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ORIGINAL ARTICLE

Amino-modified polystyrene nanoparticles affect signalling pathways of the sea urchin (*Paracentrotus lividus*) embryos

Annalisa Pinsino^a, Elisa Bergami^b, Camilla Della Torre^c, Maria Luisa Vannuccini^b, Piero Addis^d, Marco Secci^d, Kenneth A. Dawson^e, Valeria Matranga^a and Ilaria Corsi^b

^aCNR – Institute of Biomedicine and Molecular Immunology “A. Monroy”, Palermo, Italy; ^bDepartment of Physical, Earth and Environmental Sciences, University of Siena, Siena, Italy; ^cDepartment of BioScience, University of Milan, Milan, Italy; ^dDepartment of Environmental and Life Sciences, University of Cagliari, Cagliari, Italy; ^eCentre for BioNano Interactions, School of Chemistry and Chemical Biology, University College Dublin, Dublin, Ireland

ABSTRACT

Polystyrene nanoparticles have been shown to pose serious risk to marine organisms including sea urchin embryos based on their surface properties and consequently behaviour in natural sea water. The aim of this study is to investigate the toxicity pathways of amino polystyrene nanoparticles (PS-NH₂, 50 nm) in *Paracentrotus lividus* embryos in terms of development and signalling at both protein and gene levels. Two sub-lethal concentrations of 3 and 4 µg/mL of PS-NH₂ were used to expose sea urchin embryos in natural sea water (PS-NH₂ as aggregates of 143 ± 5 nm). At 24 and 48 h post-fertilisation (hpf) embryonic development was monitored and variations in the levels of key proteins involved in stress response and development (Hsp70, Hsp60, MnSOD, Phospho-p38 Mapk) as well as the modulation of target genes (*PI-Hsp70*, *PI-Hsp60*, *PI-Cytochrome b*, *PI-p38 Mapk*, *PI-Caspase 8*, *PI-Univin*) were measured. At 48 hpf various striking teratogenic effects were observed such as the occurrence of cells/masses randomly distributed, severe skeletal defects and delayed development. At 24 hpf a significant up-regulation of *PI-Hsp70*, *PI-p38 Mapk*, *PI-Univin* and *PI-Cas8* genes was found, while at 48 hpf only for *PI-Univin* was observed. Protein profile showed different patterns as a significant increase of Hsp70 and Hsp60 only after 48 hpf compared to controls. Conversely, P-p38 Mapk protein significantly increased at 24 hpf and decreased at 48 hpf. Our findings highlight that PS-NH₂ are able to disrupt sea urchin embryos development by modulating protein and gene profile providing new understandings into the signalling pathways involved.

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Introduction

Polystyrene (PS), an aromatic synthetic compound obtained by polymerisation of styrene monomers, represents one of the most widely used plastic, which has been released in the environment at an uncontrollable rate (Plastics – The Facts, 2015). PS is not readily degraded or processed by natural biological mechanisms (Andrady, 2011) but, instead, is broken up either mechanically or by photo- and thermal-oxidative degradation into plastic debris at the micrometric and nanometric size (Lambert & Wagner, 2016; Zhang et al., 2012).

Due to their small size, nanoplastics (<1 µm, as proposed by Browne et al., 2007; Hartmann et al., 2015) can be easily ingested and accumulated by organisms belonging to different trophic levels, with uncertain consequences for the health of the wildlife (Wright et al., 2013). Given the increasing evidence of plastic pollution in the ocean worldwide, their presence has become a significant environmental concern for governments, scientists and non-governmental organisations. For example, the EU Member States are committed to achieve or maintain good environmental status (GES) of their marine waters by 2020 through 11 qualitative descriptors, one of which (descriptor 10) is focused on marine litter, including plastics (Galgani et al., 2010; MSFD 2008/56/CE).

Several initiatives at scientific and industrial levels have improved the current knowledge in terms of risk assessment and sustainable productions. However, most of current information is

on macro- and microparticles or fragments ingested by marine wildlife, possibly leading to damage to the digestive system and starvation or causing entanglement (Andrady, 2011; Kühn et al., 2015; Wright et al., 2013). Smaller fragments as nanoplastics are by far less studied, despite they can originate from the weathering and/or fragmentation of larger debris (macro and micro) mainly driven by movements of the oceans (horizontally and vertically) and by physico-chemical as well as biological parameters of the receiving sea waters. Due to their smaller size, nanoplastics can be taken up and accumulated by organisms. This makes their potential biological impact an important issue, given the current concerns regarding the environmental behaviour and ecotoxicity of nano-sized particles (Koelmans et al., 2015).

Nanoparticles (NPs) are well-known to affect marine organisms causing a wide variety of biological injuries in species belonging to different trophic levels, from planktonic to benthic species, including fish (Baker et al., 2014; Blasco et al., 2015; Canesi & Corsi, 2016; Matranga & Corsi, 2012; Minetto et al., 2014). Among them, nanoplastics recently attracted the attention of the scientific community since they have been shown to cause severe damages in marine invertebrates affecting immune system, survival and reproduction (Bergami et al., 2016; Canesi et al., 2015, 2016; Della Torre et al., 2014).

