

Dynamics of Aromatase and Physiological Indexes in Male Fish as Potential Biomarkers of Anthropogenic Pollution

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Abstract Endocrine disruption on aquatic wildlife is being increasingly reported, and the changes in gene aromatase expression are used as indicators. However, natural fluctuations in brain and gonadal aromatase expression and physiological indexes have not been previously measured in a fish species (Jenynsia multidentata) throughout a complete reproductive cycle, nor the biological effects of anthropogenic inputs on these responses. Accordingly, males were monthly collected over a year in both, a reference and a contaminated site. Physicochemical analyses of water samples were done and reflected a strong anthropogenic impact. Brain aromatase fluctuates along the reproductive cycle of this species and, noticeably, the increase of brain gene expression begins with a 1 month delay in the contaminated site. This mismatch is also evidenced for testes weight. Hepatosomatic index also revealed adverse

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effects in the polluted site. In turn, the alterations observed in biological responses could be affecting the reproduction of this fish species.

Keywords Water pollution · Sewage effluents · Biomarkers · Aromatase · Somatic indexes · Fish

Endocrine disruption in wildlife is being increasingly reported worldwide, mainly on aquatic environments. Endocrine disrupting compounds (EDCs) includes many natural or manmade substances that are able to alter the functioning of the endocrine system, and therefore, represents a significant threat for aquatic organisms and human health. Wastewater treatment plants (WWTP) are commonly not designed to eliminate EDCs and therefore suppose unavoidable chemical pollution sources in freshwater ecosystems because of the incomplete elimination of pollutants during water treatment processes (Gomes et al. 2003). Among the environmental contaminants that have been identified in sewage effluents, xenoestrogens (e.g. phytoestrogens, mycoestrogens and $17-\alpha$ -ethynylestradiol) are of special interest because of their ability to induce biological responses similar to those caused by the natural hormone 17-β-estradiol. Additionally, there are other chemicals frequently found in WWTP discharges that have a weak estrogenic activity (e.g. alkylphenols, bisphenol A and pesticides). In most South American rivers, due to insufficient infra-structure, untreated or poorly treated wastewaters are released into the environment (Bertin et al. 2011).

Biomarkers are early warning signals for detecting environmental stress and long-term negative effects on populations and communities. They also allow achieving cleaning actions before irreversible effects occur (Van der Oost et al. 2003). The general state of health individuals is used as

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marker to individual level since they are influenced by generic stressors ranging from physicochemical parameters to contaminants (Kime 1997). At lower level of biological organization, the study of changes in gene expression allows an early detection of toxic effects occurring due to pollution charges (Contardo-Jara and Wiegand 2008). In this regard, aromatase genes are increasingly used as indicators of endocrine disruption in ecotoxicological studies. Cytochrome P450 aromatase is the steroidogenic enzyme that catalyzes the conversion of androgens to estrogens (Gonzalez and Piferrer 2003). Euteleosts fish express two structurally and functionally different P450 aromatase isoforms, termed Cyp19a1a (preferentially expressed in gonads), and Cyp19a1b (preferentially expressed in brain). Interestingly, both aromatase genes are potential target for EDCs, mainly xenoestrogens. Many environmental contaminants can modulate its expression or activity altering the rate of estrogen production and disturbing local and systemic levels (Cheshenko et al. 2008). In fish species collected in rivers that receive WWTP discharges an increase in brain aromatase was observed (Geraudie et al. 2011), suggesting the presence of estrogenic compounds. However, the effects of this type of anthropogenic inputs on aromatase expression and physiological indexes throughout a complete reproductive cycle have not been previously analyzed in fish species.

Given this background, the aim of the present study was: First, to characterize the natural monthly fluctuations during a year in a reference site in aromatase expression and fish health indexes in the fish *Jenynsia multidentata*. Second, evaluate the effects of anthropogenic contamination on aromatase and condition indexes. *J. multidentata* is a fish widely distributed in the Neotropical Region (Malabarba et al. 1998) and it has been used as an excellent model in both laboratory and field studies mainly because of its ability to adapt to a wide variety of environments, including poor water quality conditions (Hued and Bistoni 2005). To the best of our knowledge; this is the first characterization of natural seasonal fluctuations in aromatase expression in *J. multidentata*. Furthermore, this work contributes to the understanding about the effects of anthropogenic pollution on this specific biomarker of endocrine disruption considering a complete reproductive cycle.

