and longer exposures on gill ultrastructure taking into account that PMG is considered a pseudo-persistent pesticide.

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