The enormous diversity of life in the sea offers a rich selection of organisms, especially invertebrates, with specialised adaptations

that enable researchers to learn more about the conserved molecular signalling pathways involved in protection, robustness, resistance and plasticity towards exposure to contaminants (Bodnar, 2016). Sea urchins have served as elective model in developmental biology research for more than a century, contributing to understand not only development and differentiation, but also the basis of cellular and molecular biology as well as evolutionary biology and most recently toxicology and ecotoxicology (Lawrence, 2013). Part of the sea urchin's value as a model organism is its close genetic relationship to humans, having an estimated 23,300 genes, with representatives of nearly all vertebrate gene families (Sea Urchin Genome Sequencing Consortium, 2006). Following the publication of the sea urchin genome, a new prospect seems to be opened in science: the use of sea urchin as an alternative *proxy to human* non-mammalian model to contribute to highlight the key protective molecules that could be used in innovative applications in environmental and biomedical fields. Considering its high sensitivity to low concentrations of contaminants and its discriminatory capability, the sea urchin is considered an excellent tool to assess the toxicity of many chemical compounds and environmental stressors, especially during its embryonic life stages (Morroni et al., 2016).

With this in mind, it provides an attractive and amenable model for exploring the molecular signalling pathways engaged in NPs recognition and interaction. Sea urchin embryo has been successfully used for assessing the toxicity of metal, metal-oxides and carbon-based NPs, showing to develop countless abnormalities depending on type, exposure conditions and concentrations (Burić et al., 2015; Canesi & Corsi, 2016; Corsi et al., 2014; Larson & Hutz, 2010; Pinsino et al., 2015; Šiller et al., 2013; Torres-Duarte et al., 2016; Wu et al., 2015).

For instance, Ag and SiO₂ NPs have been reported to cause dose-dependent developmental defects such as delayed development, bodily asymmetry, shortened or irregular arms and alterations in swimming patterns in embryos of the Mediterranean sea urchin *Paracentrotus lividus* (Gambardella et al., 2015; Siller et al., 2013). Conversely, in embryos of the white sea urchin *Lytechinus pictus*, both TiO₂ and CeO₂ NPs do not elicit visible harmful effects (Fairbairn et al., 2011).

We recently showed that PS NPs having cationic (-NH₂) or anionic (-COOH) surface groups undergo different aggregation in natural sea water (NSW), thus affecting their bio-distribution and ultimate toxicity in embryos of *P. lividus* (Della Torre et al., 2014).

Cationic PS-NH₂ have been shown to be cytotoxic for mammalian cells *in vitro*, while anionic PS-COOH exhibited greater and rapid accumulation in selected internal organs (Bexiga et al., 2011; Wang et al., 2013). Specifically, authors reported that PS-NH₂ toxicity depends on cell-specific endocytic and mitochondrial injury pathways involving caspases (i.e. caspase 3, 7, 9) (Bexiga et al., 2011; Xia et al., 2006). We hypothesised a similar mechanism of toxicity in sea urchin *P. lividus* embryos, since exposure to cationic PS-NH₂ induced developmental toxicity. Our first results on sea urchin embryos exposed to PS NPs suggested that PS-NH₂ (3 µg/mL nominal concentration) do not accumulate within cells/tissues during early development but cause teratogenesis, and stimulate an up-regulation of the *caspase 8* gene expression (24 h post fertilisation, hpf) (Della Torre et al., 2014). On the contrary, PS-COOH massively accumulates in the gut without producing any toxic effect to embryos. Studies in other marine invertebrate species confirm a significant accumulation of anionic PS NPs in the gut of rotifers (Snell & Hicks, 2011) and in bivalves (Ward & Kach, 2009; Wegner et al., 2012) as well as low toxicity.

Therefore, molecular mechanisms underlying the observed developmental disorder remain largely unknown and need to be

investigated looking at signalling pathways elicited by PS-NH₂ during embryonic development. Since sea urchin embryos showed a discrete resistance and plasticity towards a persistent and direct interaction with PS-NH₂, we hypothesised that these particles are able to elicit a signalling regulating the balance between cell survival and death.

In order to test such hypothesis, in the present study, we investigated the effects of PS-NH₂ (50 nm) on sea urchin *P. lividus* embryonic development coupled with the variations in the levels of key proteins triggering stress response and development (Hsp70, Hsp60, MnSOD, P-p38 Mapk) as well as their gene expression profile (*Pl-Hsp70*, *Pl-Hsp60*, *Pl-Cytochrome b*, *Pl-p38 Mapk*, *Pl-Caspase 8*, *Pl-Univin*) under two PS-NH₂ sub-lethal concentrations of 3 and 4 µg/mL, previously shown to be able to affect embryonic development.

Materials and methods

NPs source and characterisation

Unlabelled PS-NH₂ (nominal size of 50 nm, positive surface charge) were purchased from Bangs Laboratories Inc (9025 Technology Dr, Fishers, IN 46038-2886 USA) and previously characterised both in Milli Q water (MilliQ, Merck KGaA, Darmstadt, Germany) and NSW by transmission electron microscopy (TEM) and dynamic light scattering (DLS), as described by Della Torre et al. (2014). TEM was applied for primary particle diameter identification of PS NPs (Philips Morgagni 268D electronics, at 80 KV and equipped with a MegaView II CCD camera). Dynamic Light Scattering (Malvern instruments) was used to determine size distribution parameters (Z-average and polydispersity index, PDI) (Zetasizer Nano Series software, version 7.02, Particular Sciences, UK). Measurements have been performed in triplicate, each containing 11 runs of 10 s.