Materials and Methods

The Suquía River basin (Córdoba province, Argentina) originates at the San Roque dam, flows across Córdoba city, and drains into the depression of Mar Chiquita Lake. This lake and the mouth of its tributaries are considered as a Ramsar site (wetland of international concern included in the list of the Ramsar Convention) and it is located 150 km downstream from Córdoba City (Fig. 1). At the lower basin, the Suquía River receives sewage discharge from the only WWTP of Bajo Grande. The clearance capacity of this plant is overreached and, as a consequence, discharges wastewaters without previous treatment directly into the river (Hued and Bistoni 2005). The inputs of urban and industrial wastes as well as agrochemicals from many industries and crops established on the margins of this river have contributed to an increased amount of toxic effluents entering the river.

Sampling sites upstream and downstream of the WWTP discharge area were established (Fig. 1) considering previous reports on the water quality of the basin (Wunderlin et al. 2001; Valdés et al. 2014). Yuspe River is located 30 km upstream from the San Roque dam and is an already established reference sampling site. Primero River, the downstream site, is located at 70 km from the WWTP discharge and receives varying concentrations of contaminants from different sources.

Every 2 months, a water sample was collected at each sampling site to describe the water quality during 2010. Dissolved oxygen, conductivity, pH and water temperature were monitored *in situ* using a WTW multiparametric

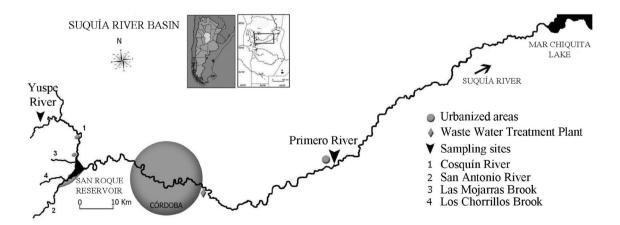


Fig. 1 Sampling stations selected along the Suquía River basin

equipment, while 5-day biological oxygen demand (BOD-5), ammonia-nitrogen, nitrite-nitrogen, nitrate-nitrogen, orthophosphate-phosphorous, chloride, sulfates, hardness, calcium, magnesium, total solids, and total coliforms were measured at the laboratory, according to APHA et al. (2005) methodologies. These physicochemical parameters were integrated into a Water Quality Index (WOI) (Pesce and Wunderlin 2000). Thirteen specimens of J. multidentata adult males were monthly caught at each site using a standard backpack electrofisher. The fish were transferred alive to the laboratory in oxygenated water tanks (20 L) and sacrificed with an overdose of tricaine methanolsulfonate (MS-222). Fish were weighted and total length recorded. After that, they were dissected and liver and gonad were weighed to determine of general state of health of individuals (van der Oost et al. 2003): Condition factor (CF = [(body)weight/total length) \times 100.000]); Gonadosomatic index as: $GSI = [(gonad weight/(total body weight) \times 100)]$ and Hepato-somatic index as: HSI=[(liver weight/(total body weight) × 100)]. Brains and gonads were quickly dissected and stored directly in RNAlater (QIAGEN) at -80°C for qPCR analysis. For cyp19a1 mRNA quantification, total RNA was extracted from brain and gonadal tissue by the guanidine thiocyanate-phenol chloroform extraction method in accordance with Chomczynski and Sacchi (1987). Nonspecific reverse transcription was performed from individual tissue total RNA. Quantitative polymerase chain reaction was performed with a Bio-Rad iO cycler and was used to amplify and measure the transcript abundance of cyp19a1a in testis and cyp19a1b in brain using specific J. multidentata primers for real-time polymerase chain reaction (Guyón et al. 2012a). Normalized expression levels for target genes were generated using the standard curve method with J. multidentata B-actin as reference gene (Larionov et al. 2005). Relative fold changes to reference sampling site were also calculated.

Statistical analyses were carried out using Infostat Software Package (Di Rienzo et al. 2011). All results are expressed as mean \pm standard deviation. Normal distribution of data was controlled using a Shapiro-Wilk's test ($p \le 0.05$), and Levene test was used to check the homogeneity of variance. In order to find significant differences between sites, months and to see if the pattern of results is the same between sites over the months, a two factors variance analyses on ranks transformed variables was performed (Hollander and Wolfe 1973).

