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Bisphenol-A alters gene expression and functional parameters in molluscan hepatopancreas

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Key words: xenoestrogens, bivalve molluscs, hepatopancreas, estrogen receptors, metallothionein,

redox balance,

Abstract

Bisphenol A (BPA) is a well-known xenoestrogen in mammalian systems that can affect reproduction also in aquatic organisms. In this work the possible effects of BPA were investigated in the hepatopancreas of the bivalve mollusc *Mytilus galloprovincialis*: mussels were injected with different amounts of BPA (3-60 ng/g dw tissue) and tissues sampled at 24 h post-injection. Expression of different *Mytilus* genes was evaluated by RT-Q-PCR: BPA exposure increased the expression of MeER2 and induced downregulation of antioxidant genes, catalase and metallothioneins. Moreover, BPA induced changes in activity of catalase, GSH transferase (GST) and GSSG reductase (GSR), and in total glutathione content. A decrease in lysosomal membrane stability and increased neutral lipid accumulation were also observed. The results were compared with those obtained with similar concentrations of 17β -estradiol. These data demonstrate that BPA can alter gene expression, activities of enzymes involved in redox balance, and lysosomal function in molluscan hepatopancreas, a tissue involved in the control of metabolism and gamete maturation. Overall, these data indicate that BPA, at environmentally relevant concentrations, can have both estrogen-like and distinct effects in invertebrates like in vertebrates.

Keywords: xenoestrogens, molluscs, *Mytilus* digestive gland, gene expression, estrogen receptors, antioxidant defences, lysosomes.

1. Introduction

Xenoestrogens are environmental chemicals that mimic the natural steroid 17β-estradiol, potentially interfering with the physiological action of the hormone. The concept of estrogenicity is generally based on the property of these compounds to bind estrogen receptors (ERs), and to act as transcription factors, when binding to the estrogen response elements (ERE) in the DNA, thus affecting gene expression (Soto et al., 2006). However, estrogens can also act through rapid, 'non genomic' mechanisms of action that may be initiated at either membrane or cytosolic locations and can result in both direct local effects (such as changes in ion fluxes) and regulation of gene transcription secondary to activation of cytosolic kinase cascades (Levin, 2005; Vasudevan and Pfaff, 2007). Recent studies demonstrated that xenoestrogens can rapidly and potently elicit signaling through non genomic pathways culminating in functional endpoints (Walsh et al., 2005; Ropero et al., 2006; Tabb and Blumberg, 2006; Waring and Harris, 2005; Watson et al., 2007).

For invertebrates, which represent 95% of animal species, information on the effects and mechanisms of action of estrogens is scarce compared to that available in vertebrates, given the complexity of endocrine systems in different phyla (Oehlmann et al., 2006; Porte et al., 2006). However, this subject is receiving increasing attention due to the concern posed by the presence in the environment of both natural and synthetic steroid estrogens, as well as a variety of estrogen-mimicking chemicals, that represent a potential hazard for humans and wildlife species (McLachlan, 2001; Rotchell and Ostrander, 2003; Waring and Harris, 2005). The presence of most vertebrate sex steroids and steroidogenic pathways has been described in different invertebrate groups, molluscs in particular, where the role of estrogens has been mainly investigated in the control of gametogenesis (Porte et al., 2006; Janer and Porte, 2007). Estrogen-like receptors have been recently characterized in gastropods (Thornton et al., 2003; Kajiwara et al., 2006) and cephalopods (Keay et al., 2006) and ER-like sequences have been identified in bivalves (Puinean et al., 2006). Molluscan ERs, although phylogenetically clustered with other steroid receptors,

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appear to be functionally different: their transcriptional activity is constitutive, and not activated by estrogens (Thornton et al., 2003; Keay et al., 2006). However, in ganglia and immune cells of the bivalve *Mytilus*, 17β-estradiol (E_2) has been shown to act through rapid, 'alternative' modes of action through modulation of Ca²⁺- and kinase-mediated cascades (Stefano et al., 2003; Canesi et al., 2004a, 2006). Moreover, we have recently demonstrated that in *Mytilus* hepatopancreas (or digestive gland), a tissue that plays a central role in metabolism, E_2 modulates different functional parameters and increases the expression of antioxidant genes (Canesi et al., 2007). These data support the hypothesis that, although the molluscan ERs do not appear to mediate genomic estrogen signaling, conserved nongenomic mechanisms via 'alternative' modes of action are likely candidates for this role (Keay et al., 2006).