PS-NH₂ suspensions were prepared in 0.45 µm filtered NSW (T 18 ± 1 °C, salinity 38‰, pH 8.3) and it was quickly vortexed prior to use. PS-NH₂ concentrations (3 and 4 µg/mL) used for embryotoxicity tests were chosen based on our previous findings on embryos of *P. lividus* (Della Torre et al., 2014), in which we observed several abnormalities at these low exposure concentrations (EC₅₀ values of 3.85 and 2.61 µg/mL at 24 and 48 hpf, respectively) and also an apoptotic pathway at 3 µg/mL.

The NSW, obtained from a marine area located in South Sardinia (Italy), was previously 0.45 µm filtered and treated with Ultra Violet light before being used for experiments.

All exposure studies were carried out at the sea urchin hatchery facilities of the Department of Environmental and Life Science of the University of Cagliari (Italy). The following parameters of the NSW were maintained during experiments (0.45 µm filtration, 18 °C, 38‰, pH 8) and DLS analysis was performed in order to confirm that the aggregation state of PS-NH₂ was similar to that obtained in our previous study (Della Torre et al., 2014).

Sea urchin embryo culture and PS-NH₂ exposure

Paracentrotus lividus broodstock was induced to spawn by injection of 1 M KCl (40 µL per g of body weight) into the coelom via the peristomial membrane. The released eggs were fertilised by addition of 10 mL of diluted sperm (Carboni et al., 2013). Sea urchin embryos (600 embryos/mL) were exposed to 3 and 4 µg/mL PS-NH₂ suspensions soon after the fertilisation following the same procedure used in our previous experiments (Della Torre et al., 2014). Control embryos were exposed to NSW.

Table 1. List of primer sequences used for gene expression profile in sea urchin embryos investigated through real-time qPCR, including amplicon size (bp), efficiency (%), accession number and annealing temperature (°C).

Gene	Forward primer 5'-3'	Reverse primer 5'-3'	Amplicon size (bp)	Efficiency (%)	Acc number	Annealing T (°C)
<i>PI-Hsp70</i>	CCAGTCGAAAGGGCAATCCT	CAACGTCGACGAGGAGAACA	244	100.5	X61379	59
<i>PI-Hsp60</i>	GACGGAACCACAACAGCAAC	CCTGGGCGATTCTTCTGGT	181	105.4	AJ249625	59
<i>PI-p38 Mapk</i>	GTGATCAGCTTGCTTGACTG	GATAGATGAGGAACTGGACGTG	144	105.4	Marrone et al., 2012	58
<i>PI-14-3-3α</i>	GATCCCCTCTGCCAAAGACC	CCTGATAGGGTCTGTCCGGT	187	101.8	AJ493680	61
<i>PI-Univin</i>	GATCATCGCTCCGATGGGTT	GCTCGGCTATCGATGGAGTT	129	94.5	AJ302364	59
<i>PI-Caspase 8</i>	TCCTGCCCAAAGTCTACTG	CCTCGCAGTTGGGTCTTTCT	232	99.0	EU078681	59
<i>PI-Cytb</i>	GGGATACGTATTAGTCTGAGG	CGAGTTAGGGTGGCATTGTC	144	108.8	Marrone et al., 2012	58
<i>PI-18S</i>	GAATGTCTGCCCTATCAACTTTCG	TTGGATGTGGTAGCCGTTTCTC	119	104.0	Ragusa et al., 2013	60

At key time-points of the development as 24 hpf (gastrula stage) and 48 hpf (pluteus stage), embryo cultures (40 mL; about 24000 embryos) were collected by centrifugation (2000 g for 8 min), stored in liquid nitrogen and further at -80°C for protein and gene expression analyses. Embryos were monitored to assess the occurrence and timing of several morphological events: endoderm, ectoderm and mesoderm (germ layers) development and differentiation. Aliquots of 5 mL of embryos at both 24 hpf and 48 hpf were fixed in 1:1 lugol:ethanol 70%, inspected by optical microscopy (Olympus BX51) and photographed by a DP50 digital camera.

SDS-PAGE, immunoblotting

Lyophilised pellets were Dounce-homogenized on ice in about 250 μL of lysis buffer [20 mM Tris, 150 mM NaCl, 1 mM EDTA, 1 mM EGTA, 1% Triton], supplemented with a protease inhibitor cocktail (Roche) and phosphatase inhibitor cocktail (SIGMA). Total cell lysate (30 μg) per each cell extract was run on a 4–15% Mini-PROTEAN TGX precast polyacrylamide gels (Bio-Rad) and transferred to nitrocellulose membranes (GE Healthcare Life Sciences, Amersham UK). After blocking, replicate membranes were incubated overnight at 4°C with either one of the following primary antibodies: (i) Heat Shock Protein 70 (SIGMA, Cat N. H-5147) 1:1000; (ii) Heat Shock Protein 60 (SIGMA, Cat N. H-3524) 1:250; (iii) Phospho-p38 MAP Kinase (Thr180/Tyr182) (Cell Signalling, 9211) 1:250; (iv) Manganese superoxide dismutase (Enzo Life Sciences, ADI-SOD-110) 1:500. Membranes were washed three times with PBS-Tween 20 prior to incubation with a fluorescein-labelled secondary anti-mouse and/or -rabbit antibody (LI-COR Biosciences). Protein levels (Hsp70, Hsp60, MnSOD, P-p38 Mapk) were normalised using α -tubulin as internal control. Results were reported in arbitrary units \pm SE obtained from the volumetric analysis of the normalised bands (sum of the two independent experiments).