Results and Discussion

During the last decades the Suquía River basin has been intensively studied not only for the detection of detrimental changes in water quality due to anthropogenic activities, but also to determine the deleterious effects on aquatic biota (e.g. Pesce and Wunderlin 2000; Hued and Bistoni 2005; Guyón et al. 2012b; Hued et al. 2012). The historical reports of water quality reveal the progressive deterioration of this aquatic system and it was confirmed in this study. Water quality was severely affected downstream the WWTP showing significant differences for all the measured parameters between sites (Table 1). The drastic diminution of dissolved oxygen, the increase in total coliforms and the high concentration of nitrogen species (nitrate, nitrite and ammonia) were the main factors affecting water quality, and they are frequently associated with sewage inputs (Wunderlin et al. 2001). Other parameters such as conductivity, total solids, hardness, calcium, magnesium, chloride and sulfates were significantly increased in the polluted site. It has been proposed that these parameters are linked to urban, industrial and agricultural runoff (Wunderlin et al. 2001; Pasquini et al. 2012). The presence of pesticides, hormones, pharmaceuticals and heavy metals confirms those sources of pollution (Monferrán et al. 2011; Maggioni et al. 2012; Bonansea et al. 2013a; Valdés et al. 2014). WOI registered values ranged from 51 to 60 in Primero River. Hued and Bistoni (2005) pointed out a WQI value close to 50 seriously difficult aquatic life. The CF was not significantly different between sites when analyzing all the studied period (F = 0.27; p = 0.607; Fig. 2a). However, differences among months were observed when considering both sites together (F=5.73; p < 0.001), indicating variations in the weight/ length relationship along the year. The pattern of change was similar in both sites, as no interaction between months and sites was observed (F = 0.50; p = 0.901). The fact that there was no decrease in the CF suggests that the fish able to inhabit Primero River locality were not overtly affected by exposure to occurring contaminants with respect to somatic growth. This result is in line with studies showing not changes in this index in fish sampled in polluted places (Hinfray et al. 2010).

The contrast between the low HSI values recorded in males coming from the reference site (Yuspe River) and the high HSI values in samples from Primero River locality (F = 76.31; p < 0.001; Fig. 2b) may be linked to the presence of compounds capable of inducing metabolic activity in fish livers in the polluted site. Accordingly, elevated HSI have been detected in species exposed to WWTP effluents (Barber et al. 2007; Vajda et al. 2011). This increase in liver weight could be an unspecific response to contaminants, induced by an augmented demand of enzymatic activity. In this regard, Maggioni et al. (2012) reported increased levels of antioxidant enzymes in female livers of J. multidentata collected in the same polluted site (Primero River). Moreover, the same authors detected histopathological liver damages such as hypertrophia, hydropic degeneration, necrosis and fibrosis, all damages that have been associated with