The xenoestrogen Bisphenol A-BPA (2,2-bis-(4-hydroxyphenyl)-propane) is a monomer used in the manufacture a variety of plastic products. In mammalian systems, BPA induces estrogenic activity through both genomic and rapid, non genomic pathways, involving both ER-dependent and ER-independent mechanism of action, depending on both cell and tissue type (Gould et al., 1998; Quesada et al., 2002; Singleton et al., 2004; Maffini et al., 2006; Ropero et al., 2006). BPA is produced in large amounts and it is detected at high levels in human samples as well as in aquatic organisms (vom Saal and Hughes, 2005), where it has been shown to induce significant disruption of reproductive output, with molluses being apparently more sensitive than fish (Jobling et al., 2003; Ohelmann et al., 2006). Concentrations of BPA ranging from ng to $\mu g/l$ can be found in the coastal marine environment (Pojana et al., 2004; Anderson et al., 2006).

In *Mytilus* immunocytes, BPA rapidly affected the lysosomal function both *in vitro* (Canesi et al., 2004b) and *in vivo* (Canesi et al, 2005); these effects were mediated by interference with components of rapid kinase-mediated cell signalling that are activated by E_2 (Canesi et al., 2004a; 2006; Porte et al., 2006). In this work, the possible effects of BPA on gene expression and functional parameters of mussel hepatopancreas were investigated and the results compared with those obtained with E_2 in the same experimental conditions (Canesi et al., 2007). Mussels were

injected with different amounts of BPA and tissues sampled at 24 hrs post-injection. In bivalves, direct administration into the blood (hemolymph) by injection into the adductor muscle sinus has been widely utilised for identifying the possible targets of both natural and environmental estrogens (Gagnè et al., 2001; Canesi et al., 2005, 2006, 2007; Le Curieux-Belfond et al., 2005; Gauthier-Clerc et al., 2006; Puinean and Rotchell, 2006). The concentrations of BPA (ng/g dry weight/mussel) were in the same range as those previously shown to affect mussel immune parameters in the same experimental conditions (Canesi et al., 2005) and comparable to environmental exposure levels (Pojana et al., 2004).

Expression of *Mytilus* ERs (MeER1, MeER2) as well as of other genes (catalase, metallothionein isoforms MT10 and MT20, p53, cathepsin L) was evaluated by quantitative RT-PCR. Moreover, the activity of enzymes involved in tissue redox balance (catalase, GSH transferase, GSSG reductase), and the concentration of total glutathione (GSH+GSSG), as well as lysosomal parameters (lysosomal membrane stability-LMS and accumulation of neutral lipids-NL, lipofuscin) were evaluated.

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2. Materials and methods

2.1 Animals and treatments

Mussels (*Mytilus galloprovincialis* Lam.) 4-5 cm long, were obtained from a fishermen cooperative (SEA-Gabicce Mare, PU) in October 2006. Mussels were kept for 1-3 days in static tanks containing artificial sea water (ASW) (1 l/mussel) at 16°C. Sea water was changed daily. In mussels of this size, average dry weight of soft tissues was 1 g, the average volume of hemolymph that can be withdrawn from each animal was about 1 ml; average concentration of free E_2 in the hemolymph of control mussels was 0,8-1 ng/ml (evaluated by a commercial competitive chemiluminescent enzyme immunoassay kit (Immulite 2000 Estradiol).

BPA was injected into the posterior adductor muscle of groups of 10-12 mussels each, using a sterile 1 ml syringe with a 18G1/2" needle as previously described (Canesi et al., 2005). In bivalve molluses, injection into the adductor muscle sinus allows for direct administration of estrogens into the animal open circulatory system leading distribution from the blood (hemolymph) to the tissues, including the hepatopancreas (Le Curieux-Belfond et al., 2005). This method has been largely utilised for evaluating the effects of natural and environmental estrogens in bivalves (Gagnè et al., 2001; Canesi et al., 2005, 2006, 2007; Gauthier-Clerc et al., 2006; Puinean and Rotchell, 2006). Each mussel was injected with 50 µl of BPA solutions (from a 10 mM stock solution in ethanol diluted in ASW), containing, respectively 3, 15, and 60 ng BPA, corresponding to a nominal concentration of BPA 3, 15, and 60 ng/g dry weight or per mussel. Similarly, groups of 10-12 mussels were injected with 1,35 and 6,75 ng 17 β -estradiol (E₂) (nominal concentrations of 1,35 and 6,75 ng/g dw or per mussel), as previously described (Canesi et al., 2006, 2007). For each experiment, a parallel set of control mussels were injected with 50 µl of a solution of ASW containing an equal amount of ethanol (≤0.005%). Mussels were then placed in plastic tanks containing ASW at 16 °C (0.5 l/mussel). Experiments were repeated three times. At 24 hrs post-