RNA extraction, cDNA synthesis, real-time quantitative PCR

Total RNA was extracted from pellets of control and exposed embryos (3 and 4 $\mu\text{g}/\text{mL}$ PS-NH₂) from specimens obtained from two independent experiments. Total RNAs from each specimen (250 ng) were reverse transcribed using I-script cDNA Reverse Transcription Kit (Biorad), according to manufacturer's instructions. The real-time qPCRs were run using a StratageneMx3000P thermal cycler, as previously described (Della Torre et al., 2014). The expression of selected genes, which are known to be involved in the development and stress-response (*PI-Hsp70*, *PI-Hsp60*, *PI-Cytb*, *PI-p38 Mapk*, *PI-Caspase 8*, *PI-Univin*) was investigated. The *PI-18S* was used as internal endogenous reference gene. *PI-p38 Mapk*, *PI-Cytochrome b* and *PI-18S* primer sequences were already published (Marrone et al., 2012; Ragusa et al., 2013), whereas others were designed using NCBI Primer-BLAST (<http://www.ncbi.nlm.nih.gov/tools/primer-blast/>). Primer sequences used for real-time qPCRs

and the amplicon lengths (ranging from 129 to 244 bp) are summarised in Table 1. Data were analysed by the $\Delta\Delta\text{Ct}$ method (Pfaffl 2001). The levels of expression of the genes were considered significantly different when they were two-fold higher or lower than those measured in controls ($0.5 < \text{not significant} < 2$).

Statistical analysis

All data were analysed by the one-way analysis of variance (one-way ANOVA) compared with the respective control group, followed by the multiple comparison test of Tukey's, using the OriginPro 7.5 statistical programme (the level of significance was set as $p \leq 0.05$).

Results and discussion

PS-NH₂ characterisation

PS-NH₂ nominal diameter of 50 nm in MilliQ was confirmed by TEM. Secondary characterisation of PS-NH₂ by DLS revealed the formation of a number of small aggregates in NSW (Z-average of 143 ± 5 nm SD and PDI of 0.413), slightly higher with respect to the average aggregate size reported in our previous study for the same particle (89 ± 2 nm, Della Torre et al., 2014). The limited increase in the average size of PS-NH₂ aggregates could be related to the different NSW used (see Materials and Methods section). However, no differences in phenotypes of sea urchin embryos at 24 and 48 hpf previously exposed (Della Torre et al., 2014) and those obtained in the present study were observed.

PS-NH₂ effects on sea urchin embryonic development

At 24 hpf, at both concentrations used, embryos displayed abnormal proliferation of ectodermal membrane despite they maintained regular time schedule and proper sites of spicule elongation (Not shown). Such phenotypes were in agreement with our previous findings in which the thickening and abnormal proliferation was observed at this developmental stage (Della Torre et al., 2014).

At the pluteus stage (48 hpf), PS-NH₂ exposed embryos displayed a high number of abnormalities, defined as delayed in the developmental schedule or impairment of the correct differentiation of the derivatives germ layers (Figure 1).

Specifically, more than 70% of embryos exposed to 3 $\mu\text{g}/\text{mL}$ PS-NH₂ belonged to the delayed category, showing a correct morphology with a pyramidal shape although they were late in skeleton elongation (see black and pink arrow heads in Figure 1(B-B')) compared to controls (Figure 1(A-A')).

An increased incidence of potential teratogenic effects was observed at 4 $\mu\text{g}/\text{mL}$ PS-NH₂. At this concentration, 90% of embryos exhibited poorly developed skeleton rods,