Variables	January		March		May		August		October	
	Yuspe River	Primero River	Yuspe River	Primero River	Yuspe River	Primero River	Yuspe River	Primero River	Yuspe River	Primero River
Ammonia-nitrogen 0.39 ± 0.54 a	0.39±0.54 a	5.00±0.22 b	1.65±0.67 a	1.52±0.52 a	0.18±0.03 a	1.31 ±0.22 b	0.087±0.003 a	11.69±0.99 b	0.87±0.64 a	0.86±0.32 a
Biological oxygen demand after 5 days (BOD)	1.35±0.07 a	4.50±0.14 b	0.30±0.14 a	4.11 ± 0.14 b	1.00±0.14 a	1.58±0.01 b	1.35±0.01a	1.22±0.01 b	1.71±0.14 a	2.33±0.01 b
Chloride	3.91 ±0.02 a	89.33±0.69 b	3.91±0.03 a	$77.61 \pm 0.69 \text{ b}$	3.91±0.01 a	$80.54 \pm 0.69 \text{ b}$	3.91 ±0.04 a	$112.76 \pm 0.69 \text{ b}$	3.91±0.02 a	79.08 ± 1.38 b
Conductivity [µS cm ⁻¹] ^a	72.00±1.41 a	980.00 ± 7.07 b	49.00±1.41 a	978.00±2.83 b	82.00±1.41 a	967.00±1.56 b	148.00±1.71 a	1351.00 ± 1.72 b	131.01 ±2.83 a	1335.00±1.78 b
Dissolved oxygen ^a	10.95±0.07 a	7.56±0.03 b	10.17 ± 0.03 a	7.36 ± 0.01 b	10.70 ± 0.14 a	6.88 ± 0.01 b	10.03±0.01 a	4.84 ± 0.01 b	9.02±0.03 a	5.57 ± 0.03 b
Nitrates-nitrogen	0.88±0.09 a	31.07 ± 2.20 b	0.25±0.03 a	7.24±2.45 b	0.51±0.12 a	$9.11 \pm 1.47 \text{ b}$	0.30±0.01 a	2.65 ± 1.48 b	0.36±0.08 a	$7.50 \pm 0.06 \text{ b}$
Nitrites-nitrogen	$0.09 \pm 0.04 \text{E-8}$ a	$0.28 \pm 0.01 \text{ b}$	0.09±0.01 a	$0.18 \pm 0.09 \text{E-2 b}$	$0.01 \pm 0.01 \text{E-8}$ a	1.75 ± 0.02 b	0.011 ± 0.003 a	$0.79 \pm 0.04 \text{ b}$	0.076 ± 0.001 a	0.177 ± 0.002 b
Orthophosphates phosphorous	<dl< td=""><td>0.74 ± 0.16</td><td><dl< td=""><td>0.28 ± 0.04</td><td><dl< td=""><td>0.25 ± 0.03</td><td><dl< td=""><td>1.37 ± 0.02</td><td><dl< td=""><td>0.07 ± 0.04</td></dl<></td></dl<></td></dl<></td></dl<></td></dl<>	0.74 ± 0.16	<dl< td=""><td>0.28 ± 0.04</td><td><dl< td=""><td>0.25 ± 0.03</td><td><dl< td=""><td>1.37 ± 0.02</td><td><dl< td=""><td>0.07 ± 0.04</td></dl<></td></dl<></td></dl<></td></dl<>	0.28 ± 0.04	<dl< td=""><td>0.25 ± 0.03</td><td><dl< td=""><td>1.37 ± 0.02</td><td><dl< td=""><td>0.07 ± 0.04</td></dl<></td></dl<></td></dl<>	0.25 ± 0.03	<dl< td=""><td>1.37 ± 0.02</td><td><dl< td=""><td>0.07 ± 0.04</td></dl<></td></dl<>	1.37 ± 0.02	<dl< td=""><td>0.07 ± 0.04</td></dl<>	0.07 ± 0.04
pH ^a	8.75±0.35 a	8.22±0.03 a	7.92±0.03 a	$8.58 \pm 0.02 \text{ b}$	8.15±0.03 a	7.44 ± 0.01 b	8.94±0.03 a	7.53±0.01 b	8.80±0.14 a	7.85 ± 0.14 b
Solids: total	76.40±0.14 a	1008.40 ± 0.14 b	56.80±0.28 a	701.60±0.28 b	40.00±1.41 a	400.00 ± 1.41 b	96.40±0.14 a	891.60 ± 0.14 b	58.40±0.28 a	676.00±2.83 b
Sulfates	0.29 ± 0.33 a	132.90 ± 1.61 b	0.37±0.03 a	132.27 ± 0.89 b	0.35 ± 0.06 a	67.14±8.02 b	0.25 ± 0.04 a	121.99±1.34 b	0.35 ± 0.06 a	67.14 ± 8.02 b
Temperature ^a [°C]	30.50±0.28 a	24.40 ± 0.24 b	16.71 ±0.49 a	22.60 ± 0.57 b	12.30±0.28 a	11.70±0.49 a	18.20±0.57 a	15.00 ± 0.57 b	18.01±0.35 a	17.30±0.42 a
Total coliforms	$1.50E + 04 \pm 7.07 a$		$8.80E + 05 \pm 7.07 b$ $2.00E + 02 \pm 2.83 a$	$1.20E + 05 \pm 2.83 b$	$5.00E + 02 \pm 1.41 a$	$2.40E + 05 \pm 1.41 b$		$2.20E + 03 \pm 1.41 a$ $1.50E + 05 \pm 1.41 b$	$5.00E + 02 \pm 2.83 a$	$5.00E + 03 \pm 2.83 b$
Hardness	29.33 ± 7.32 a	337.53 ± 15.45 b	20.70 ± 0.07 a	317.41 ± 6.51 b	28.75±1.63 a	281.75±4.88 b	57.5±3.3 a	292.68 ± 0.81 b	50.6±0.1 a	289.23 ± 0.81 b
Calcium	7.14±1.63 a	98.41 ± 6.84 b	6.45±1.30 a	92.18±3.42 b	8.76±0.65 a	88.96 ± 0.65 b	14.75±3.63 a	100.02 ± 1.96 b	8.30±1.30 a	87.81 ± 0.33 b
Magnesium	2.79±2.77 a	22.36±7.90 b	1.12±0.79 a	21.24 ± 1.58 b	1.68±0.79 a	14.53±1.58 b	5.03±0.79 a	10.48 ± 0.99 b	7.27±0.79 a	17.05 ± 0.88 b
MQI	74.00 ± 2.33	51.70 ± 1.89	82.70 ± 0.56	54.30 ± 1.00	88.70 ± 1.00	57.00 ± 1.80	88.70 ± 2.00	51.70 ± 1.13	78.70 ± 0.99	62.70 ± 0.71