injection, digestive glands were rapidly dissected fro each group of mussels, pooled and frozen in liquid nitrogen and maintained at -80°C. For histological analyses, small pieces of digestive glands were placed on aluminium chucks, immersed in hexane pre-cooled to -70°C in liquid nitrogen and maintained at -80°C (Moore, 1976). Animals were not fed during the experiment.

2.2 RNA isolation and quantitative RT-PCR

Total RNA was extracted from 0.5 grams of frozen pools of digestive glands using Trizol Reagent (Sigma) following the manufacturer's protocol. The quality of the isolated RNA was checked by gel electrophoresis. The concentration and purity of the isolated RNA were assessed by absorption spectroscopy. Only samples with $OD_{260/280} > 1.8$ were processed. Aliquots of 1 µg of total RNA were reverse-transcribed into cDNA using 200 units of RevertAid H-Minus M-MuLV Reverse Trascriptase (Fermentas, Hannover MD, USA), in the presence of 250 ng of random examers (Invitrogen, Milan, Italy), 1 mM dNTPs (Promega, Milan, Italy) at 42°C for 60 min in a reaction volume of 20 µl. cDNA was then utilized to amplify the target sequences of interest using a Chromo 4TM System real-time PCR apparatus (Biorad, Milan, Italy). Aliquots of cDNA samples were suitably diluted in a final volume of 20 µl containing 1x iTaq SybrGreen Supermix with Rox (Biorad, Milan, Italy) and 0.25 µM of each primer (TibMolBiol, Genoa, Italy). The primer pairs utilized and their references are shown in Tab. 1. The thermal protocol consisted of 3 min initial denaturation at 95 C, followed by 40 cycles: 15 s at 95 C; 30 s at 55°C (30 s at 51°C and 54°C for MeER1 and MeER2 respectively); 20 s at 72°C. A melting curve of PCR products (55–94 °C) was also performed to ensure the absence of artefacts. Relative expression of target genes in comparison with the 18S reference gene was evaluated by the comparative C_T threshold method (Pfaffl, 2001) using the Biorad software tool Genex-Gene Expression MacroTM (Vandesompele et al, 2002). The normalized expression obtained was expressed as relative quantity of mRNA with respect to control samples. Data are the mean \pm SD fo three experiments in quadruplicate.

2.3 Determination of enzyme activities and glutathione content

Catalase: tissues were homogenized in 5 vol of homogenization buffer (20 mM Tris buffer, 0.5 M sucrose, 0,15 M NaCl, pH 7.6) and centrifuged at 500 x g for 15 min at 4°C. The supernatants were then centrifuged at 12,000 x g for 30 min. Both supernatant and pellet, (containing mitochondria and peroxisomes) were utilised for the spectrophotometric evaluation of catalase activity following the decomposition of H_2O_2 at pH 7, 25°C, at 240 nM as previously described (Viarengo et al., 1991).

GSH transferase (GST) and GSSG reductase (GSR) activities were evaluated as previously described (Canesi and Viarengo, 1997). Digestive glands were homogenised in 4 vol of homogenization buffer and centrifuged at 100,000 x g for 90 min at 0-4 °C. Aliquots of the supernatants were utilized for the spectophotometric determination of enzyme activities. GST activity was evaluated with CDNB (1-chloro-2,4-dinitrobenzene) as a substrate. The reaction mixture (1 ml) contained 125 mM K-phosphate buffer, pH 6.5, 1 mM CDNB, 1 mM GSH. The formation of S-2,4-dinitro phenyl glutathione conjugate was evaluated by monitoring the increase in absorbance at 340 nm. GSR activity was estimated in 1 ml of a reaction mixture containing 125 mM K-phosphate buffer, pH 7.5, 0,05 nM NADPH, 1 mM GSH. The NADPH oxidation was evaluated by monitoring the increase in absorbance at 340 nm.