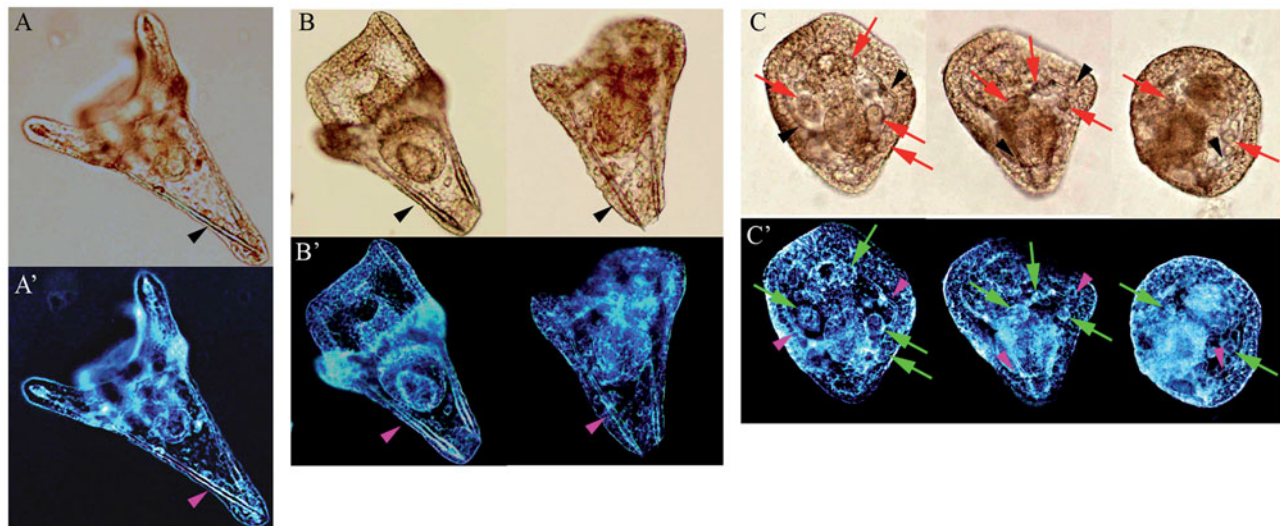


Figure 1. PS-NH₂-exposed embryos under optical microscopy. Pictures of representative sea urchin *P. lividus* embryos exposed to PS-NH₂ (3 and 4 µg/mL) are shown. Control (A–A') and 3 and 4 µg/mL PS-NH₂-exposed (B–B' and C–C') embryos at pluteus stage (48 hpf). PS-NH₂ exposure produced teratogenic effects in the sea urchin embryos. Embryos exposed to 3 µg/mL PS-NH₂ belonged to the delayed category, showing a correct morphology with a pyramidal shape although they were late in skeleton elongation (black and pink arrows). Embryos exposed to 4 µg/mL PS-NH₂ revealed poorly developed skeleton rods, underdeveloped/poorly developed arms, poor symmetry and atypical cells/masses randomly distributed, which appeared bigger than expected (red and green arrows). Arrows indicate skeleton (black and pink arrows) and atypical cells/masses (red and green arrows). (For the interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article).

underdeveloped/poorly developed arms, poor symmetry and a typical cells/masses randomly distributed, which appeared bigger than expected (see red and green arrows Figure 1(C–C')).

Similar skeleton-defective embryos have been previously reported for *P. lividus* exposed to other types of NPs suggesting that such abnormalities in embryo phenotype could be used as sensitive target upon NP exposure (Gambardella et al., 2013, 2015). A competition with calcium uptake, internalisation and consequently with bio-mineral formation during development could be responsible for such abnormalities as previously postulated in manganese-exposed embryos (Pinsino et al., 2011a).

Besides, the formation of typical cells/masses as shown upon PS-NH₂ exposure is in agreement with results previously obtained by Magesky and Pelletier (2015) in *Strongylocentrotus droebachiensis* embryos exposed to Ag NPs.

Due to their random distribution, it can be hypothesised that, in 4 µg/mL PS-NH₂ exposed embryos, these big cells/masses could be putatively intoxicated blastocoelar cells or other non-skeletogenic mesodermal cells appearing malformed as though they were vacuolated or swelled. Normal blastocoelar cells are embryonic phagocytes with a fusiform aspect residing in the blastocoel, particularly concentrated around the gut (Tamboline & Burke, 1992). As phagocytosis is an early line of defence against host invasion, and is known mechanism of NP internalisation (Petros & De Simone, 2010), blastocoelar cells could represent a primary cellular target of PS-NH₂. However, our results do not rule out the possibility that other embryonic cell types are also able to internalise PS-NH₂ involving phagocytosis or other cellular pathways.

On the other hand, PS-NH₂ could interfere indirectly with the normal embryonic development altering the biological culture media in which the embryo grows. For example, NP agglomerates/aggregates surrounding embryonic tissues could be able to interact with the cell membranes, disrupting the membrane functions (i.e. ion transport or signal transduction) (Fleischer & Payne, 2012; Rossi et al., 2014; Salvati et al., 2011) and inducing teratogenic effects (Bexiga et al., 2011; Canesi et al., 2015; Della Torre

et al., 2014; Wang et al., 2013). Further studies are needed to clarify at least one of these hypotheses.

During embryonic development, cells proliferate rapidly and differentiate to form specialised cells, most of which enter and stay into the form of G₀ phase. These cells are able to restart proliferation when required to replace cells after injury or cell death, activating alternative signalling pathways as a defence strategy for embryonic survival (Hamdoun & Epel, 2007).

In the sea urchin embryo, an integrated network of genes, proteins and pathways defends itself against chemical agents (Defensome). This network either buffers stress or produces alternative developmental phenotypes (Hamdoun & Epel, 2007).

Since PS-NH₂ exposure up to 4 µg/mL significantly interfered with the sea urchin embryonic development, producing teratogenic but not lethal effects, it can be postulated that PS-NH₂ are able to elicit signalling pathways regulating the balance between embryonic cell survival and death.

Molecular machinery engaged in PS-NH₂ exposure

To get a deeper insight into the molecular mechanisms underlying the effects of PS-NH₂ on sea urchin embryonic development, variations in the levels of key proteins triggering stress response and development (Hsp70, Hsp60, MnSOD, P-p38 Mapk) as well as in the expression of target genes (*PI-Hsp70*, *PI-Hsp60*, *PI-Cytb*, *PI-p38 Mapk*, *PI-Caspase 8*, *PI-Univin*) were analysed.