 Table 1
 Water quality parameters of selected sampling sites in Suquía River basin

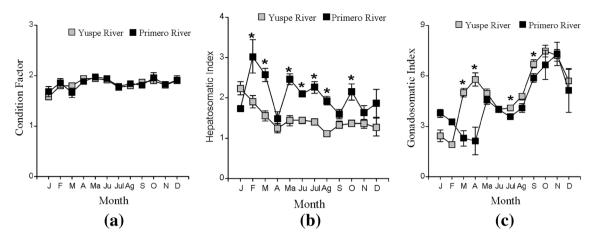
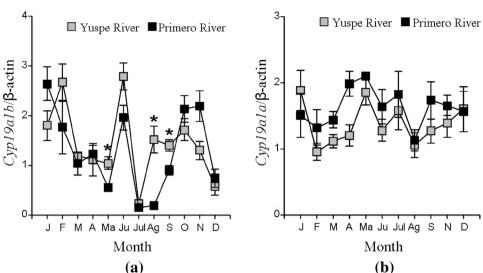


Fig. 2 Condition factor (a), Hepatosomatic (b) and Gonadosomatic (c) index measured in two sites with different water quality during an annual cycle. Significant differences between sites are indicated with *asterisk*

pesticide and wastewater exposure (Ballesteros et al. 2007; Hued et al. 2012). When considering both sites together, significant differences among months were observed (F = 6.86; p < 0.001), showing the highest values of HSI during summer (January, February, March). The presence of interaction between months and sites (F=3.40; p < 0.001) indicated variations in the pattern of change at each site along the year.

When considering all the studied period, the gonad growth seems to be affected by exposure to the contaminants as is shown by the GSI reduction at the polluted site (F = 19.31; p < 0.001). This result is similar to the reported for other species after exposure to WWTP discharges (Hinfray et al. 2010; Vajda et al. 2011). When considering both sites together, significant differences were observed among months (F=31.28; p < 0.001), showing the highest values of GSI during spring (September, October, November, December). Both sites showed two peaks in GSI during the breeding season that goes from August to April (Fig. 2c); however, the pattern of change was significantly different for each site, as indicated by an interaction between month and site (F = 5.01; p < 0.001). In the reference site, an increase in GSI was observed since July reaching a maximum in October, while the other increase started in March with maximum in April. At Primero River, a marked delay in the GSI increase was observed. The first peak started in August with maximum in November, while the second peak was in May. Consistent with this result, Bianco (2011) observed the same pattern when analyzing the reproductive cycle of J. multidentata females in the same year and sampling sites than this study. This author found a 1 month delay in the presence of embryos in the ovary in the site with elevated pollution. It might be possible that the amount of chemical compounds present in polluted environments can cause endocrine disruption and negative effect on fish reproduction (Jobling et al. 2002) and could also modify the beginning and length of the reproductive cycle.

