Supplementary Information

Functionalized polystyrene nanoplastics induced energy homeostasis

imbalance and immunomodulation dysfunction of marine clams

(Meretrix meretrix) at environmentally relevant concentrations

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Environmentally relevant concentrations of MPs and NPs

The exposure concentration of NPs $(0.02-2.0 \text{ mg } \text{L}^{-1})$ in the present study was selected according to the reported and predicted environmentally relevant concentrations of MPs and NPs. As shown in Table S2, the reported abundance of MPs in seawater ranged $7.6 \times 10^{-12} - 1.6 \times 10^4$ mg L⁻¹. Weathering or ageing of a single MP particle can yield millions to billions of NPs.¹ Our previous study reported that the mass yield of PS NPs (~75.2 nm) from photodegradation (a common ageing process in marine ecosystem) of PS MPs (~38.6 μ m) was 7.03 ± 0.37% (w%).² Accordingly, the estimated concentration of PS NPs in marine environment could be up to 5.3×10⁻¹⁰- 1.1×10^3 mg L⁻¹. In addition,³ predicted that the environmental concentration of NPs (50 nm) was $1.0 \times 10^{-9} - 1.5 \times 10^{-2}$ mg L⁻¹ using a theoretical 3D fragmentation model. The highest predicted concentration of 0.015 mg L⁻¹ was at the same magnitude order of the lowest exposure concentration of 0.02 mg L⁻¹ used in this study. Moreover, the concentrations of NPs used in previous toxicological investigations ranged 0.1–100 mg L^{-1} (Table S1). Therefore, to be environmentally relevant and comparable with previous studies, we selected 0.02, 0.2, and 2 mg L⁻¹ as the exposure concentrations of two NPs in the present study.

Determination of the clam ingestion rate and oxygen consumption rate

Ingestion rate, an indicator of feeding activity of clam,⁴ is defined as the quantity of microalgae ingested by per unit body weight of clams at given time.⁵ In order to determine the ingestion rate, one clam was randomly selected from each beaker after 7 days of exposure and maintained in a 500 mL glass beaker containing 400 mL filtered seawater. The clam was fed with the 2×10^5 cells mL⁻¹ mixed algae of *C. meülleri* and

I. zhanjiangensis (1: 1, v/v). After 30 minutes of ingestion, 5 mL water sample was collected from the beak using pipette to determine the number of residual algae cells using a hemocytometer. The ingestion rate was calculated according to the following equation:

Ingestion rate =
$$V \times (C_0 - C_T) / (W \times T)$$
 (1)

where V (mL) is the volume of seawater, W (mg) is the dry weight of clam soft tissue, T (h) is the ingestion time of 30 minutes, C_0 and C_T (cells mL⁻¹) is the algae density at initial and T time during the ingestion, respectively.

For measuring the oxygen consumption rate, an indicator of metabolic rate,⁴ one clam was randomly selected from each beaker and was placed into a wild-mouth bottle with filtered full aeration seawater (dissolved oxygen $8.53 \pm 0.05 \text{ mg L}^{-1}$). After 2 hours of cultivation at 17 °C in an illumination incubator (GXZ- 500C-LED, Ningbo, China), the content of dissolved oxygen in the seawater was measured by an oximeter (YSI-5000, Yellow Spring, Ohio, USA). Then the OCR was calculated as the following equation:

Oxygen consumption rate =
$$(D_0 - D_T) / (W \times T)$$
 (2)

where D_0 and D_T (mg) is the oxygen content at initial and T time; T (h) is the oxygen consumption time; W (mg) is the dry weight of calm soft tissue at time T.