Assay for total glutathione: digestive glands were homogenised in 4 vol of 1 N perchloric acid, 2 mM EDTA and then centrifuged at 16,000 x g for 20 min at 4 °C. The supernatants were neutralized with 2 M KOH, 0.3 M morpholin ethanol sulphonic acid (MOPS) and centrifuged at 1,000 x g for 10 min at 4 °C. Aliquots of the neutralized supernatant were utilized for the evaluation of the total glutathione concentration (i.e. molar sum of GSH+1/2 GSSG) as previously described (Canesi and Viarengo, 1997), utilizing the enzymatic GSSG reductase assay, in 1 ml of a reaction mixture containing 0.1 M K-phosphate buffer, 1 mM EDTA, pH 7.0, 0.2 mM NADPH, 0.06 mM dithionitrobenzoic acid and 0.12 U of GSSG reductase. Calibration was performed utilizing known

concentrations of GSSG instead of samples. Protein content was determined according to the Lowry method using bovine serum albumin (BSA) as a standard.

2.4 Lysosomal parameters

Lysosomal membrane stability was evaluated in duplicate cryostat section of 5 digestive glands according to Moore (1988) and sections (10 µm) were cut with a cryostat (Bright CM3050), flashdried by transferring them to room temperature, and then stained for N-acetyl-β-hexosaminidase activity (Moore, 1976). Digital images of the tissue sections were acquired by an Olympus BX60 microscope light microscope equipped with a scientific grade Color ViewII CCD Camera (Olympus, J). Staining intensity of lysosomes was determined at 400× magnification using the Scion Image software package (Scion Corp). Neutral lipid content was evaluated in cryostat sections of digestive glands fixed in calcium-formaldehyde (2% Ca-acetate (w/v), 10% formaldehyde (v/v)) for 15 min at 4 °C, followed by a rinsing step with de-ionised water, and incubation with 60% triethylphosphate (TEP) for 3 min (Moore, 1988). The sections were then stained with Oil Red-O (1% in 60% TEP) for 30 s, rinsed with de-ionised water, and mounted in 20% (v/v) glycerol. Lipofuscin content was determined using the Schmorl reaction on cryostat sections fixed in calcium-formaldehyde and rinsed with de-ionised water, as described for the neutral lipid assay, followed by a 5 min incubation step with 1% Fe₂Cl₃, 1% potassium ferrocyanide in a 3:1 ratio (Moore, 1988). The sections were rinsed with 1% acetic acid and mounted in 20% (v/v) glycerol. Neutral lipid and lipofuscin content were quantified by digital image analysis of stained sections, as described for the LMS assay. Results are reported as % optical densities with respect to controls.

2.5 Data analysis

The results are the mean±SD of at least 3 experiments, each conducted with 10-12 mussels per experimental group, and analyses were performed in triplicate. Statistical analysis was performed

by using the Mann-Whitney U test with significance at P \leq 0.05.

2.6 Chemicals

All reagents were of analytical grade. Bisphenol A was from Sigma (St. Louis, MO). All other reagents were purchased from Sigma (St. Louis, MO). The PCR reagents were from Fermantas (St.Leon-Rot Germany), and Promega, Sigma, or Biorad (Milan, Italy).

3. Results

3.1 Effects of BPA on gene expression

The possible effects of BPA exposure on the expression of different genes in mussel hepatopancreas were evaluated by quantitative RT-PCR, utilising the primer pairs reported in Tab. 1. Expression of the *Mytilus* ER gene sequences MeER1 and MeER2 was first evaluated in control animals by RT-PCR: both sequences were expressed in mussel hepatopancreas, with MeER1 showing a higher basal expression with respect to MeER2 (Fig. 1A). As shown in Fig. 1B, BPA induced a significant decrease in the expression of MeER1 at 15 ng/g dw (-54% with respect to controls; P \leq 0.05), whereas other concentrations were ineffective. On the other hand, a two-fold increase in the transcription of MeER2 was observed at all the concentrations tested, that was significant (P \leq 0.05) at the lowest concentration (3 ng/g). When mussels were exposed to E₂, at concentrations previously shown to affect the expression of other genes in the hepatopancreas in the same experimental conditions (Canesi et al., 2007), a small increase in the amount of both MeER1 and MeER2 transcripts was observed, that was significant for MeER1 at 1.35 ng/g (+27%; P \leq 0.05) and for MeER2 at 6,75 ng/g (+39%; P \leq 0.05), respectively (Fig. 1C).