Proteins and genes were chosen taking into consideration: (i) the protein functions; (ii) the cellular localizations; (iii) the processes in which proteins participate during embryonic development; (iv) the biological and molecular results obtained from studies performed in the sea urchin embryos exposed to several chemical and physical stressors, including NPs.

For example, the stress proteins belonging to the Hsp70 and Hsp60 family are a collective of specialised proteins, molecular chaperones, which bind non-native states of other proteins and assist them to reach a functional conformation (Bukau & Horwich, 1998). MnSOD is a nuclear-encoded and mitochondria-matrix-

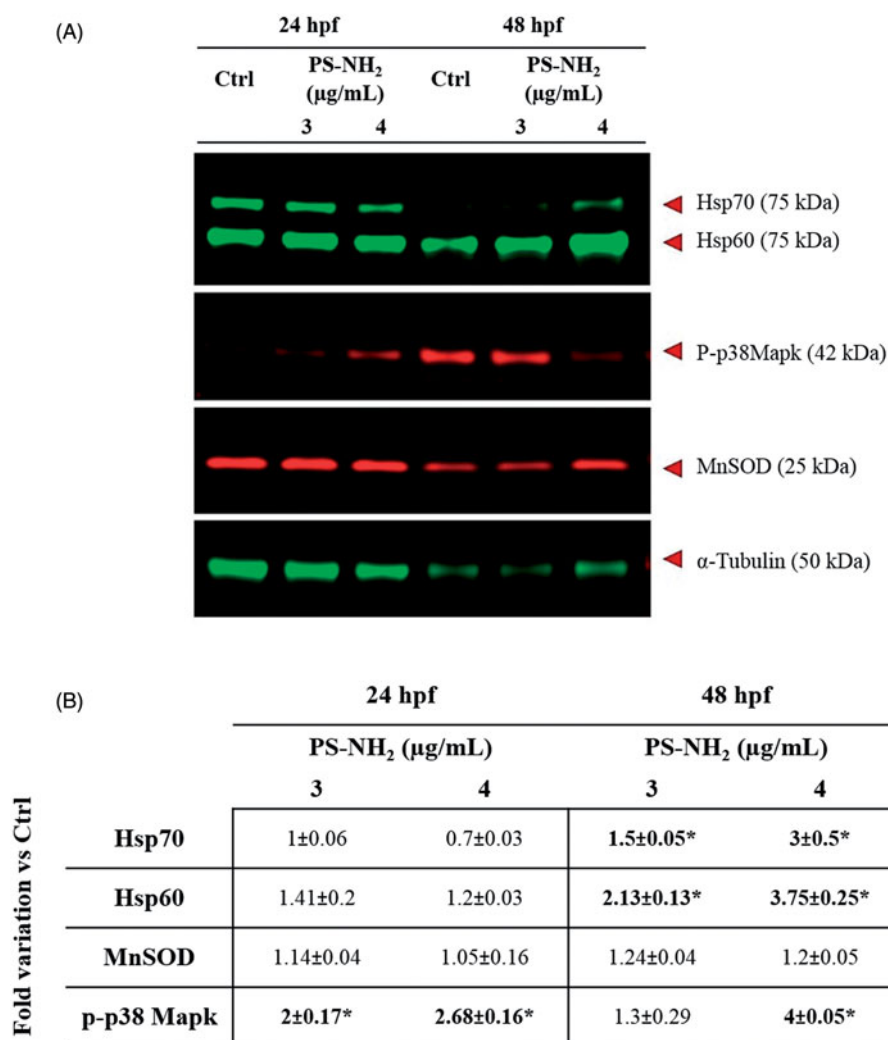


Figure 2. Levels of key proteins triggering stress response and development after PS-NH₂ exposure. (A) Representative image of the Hsp70, Hsp60, P-p38 MAPK and MnSOD protein levels of control and PS-NH₂ exposed embryos (3 and 4 µg/mL) at 24 and 48 hpf evaluated by immunoblotting. (B) Table showing fold variation (as mean ± SE) of the proteins by increasing PS-NH₂ concentration from 3 to 4 µg/mL, compared to control. Data are representative of two independent experiments after normalisation with α-tubulin levels (assumed constant). *Significant difference with respect to controls.

localized oxidation-reduction (redox) enzyme that regulates cellular redox homeostasis (Sarsour et al., 2014), while the Cytb is one of the major enzymes responsible for mitochondrial superoxide production (Zhu et al., 1999). A number of published studies demonstrated that, in the sea urchin embryos, these molecules are involved in the activity of intracellular signalling molecules, determining stress resistance, immune resistance and apoptosis (Bonaventura et al., 2011; Korkina et al., 2000; Marrone et al., 2012; Pinsino et al., 2010, 2011b; Roccheri et al., 2004). However, to our knowledge, only our previous study aimed to analyse gene expression profile upon PS NPs exposure in sea urchin embryo (Della Torre et al., 2014). Here, we extended this aim by focusing on key components of the signalling regulating growth and survival in sea urchin embryos at 24 and 48 hpf using sub-lethal concentrations of the PS-NH₂ (3 and 4 µg/mL).