	Organisms	NP type	NP size (nm)	Exposure concentration (mg L ⁻¹)	Reference	
Ph	Blue-green alga (Microcystis aeruginosa)	PS-NH ₂	200	0.5-7	6	
iytoj	Microalgae	PS	100, 500		-	
olan	(Scenedesmus obliquus)	PS-NH ₂	100	1-100	1	
kton	Cyanobacteria	DC	100	5	0	
	(Synechococcus elongates)	13	100	5	8	
	Sea urchin embryo	PS-NH ₂	50	1-50	Q	
	(Paracentrotus lividus)	PS-COOH	40	1 50)	
	Rotifer	PS	50 500	10	10	
Zooplan	(Brachionus koreanus)	10	50,500	10	10	
	Daphnia magna	PS	100	1-75	11	
kton	Daphnia magna	PS	100-120	1-400	12	
	Brine shrimp	PS-NH ₂	50	0.1-10	13	
	(Artemia franciscana)					
	Daphnia magna	PS	20	1, 50		
	Blue mussel	PS	30	100-300	15	
	(Mytilus edulis)					
	Mussels	DC MIL	50	1 50	16	
	Mytilus galloprovincialis	PS-NH ₂	50	1-50	16	
	(neamocytes)					
	(Pacton maximus)	PS	25, 250	0.015	17	
ω	Pacific ovsters	PS-NH-				
ival	(Crassostrea gigas)	15-1112	100	0.1-100	18	
ve	(gametes)	PS-COOH	100	0.1 100	10	
	Pacific ovsters	PS	50, 500			
	(Crassostrea gigas)	PS-NH ₂	50	0.1-25	19	
	(gametes)	PS-COOH	50			
	Blood clams					
	(Tegillarca granosa)	PS	500	0.26	20	
	C. fluminea	PS	80	0.1-5	21	
	Zebrafish	PS	25	10	22	
	Zebrafish	DC	2 0 - 200	100	20	
Fis	(Danio rerio)	PS	20-500	100	23	
÷	Yellow croaker	DC	100	5.50×10 ⁻¹²	24	
	(Larimichthys crocea)	٢٥	100	-5.50×10-7	2 4	

Table S1. The summarized exposure concentrations of NPs in previous studies.

Table S2. Reported abundance of MPs in global seawa	iter.
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Region	Reported concentration	Size (µm)	Normalized concentration $(mg \ L^{\text{-1}})^{\alpha}$	Reference
North Pacific Central Gyre	64-3×10 ⁴ g km ⁻²	330–5000	4×10 ⁻⁴ -2.0×10 ⁻¹	25
Australian vessels	9×10 ³ -1×10 ⁵ particles km ⁻²	330–5000	1.1×10 ⁻⁹ -4.7×10 ⁻²	26
South Pacific subtropical gyre	71–732 g km ⁻²	330–5000	5.0×10 ⁻⁴ -4.9×10 ⁻³	27
Kuril–Kamchatka Trencharea	60–2×10 ³ particles m ⁻²	330-1000	7.6×10 ⁻¹² -7.3×10 ⁻⁶	28
Yangtze Estuary	9.8×10 ⁴ -2.6×10 ⁵ particles L ⁻¹	330–5000	1.9×10 ⁻⁷ -1.8×10 ¹	29
Southeast coast of India	10–30 particles L ⁻¹	500-1000	6.9×10 ⁻¹ -1.6×10 ¹	30
Gulf of Mannar	6-223.6 particles L ⁻¹	1000–5000	3.3-1.6×10 ⁴	31
Hangzhou Bay	100–500 particles L ⁻¹	330–5000	1.9×10 ⁻⁹ -3×10 ⁻⁴	32
Marmara Sea	3–124 particles L ⁻¹	50-5000	2.1×10 ⁻⁵ -8.7×10 ³	33
South China Sea	6.6–36.6 particles L ⁻¹	330-5000	1.0×10 ⁻⁴ -2.6×10 ³	34

 $\overline{\alpha}$ Normalized concentration (mg L⁻¹) was calculated using the spherical volumes and density (1.04 g cm⁻³) of polystyrene beads.³

Reaction pattern ^{α}	Histopathological alteration	Weight (w) ^{α}
Tubule alterations	Necrosis (nes)	3
	Epithelial cell hypertrophy (ech)	2
	Widening of the tubular lumen (wtl)	2
	Epithelial cell exfoliation (ece)	2
Intertubular tissue changes	Fibrosis (fis)	2
	Haemocytes infiltrate (hai)	1

Table S3. Histopathological alterations in the digestive gland of clams and their weight values.