The effects of BPA on *Mytilus* gene sequences whose transcription has been previously shown to be modulated by E_2 in mussel hepatopancreas in the same experimental conditions (Canesi et al., 2007) were also evaluated. Fig. 2 shows the results obtained for catalase, the two *Mytilus* metallothionein isoforms MT20 and MT10, and the p53-like gene. BPA induced a decrease in the transcription of catalase that was significant at 15 ng/g (-60%; P≤0.05), followed by recovery at the highest concentration. BPA induced downregulation of the MT20 gene at all the concentration tested (about -60-70%; P≤0.05); a significant decrease in the expression of MT10 was also observed at the highest concentration of 60 ng/g BPA (-40%; P≤0.05). A small but not significant decrease was seen for the transcription of the mussel p53-like gene at 15 ng/g BPA; no significant changes were observed for the lysosomal protease cathepsin L (data not shown).

3.2 Effects of BPA on antioxidant enzyme activities and glutathione content.

BPA exposure also modulated the activities of enzymes involved in maintenance of redox balance (Fig. 3). As shown in Fig. 3A, catalase specific activity showed a significant decrease in response to the two lower concentrations of BPA (-19% and -13% with respect to controls, respectively; P \leq 0.05), followed by an increase (+24%, P \leq 0.05) at the highest concentration. BPA exposure increased GST and GSR activity at lower concentrations (Fig. 3B and 3C), up +63% and +29% of control values at 15 and 3 ng/g, respectively (P \leq 0.05) and increased the total glutathione content at all the concentrations tested (about +40% with respect to controls; P \leq 0.05) (Fig. 3D).

3.3 Effects of BPA on lysosomal parameters

Fig. 4 shows the effects of mussel exposure to BPA on lysosomal membrane stability (LMS) and neutral lipid and lipofuscin accumulation, evaluated in digestive gland cryostatic tissue sections. BPA induced a concentration-dependent decrease in LMS (Fig. 4A). At the same time, increased accumulation of NL was observed, up +240% with respect to controls (P \leq 0.05) in mussels injected with the highest concentration of BPA (Fig. 4B). BPA did not affect the lysosomal content of lipofuscin, the end-product of lipid peroxidation (Fig. 4C).

No significant differences in the response to BPA were observed in males and females and in animals sampled at different times of the year (October or June).

4. Discussion

The results demonstrate that BPA exposure significantly affected gene expression and functional parameters in *Mytilus* hepatopancreas. When compared with the results obtained in mussels exposed to exogenous E_2 in the same experimental conditions (this work; Canesi et al., 2007) these data indicate that in invertebrates the xenoestrogen can induce changes in gene expression at concentrations comparable with those of the natural estrogen.

Partial sequences of the *Mytilus edulis* MeER1 and MeER2 have been recently identified that are expressed in gonads (Puinean et al., 2006). MeER1 clusters with vertebrate ERs, and the MeER1 protein sequence has the highest identity (60%) with mouse and zebrafish Estrogen Related Receptor ERR γ compared with the invertebrate *Aplysia californica* ER (37%). MeER2 clusters with the invertebrate Aplysia ER and has the highest protein sequence identity with Aplysia and Octopus ERs (70% and 66%, respectively) and, among vertebrates, with the chicken ER α (51%). Our data indicate that both MeERs are constitutively expressed in the hepatopancreas of control mussels, with MeER1 showing a higher basal expression than MeER2, and that BPA exposure induced distinct and significant changes in the level of the mRNA transcripts for both genes. In particular, a 50% decrease for MeER1 at 15 ng/g BPA and a two-fold increase for MeER2 at all the concentrations tested were observed. Smaller but significant increases in the transcription of both MeERs were observed with E_2 at concentrations that were previously shown to affect the expression of other genes in the same experimental conditions (Canesi et al., 2007). In fish, BPA has been shown to increase expression of ER α in different tissues (Seo et al., 2006; Hayashi et al., 2007). Our data demonstrate that estrogenic compounds can modulate the expression of ER-like sequences also in invertebrates.