Specifically, after 24 hpf, PS-NH₂ exposure (3 µg/mL concentration) was not effective in inhibiting or enhancing the protein levels of Hsp70, Hsp60 and MnSOD, respectively (see Figure 2(B)). At the same time, PS-NH₂ exposed embryos displayed a two-fold increase in the P-p38 Mapk protein levels with respect to controls ($p_{\text{value}} = .00437$). PS-NH₂ exposed embryos showed levels of expression of the *PI-Hsp70* gene 3.9-fold higher than those measured in the controls (assumed as 1), while the transcripts of *PI-Hsp60* and *PI-Cytb*

did not show a significant difference (Figure 3(A)). In addition, an increase in the levels of expression of the *PI-Univin* and *PI-Cas8* genes (2- and 2.8-fold) was also found (Figure 3(A)).

At 24 hpf, embryos exposed to 4 µg/mL nominal concentration showed significantly higher levels of the *PI-Hsp70*, *PI-p38 Mapk*, *PI-Univin* and *PI-Cas8* gene expression compared to controls (3.3, 3.5, 2.2, 2.8 and 2.3 fold, respectively), while the transcripts of *PI-Hsp60* and *PI-Cytb* did not show any change (Figure 2(B)). In accordance, no significant differences in the levels of the Hsp70, Hsp60 and MnSOD proteins were observed in PS-NH₂-exposed embryos. Instead, embryos displayed an increase in the P-p38 Mapk protein levels higher than those measured in the 3 µg/mL PS-NH₂-exposed embryos (2.7-fold) ($p_{\text{value}} = 0.00047$) (Figure 2(B)).

In summary, at 24 hpf embryos exposed to 3 and 4 µg/mL show: (i) similar levels of expression of the *PI-Hsp70* gene (from 3.93- to 3.33-fold); (ii) levels of expression of *PI-p38 Mapk* and *PI-Univin* genes increased with increasing concentration (from 1.53- to 3.58-fold and from 1.98 to 2.22); (iii) levels of expression of *PI-Cas8* unchanged; (iv) any modulation of Hsp70/Hsp60/MnSOD proteins and (v) a dose-dependent increase of p38 Mapk protein phosphorylation.

Our findings indicate that at 24 hpf, PS-NH₂ were not effective in modulating proteins having stress-related function (Hsp70/

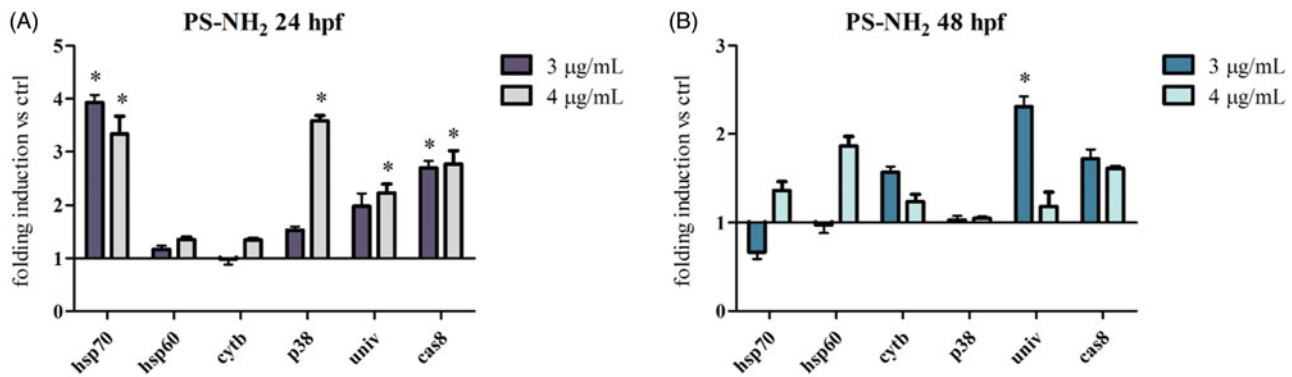


Figure 3. Misregulation of gene expression in response to PS-NH₂ exposure. Levels of expression of *PI-Hsp70*, *PI-Hsp60*, *PI-Cytochrome b*, *PI-p38 Mapk*, *PI-Caspase 8*, *PI-Univin* genes analysed by comparative real-time qPCR with total RNA isolated from control and exposed embryos at (A) 24 hpf and (B) 48 hpf. Results are expressed as fold increase compared to controls assumed as 1, using the endogenous gene *PI-18S* for normalisation. Each bar represents the mean of two independent experiments \pm SE. *Significant difference respect to controls.

Hsp60/MnSOD) in agreement with our previous observations with other types of external stimuli (i.e. metals, radiations) (Bonaventura et al., 2011; Pinsino et al., 2010). On the contrary, a significant increase of P-p38 Mapk protein was observed. The p38 Mapk is a member of a well-studied family of proteins that plays an essential role in the signalling events associated to survival response, apoptosis and autophagy, but it is also known as key regulator of both embryonic development and cancer progression (Bradham & McClay, 2006; Pinsino et al., 2014).

These results were consistent with report describing manganese effects on p38 Mapk phosphorylation during sea urchin embryonic development, in which we emphasised the role played by p38 Mapk in embryo skeleton development (Pinsino et al., 2014).