^a Reaction pattern and weight values were classified according to previous studies.³⁵⁻³⁸ The alterations were classified into three weight (w) values according to the biological significance of the lesion, represented the degree in which the lesion might affect the normal function of the tissue or organ. Marked pathological (w = 3), the lesion was generally irreversible, leading to partial or total loss of the organ function. Moderate pathological (w = 2), the lesion was easily reversible in most cases if the stressor is neutralized. Minimal pathological (w = 1), the lesion was easily reversible as exposure to irritants ends.

Table S4. Differentially expressed genes of clams related to energy homeostasis and immunomodulation in the form of fragments per kilobase per million fragments (FPKM)

Gene ID	Gene description	Gene	FPKM ^α			Log2 (fold change) ^{β}		FDR ^γ	
	Gene description	name	СК	PS-NH ₂	PS-COOH	PS-NH ₂ PS-COOH PS-NH ₂	PS-NH ₂	PS-COOH	
Unigene0 086391	NF-kappa-B inhibitor alpha [<i>Ruditapes philippinarum</i>]	NFKBIA	17.34	3.60	12.44	-2.27	-0.48	0.01	1.00
Unigene0 014347	interleukin-1 receptor- associated kinase 4-like [Aethina tumida]	IRAK4	52.17	1.18	15.60	-5.47	-1.74	0.02	0.90
Unigene0 009386	PREDICTED: tubulin alpha-3 chain-like [Chinchilla lanigera]	TUBA	3.92	1.36	0.96	-1.53	-2.03	0.48	0.05
Unigene0 066643	PREDICTED: tubulin beta chain, partial [<i>Columba</i> <i>livia</i>]	TUBB	8.07	0.50	7.48	-4.02	-0.11	0.02	1.00
Unigene0 043181	cathepsin L [<i>Meretrix meretrix</i>]	CTSL	77.58	16.45	88.77	-2.24	0.19	0.01	1.00
Unigene0 001881	PREDICTED: alpha-2 adrenergic receptor [Crassostrea gigas]	CCKAR	2.30	19.26	1.25	3.06	-0.88	0.00	0.89
Unigene0 101284	PREDICTED: carboxypeptidase A2-like [Crassostrea gigas]	CPA2	0.00	1.90	1.53	10.89	10.58	0.00	0.65

Unigene0 010921	serine protease CFSP3 [Azumapecten farreri]	CELA2	0.00	4.87	7.09	12.25	12.79	0.01	0.43
Unigene0 014649	PREDICTED: phosphoenolpyruvate carboxykinase, cytosolic [GTP] isoform X2 [Crassostrea gigas]	PCK1	8.18	24.98	6.25	1.61	-0.39	0.00	1.00
Unigene0 056199	PREDICTED: acyl-CoA desaturase-like isoform X2 [Lingula anatina]	SCD-1	13.38	27.12	19.71	1.02	0.56	0.01	1.00
Unigene0 016927	PREDICTED: fatty acid- binding protein, intestinal isoform X2 [<i>Crassostrea</i> gigas] PREDICTED: ATP-	FABP3	5.73	33.23	11.13	2.54	0.96	0.02	1.00
Unigene0 072869	binding cassette sub-family A member 1-like, partial [Saccoglossus kowalevskii]	ABCA1	0.41	2.03	0.53	2.32	0.39	0.02	1.00

^{α} The FPKM represents the gene expression by normalizing the read counts in transcript. ^{β} Log2 (fold change) represents the fold change of gene levels in the PS-NH₂ and PS-COOH groups relative to the control group, respectively.