BPA exposure downregulated the transcription of the antioxidant enzyme catalase, that is responsible for decomposition of H_2O_2 , and of the metallothionein isoform MT20; a significant decrease in the level of MT10 transcripts was also observed at the highest concentration tested. MTs play a role in protection against oxidative stress in different organisms (Vergani et al. 2005);

in *Mytilus*, the MT-20 isoform is preferentially induced by oxidant agents (such as hydrogen peroxide) and cadmium, whereas MT-10 is the constitutive isoform that preferentially responds to essential metals, such as zinc (Lemoine and Laulier, 2003; Dondero et al., 2005). The effects of BPA were distinct from those previously observed in the same experimental conditions with E_2 , that induced significant increases in the expression of both catalase and MT20 (Canesi et al., 2007). Modulation of MT expression seems to represent a significant target for xenoestrogen action in different organisms (Gerpe et al., 2000; Sogawa et al., 2001; Werner et al., 2003; Gagnè et al., 2004). BPA, like E_2 (Canesi et al., 2007), decreased the expression of *Mytilus* p53-like gene and did not affect the expression of the lysosomal protease cathepsin L (not shown).

Overall, the results demonstrate significant changes in gene transcription in mussels injected with E_2 or BPA. This route of exposure, that allows for direct administration of estrogens into the animal open circulatory system leading to distribution from the blood (hemolymph) to the tissues, including the hepatopancreas, has been largely utilised for evaluating the effects of natural and environmental estrogens in bivalve molluscs (Gagnè et al., 2001; Canesi et al., 2005, 2006, 2007; Le Curieux-Belfond et al., 2005; Gauthier-Clerc et al., 2006; Puinean and Rotchell, 2006). In Mytilus sp., exogenous E₂ added to the seawater (200-2000 ng/l) is taken up and progressively metabolised (within 7 days) to an esterified, inactive form that is stored in the tissues, this indicating that mussels can tightly regulate the concentration of free E_2 (Janer et al., 2005). In mussels exposed for 10 days to E_2 (200 ng/l) in the seawater to ensure significant estrogen accumulation in the tissues, no significant changes were observed in the expression of ER- and Vitellogenin-like sequences in the gonad (Puinean et al., 2006). The lack of effect was mainly ascribed to homeostatic mechanisms of conversion of the excess E_2 into inactive esters in these organisms (Janer et al., 2005; Janer and Porte, 2007). On the other hand, changes in Vtg expression were observed in the gonad of mussels 48 hr post-injection with 25 μ g E₂ (Puinean and Rotchell, 2006). Administration by this route probably allows to bypass the homeostatic mechanism of progressive metabolism to inactive compounds that occurs in the presence of exogenous E_2

(Puinean et al., 2006; Puinean and Rotchell, 2006). Irrespectively of the route of administration, the concentrations of E_2 and BPA utilised in the present work were much lower than those utilised by these Authors, and comparable to those found in environmental samples from marine coastal areas (1,2-53 ng/l for E_2 and 1,5-1,88 ng/l for BPA, respectively; Pojana et al., 2004). Therefore, the effects observed after 24 hr injection may be due to direct effects of the estrogens administered into the blood rather than to their large accumulation in the tissues.

BPA also significantly affected the activity of enzymes involved in maintenance of tissue redox balance. A decrease in catalase activity was observed at the two lower concentration tested, followed by an increase at the highest concentration; such a trend in enzyme activity partially reflected that observed for transcription of catalase mRNA and was opposite to that observed with E_2 (Canesi et al., 2007). BPA stimulated GSR activity at the two lower concentrations, and such an increase may compensate for the decrease in catalase activity, allowing the continuous recycling of GSSG to GSH; moreover, BPA exposure resulted in an increase in total glutathione content. BPA also stimulated the activity of the phase II enzyme GST, this suggesting that some metabolism of BPA may occur by this route in the hepatopancreas. BPA bioaccumulation and clearance have been demonstrated in bivalves, with shorter half-lives at higher temperatures (Heinonen et al., 2002).

Overall, the results indicate that BPA affects components involved in the maintenance of the redox status by downregulating the expression of antioxidant genes, at the same time upregulating the activities of enzymes involved in glutathione metabolism. As a result of this, BPA exposure did not induce oxidative stress conditions in the tissue, as indicated by the absence of changes in the lysosomal content of lipofuscin, the end-product of lipid peroxidation, or in the tissue level of malondialdehyde (not shown). On the other hand, in the same experimental conditions, E₂ mainly showed anti-oxidant effects, as indicated by upregulation of catalase and MT20 genes, increase in catalase specific activity and slight decrease in lipofuscin content (Canesi et al., 2007). The results obtained with BPA are in line with those obtained in mammalian systems, where the xenoestrogen can act both as a pro-oxidant and an antioxidant because BPA has antioxidant activities structurally

(Kadoma and Fujisawa et al., 2000), while it shows pro-oxidant activity through the estrogen receptor or its metabolites (Knaak and Sullivan, 1966; Atkinson and Roy, 1995; Obata and Kubota, 2000; Kabuto et al., 2003).