WWTP discharges are linked with EDCs mainly xenoestrogens (Vajda et al. 2011). As a consequence, an increase in brain aromatase could be expected since in laboratory studies estrogenic compounds up-regulate its expression and activity (Vosges et al. 2011). Accordingly, Geraudie et al. (2011) found that aromatase activity was significantly upregulated in wild fish collected in river that receives WWTP effluents, suggesting the occurrence of estrogenic compounds and their involvement in aromatase activity modulation. In the study area, downstream the WWTP of Bajo Grande discharge, Valdés et al. (2014) detected estrogenic compounds like estrone. Given this background, we hypothesized to found elevated brain aromatase expression in males collected at this site. However, we found lower levels of mean seasonal values than the reference site (1.41 for Yuspe River and 1.21 for Primero River), without statistical differences between sites (F=3.11; p=0.0801). Laboratory studies showed that the pesticide chlorpyrifos and its commercial mixture with cypermethrin inhibit brain aromatase expression in J. multidentata females (Bonansea et al. 2013b). Both compounds have been detected in Primero River at concentrations that exceeded the international limits for aquatic life preservation (Bonansea et al. 2013a). Additionally, in a previous study conducted with wild females of this species collected in a pollution gradient, a cyp19a1b inhibition was observed downstream the WWTP of Bajo Grande (Guyón et al. 2012b). In this regard, fish are exposed to a complex mixture of pollutants some of which could act inhibiting aromatase expression. Fish brain aromatase is characterized by an elevated expression and activity during the spawning period in several species, suggesting a role in the control of the reproductive cycle (Gonzalez and Piferrer 2003; Cheshenko et al. 2008). Accordingly, when considering both sites together, significant differences among months were observed (F = 20.29; p < 0.001), with maximum values of aromatase expressed



during the reproductive season of this species according to the reproductive period description of Goyenola et al. (2011). In the reference site, the gene abundance showed an enhancement from August to October, and afterwards another increase in January to February. From March to July, brain aromatase expression was lower, however a peak was observed in June. In the polluted site an increase in aromatase expression was observed from September to November and another augment in January. This result shows a significant mismatch in the increase of the gene expression beginning with 1 month delay relative to reference site, indicating that the seasonal pattern of variation was different (Fig. 3a, F = 5.18; p < 0.001). Fishes from the reference sampling site had approximately 1.6, 6.4 and 1.5-fold higher cyp19a1b expression than fishes from the polluted sites in May, August and September, respectively.

This mismatch could be related to offsets in the beginning of the reproductive cycle due to pollution. Similarly to Yuspe River, a peak in June was observed in Primero River. In this regard, Gonzalez and Piferrer (2003) observed a peak of aromatase activity in *D. labrax* in a time when an overall reorganization of the gonadal tissues occurs after the spawning season and the authors suggested that these phenomena could somehow be connected.

Gonadal aromatase (Fig. 3b) showed significant differences between sites when considering the studied period (F=5.89; p=0.017), being 1.37 the value for Yuspe River and 1.65 for Primero River, but no differences were observed when considering month to month between sites. All along the studied period there were differences between months (F=2.28; p=0.016); and no interaction was registered between months and sites (F=0.63; p<0.786), indicating that the seasonal changing pattern was similar for both sites. The effects of WWTP effluents exposure on this parameter do not appear to follow a general rule. Lavado et al. (2004) informed lower gonadal aromatase activity in males of *C*. *carpio* collected downstream a WWTP discharge, while Douxfils et al. (2007) did not find differences in males of two species coming from sites upstream and downstream WWTP outfall. In *J. multidentata, cyp19a1a* expression was significantly higher in Primero River than Yuspe River. This result suggests the presence in the river of compounds capable of modulate this variable without causing an estrogenic response in *cyp19a1b* expression.

In summary, urban wastes released from the WWTP of Bajo Grande, which has overreached its capacity, coupled with pesticides, industry and agricultural runoff from the margins of the Suquía River are the major causal factors and may have cumulatively negative effects on the environmental quality of the waters. Consequently, fish population studied is exposed to complex xenobiotic mixture, where the occurrence and activity of estrogenic and antiestrogenic compounds cannot be excluded. In J. multidentata coming from the reference site, brain aromatase fluctuates along the reproductive cycle with maximum values during the breeding season. The mismatch observed in the polluted site not only for gene expression but also for GSI values could interfere with the fish reproduction altering the beginning and length of the reproductive cycle. Moreover, the results make out the maintenance of the spatial deterioration in water quality that has been acting for more than 15 year, which not only provoke damage on organism's health but also could result in gradual loss of the reproductive function and important alterations at ecosystem levels upon long exposure.

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