The up-regulation of Univin observed in sea urchin embryos upon PS-NH₂ exposure highlighted skeletal development as potential target of toxicity, supported also by the evident skeletal defects observed in embryos exposed to 4 µg/mL. Univin is a member of the transforming growth factor beta (TGF-beta) protein superfamily, known to be one of the promoter of the skeletal growth in the sea urchin embryos (Zito et al., 2003). Specifically, Univin is a relevant Alk4/5/7 ligand for anterior skeletal patterning during gastrulation (Piacentino et al., 2015).

Furthermore, Caspase 8 is known to be implicated in the apoptosis regulation but also required for the development of specific embryonic cell types (Fu et al., 2000; Kruidering & Evan, 2000). Taken together, our results suggest that at 24 hpf, PS-NH₂ activate protein-coding genes, involved both in embryos survival and development.

Concerning effects observed at 48 hpf, embryos exposed to 3 and 4 µg/mL PS-NH₂ displayed a concentration-dependent increase in the levels of the Hsp60 protein compared to controls (2-fold and 3.8-fold, respectively) ($p_{\text{value}}=0.00283$). A slight increase in *PI-Hsp60* gene expression (1.9-fold) only at the highest concentration was observed (Figure 2(A) and 3(B)). This result suggests that the stress-induced Hsp60 could be regulated either at the transcriptional level by activation of the heat shock transcription factors (HSFs) or at the translational level, irrespectively of the HSF transcriptional activity. Gene expression studies in cancer revealed an interruption between transcription and translation of Hsp60. For instance, the increased expression of the *Hsp60* gene is observed in different types of human cancers (ovarian cancer, pancreatic cancer, large bowel carcinoma, etc.) due to the direct regulation of *Hsp60* expression by *c-Myc* (Tsai et al., 2008).

The increase of the *Hsp60* gene, responsible for the cell transformation could be linked to the presence of atypical cells/masses randomly distributed in embryos exposed to 4 µg/mL PS-NH₂.

On the other hand, the increase in the levels of the Hsp60 protein suggests the activation of a cell survival programme focused on stabilisation of mitochondrial proteins. This finding is not surprising, given that at this stage of development we also found a slight increase in the expression levels of the *PI-Caspase 8* gene (1.8 and 1.7 fold) (Figure 3(B)). Caspase 8 would serve to counteract tumorigenesis and ensure survival. Furthermore, a concentration-dependent increase in the levels of the Hsp70 protein was observed in embryos exposed to 3 and 4 µg/mL PS-NH₂ when compared to controls (1.5- and 3-fold, respectively) ($p_{\text{value}}=0.03327$). On the contrary, no appreciable variation in the expression levels of the *PI-Hsp70* gene was found (Figure 3(B)).

These findings are consistent with the notion that, under conditions of prolonged stress, HSF (the transcription factor that regulates the expression of all heat shock genes) is inactivated by negative feedback mechanisms (Voellmy, 2004). Specifically, increased levels of the Hsp70 protein cause the inactivation of own transcription factor.

In addition, in line with the results obtained at 24 hpf, PS-NH₂ exposure was not effective in inhibiting or enhancing the protein levels of MnSOD and the transcripts of *PI-Cytb*, suggesting that PS-NH₂ exposure do not provoke oxidative stress during embryonic development.

Differently, the P-p38 Mapk protein levels were found significantly decreased at the highest concentration used (4-fold) compared to controls (Figure 2(A,B)). A recent study showed that the p38 Mapk inhibition markedly reduces the tumorigenicity by modulating angiogenesis, cell motility and invasion (Marengo et al., 2013). As reported for *PI-Hsp70* gene, also in this case the levels of expression of the *PI-p38 Mapk* gene were found comparable to controls. Finally, exposed embryos showed levels of expression of the *PI-Univin* gene significantly different from those measured in controls only at the lowest concentration (Figure 3(B)).

In summary, overall results highlighted an increase in embryotoxicity towards PS-NH₂ exposure at pluteus stage (48 hpf) at relatively low concentrations with the involvement of molecular targets such as P-p38 Mapk, Hsp60 and Hsp70 proteins, *PI-Hsp60* and *PI-Univin* genes.

Conclusions

Overall, results highlighted an astounding level of developmental plasticity of the sea urchin embryos exposed to PS-NH₂, established by the ability to try to maintain the underlying developmental pathways and to activate alternative signalling

pathways as a defence strategy for survival to these nanoplastics.

Sea urchin embryos plasticity to environmental changes and emerging contaminants should not be undervalued. The importance of our study on sea urchin embryos is therefore 2-fold as it provides a powerful model for nano-toxicity/nano-safety investigations as a proxy to humans confirming themselves as elective model for ecotoxicological studies.

Our study confirmed the concern related to the presence of nanoplastics in the marine environment. PS-based NPs are able to elicit several developmental defects, despite the potent cellular defence machinery against environmental challenges present in embryos. Our findings stress the need to thoroughly characterising the occurrence and environmental fate of nanoplastics in marine ecosystems, and to assess the interactions with marine wildlife, focusing on misregulation of key signalling pathways involved in the onset of toxicity. The information thus obtained can be useful for defining predictive investigative tools for the evaluation of the biological risk related to these emerging environmental pollutants.

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Disclosure statement

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of this article.

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