BPA also induced significant changes in digestive gland lysosomal parameters that are related to the health status of bivalves (Moore et al., 2006). The effects were similar to those previously observed with E_2 (Canesi et al., 2007), showing a dose-dependent decrease in LMS and increase accumulation of neutral lipids (NL). Increased NL accumulation in bivalve tissues indicates either increased synthesis or decreased utilisation of neutral lipids, as well as impairment of lysosomal lipases, leading to enhanced autophagy of the accumulated lipids in digestive cells often associated to lysosomal membrane destabilization (Moore et al., 2006). Overall, the results further support the hypothesis that the lysosomal system represents a significant target for estrogenic compounds in molluscan cells (Moore et al., 1978; Canesi et al., 2004a,b, 2006, 2007) like in mammalian cells (Szego et al., 1975). In bivalves, the hepatopancreas plays a central role in metabolism through intracellular digestion of food particles and in distribution of nutrients to reproductive tissues during gamete maturation (Dimitriadis et al., 2004). Therefore, the overall effects induced by BPA in this tissue may result in indirect effects on reproduction. This data addresses the importance of investigating alternative endpoints of estrogenic effects also in invertebrate species.

With regards to the possible mechanisms of action of BPA, both genomic and non genomic modes of action involving ER-like receptors, as well as receptor-independent mechanisms may participate in mediating the effects (Quesada et al., 2002; Walsh et al., 2005; Ropero et al., 2006; Watson et al., 2007). In mussel immunocytes, BPA was shown to interfere with signaling components whose activation plays a key role in mediating the effect of E₂. *In vitro*, the effects of BPA were prevented or significantly reduced by the antiestrogen Tamoxifen, suggesting the involvement of ERs (Canesi et al., 2004b). Although BPA, like E₂, activated ERK MAPKs, both *in vitro* and *in vivo*, the xenoestrogen induced dephosphorylation of p38 MAPK and of transcription factors (such as STAT5 and CREB) (Canesi et al., 2004b, 2005; Porte et al., 2006) that are activated

by E_2 (Canesi et al., 2004a, 2006). Moreover, the *in vivo* effects were observed at concentrations of BPA comparable to those of E_2 and to environmental exposure levels (Canesi et al., 2005). This data suggested that BPA could lead to negative modulation of the expression of certain genes secondary to downregulation of components of cytosolic kinase-mediated cascades. In mammalian systems, E_2 and BPA exhibit both common and divergent gene regulation involving both ER-dependent and ER-independent mechanisms (Singleton et al., 2004; Welshons et al., 2006). The results here obtained seem to support the hypothesis that also in molluscan cells the distinct effects of BPA and E_2 may be ascribed to distinct modes of action of the two compounds.

Individual xenoestrogens can differ in their potency and temporal phasing of effects compared to E_2 , thus possibly disrupting some endocrine functions when acting in combination with physiological estrogens (Watson et al., 2007). In mussels, BPA exposure did not result in significant changes in the circulating levels of free E2 at 24 hr post-injection (data not shown). However, it must be considered that in vivo estrogenic chemicals are generally more stable than E_2 and circulating xenoestrogens may not be sequestered by plasma steroid-binding proteins; therefore, unmodulated action of these compounds on target tissues and cells may occur (Ropero et al., 2006; Welshons et al., 2006). Interestingly, several of the parameters affected by BPA in mussel hepatopancreas apparently showed a non monotonic dose-response curve, with stronger or more evident effects at lower concentrations. Such an effect is commonly observed with hormones, or drugs and chemicals that act via hormonal, receptor-mediated mechanisms. Although further experiments utilising a wider range of BPA concentrations, as well as comparisons with exposure to BPA dissolved in the sea water at low environmental concentrations (ng/l), are needed to confirm these observations, the results further support the hypothesis that lower, environmentally relevant doses of BPA should be used to assess the hazard posed by estrogenic chemicals to both wildlife and humans (vom Saals and Hughes, 2005; Welshons et al., 2006).

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Figure Legends

Fig. 1 – Expression of MeER transcripts in *Mytilus galloprovincialis* hepatopancreas and effects of exposure to BPA and E_2 . A) RT-PCR of MeER1 and MeER2 in the hepatopancreas of control mussels; B) and C): Quantitative RT-PCR of MeER1 and MeER2 in BPA- and E_2 -exposed mussels. Mussels were injected with different amounts of BPA (nominal concentrations 3, 15 and 60 ng/g dw, respectively) or E_2 (1.35 or 6.75 ng/g dw, respectively) or vehicle (ethanol), and tissues sampled 24 hrs post-injection. Relative expression was calculated with respect to controls as described in Methods. Data are the mean±SD from of three experiments in triplicate. * = P≤0.05, Mann-Whitney U test.

Fig. 2 – Effects of BPA exposure on the transcription of *Mytilus* genes in the hepatopancreas: catalase, MT20, MT10, p53. Mussels were injected with different amounts of BPA (nominal concentrations 3, 15 and 60 ng/g dw, respectively) or vehicle (ethanol) as described in methods and tissues sampled 24 hrs post-injection. Gene expression was determined quantitative RT-PCR and relative expression was calculated with respect to controls as described in Methods. Data are the mean \pm SD from of three experiments in triplicate. * = P \leq 0.05, Mann-Whitney U test.

Fig. 3 – Effect of BPA exposure on enzyme activities and glutathione content in *Mytilus galloprovincialis*. A) Catalase; B) GST-GSH transferase; C) GSR-GSSG reductase; D) total glutathione (GSH+1/2 GGSG). Mussels were injected with different amounts of BPA (nominal concentrations 3, 15 and 60 ng/g dw, respectively) or vehicle (ethanol) as described in methods and tissues sampled 24 hrs post-injection. Catalase activity was measured in both the 12,000 x g supernatant and pellet as described in methods, and expressed as % µmoles H_2O_2 consumed/min/mg prot with respect to controls. Data are the mean±SD of three experiments in triplicate; * = P≤0.05, Mann-Whitney U test.

Fig. 4 – Effect of BPA exposure on lysosomal parameters in *Mytilus galloprovincialis* hepatopancreas. A) Lysosomal membrane stability (LMS); B) neutral lipid accumulation; C) lipofuscin content. Mussels were injected with different amounts of BPA (nominal concentrations 3, 15 and 60 ng/g dw, respectively) or vehicle (ethanol) as described in methods and tissues sampled 24 hrs post-injection. Data are the mean \pm SD (n=10); * = P \leq 0.05, Mann-Whitney U test.

Figure

ACCEPTED MANUSCRIPT





Figure





Tab.1 – Oligonucleotide primers used for RT-Q-PCR analysis.

Gene	Forward	Reverse	GenBank accession	References
MT10	5'-GGGCGCCGACTGTAAATGTTC-3'	5'-CACGTTGAAGGCCTGTACACC-3'	<u>AY566248</u>	Dondero et al., 2005
MT20	5'-TGTGAAAGTGGCTGCGGA-3'	5'-GTACAGCCACATCCACACGC-3'	<u>AY566247</u>	Dondero et al., 2005
p53-like	5'-CCAACTTGCTAAATTTGTTGAAGA-3'	5'TTGGTCCTCCTACACATGAC-3'	DQ158079	Dondero et al., 2006
Catalase	5'-CGACCAGAGACAACCCACC-3'	5'-GCAGTAGTATGCCTGTCCATCC-3'	<u>AY743716</u>	
CathepsinL	5'-CGCAGCTAATGTTGGCGCC-3'	5'-CTACGGCGATTGGTCCCTG-3'	<u>AY618311</u>	
18S	5'-TCGATGGTACGTGATATGCC-3'	5'-CGTTTCTCATGCTCCCTCTC-3'	<u>L33451</u>	Dondero et al., 2005
MeER1	5'-TTACGAGAAGGTGTGCGTTTT-3'	5'-TCACCATAGGAAGGATATGT-3'	<u>AB257132</u>	Puinean et al., 2006
MeER2	5'-GGAACACAAAGAAAAGAAAGGAAG-3'	5'-ACAAATGTGTTCTGGATGGTG-3'	<u>AB257133</u>	Puinean et al., 2006

GAACACAAAGAA.