^γ FDR, the false discovery rate, represents the corrected *P*-value of gene levels in the PS-NH₂ and PS-COOH groups relative to the control group.



Fig. S1. TEM images of (a) PS-NH2 and (b) PS-COOH. Fluorescent images of (c) PS-
NH2 and (d) PS-COOH. The NPs were suspended in Milli-Q water (mQW). These data
were previously reported by Luan et $al.^{39}$

Fig. S2. A schematic diagram for the exposure experiment design and sampling protocols. After a 7-day exposure, two clams were randomly selected from each beaker to assess the filtration rate and oxygen consumption rate. Three of clams in each beaker were dissected to excise digestive glands and hemolymph for further analysis of toxic mechanisms of NPs on energy homeostasis and immunomodulation of clams. The weight of shell and soft calculate conditional index and tissue were measured to the water content.



Fig. S3. The spectra of PS-NH₂ and PS-COOH determined using Fourier transformed infrared spectroscopy (FTIR). The shaded parts represent the characteristic peaks of polystyrene polymer. The peaks presented in both of PS-NH₂ and PS-COOH was labeled with black arrows, whereas the characteristic peak only presented in PS-NH₂ was marked with blue arrows.



Fig. S4. SEM images of (a) PS-NH₂, (b) PS-COOH, (c) microalgae I. zhanjiangensis and (d) C. meülleri. The interaction between PS-NH2 and (e) the mixed algae solution of C. meülleri and I. zhanjiangensis (1: 1, v/v) (f) microalgae I. zhanjiangensis, and (g) C. meülleri. The interaction of PS-COOH with (h) the mixed algae solution of C. meülleri and I. zhanjiangensis (1: 1, v/v) (i) microalgae I. zhanjiangensis and (j) C. meülleri. The concentration of PS-NH2 or PS-COOH was 2 mg L⁻¹. The density of microalgae was 2 \times 10 5 cells mL $^{-1}$. The algae were incubated in an illumination incubator for 24 h at 25 °C under a 12: 12 h light: dark cycle. The panels (e1-g1) show the enlarged regions from the red frames in the panels (e-g). The red arrows indicate the NPs adsorbed the surface of algae. on



Fig. S5. Images of (a) microalgae *I. zhanjiangensis* (CK), mixed solution of *I. zhanjiangensis* and PS-NH₂ and PS-COOH at 0, 4, 8, 12, 16, 24 h. (b) Effect of PS-NH₂ and PS-COOH on the suspension stability of microalgae *I. zhanjiangensis*. The concentration of PS-NH₂ or PS-COOH was 2 mg L⁻¹. The density of microalgae was 2×10^5 cells mL⁻¹. The algae were incubated in an illumination incubator for 24 h at 25 °C under a 12: 12 h light: dark cycle. There was no significant different of OD value between CK and PS-NH₂/PS-COOH (Duncan's multiple-comparison test, n = 3, *P*>0.05). The error bars represent the standard deviation of the three replicates in each treatment.



Fig. S6. The histological observations of clam digestive glands. (a) Control digestive glands without PS-COOH exposure, (b-e) pathological digestive glands exposed with PS-COOH at 2 mg L⁻¹. The red arrows indicate the structure of digestive glands, including digestive tubule (dt), epithelial cell (ec), tubule lumen (tl), and intertubular tissues (int). The blue arrows indicate the digestive tubule lesions, including widening of the tubular lumen (wtl), epithelial cell exfoliation (ece), epithelial cell hypertrophy (ech), and necrosis (nes). The yellow arrows indicate the intertubular tissue lesions, including haemocytes infiltration (hai) and fibrosis (fis). (f) Histopathological condition indices (I_h) of the clam digestive glands, which were calculated according to the method described by Costa et al.³⁸ Different small letters indicate significant difference between different concentrations of PS-COOH (Duncan's multiple-comparison test, n = 3, P < 0.05). The error bars represent the standard deviation of the three replicates in each treatment.



Fig. S7. The histological observations of clam digestive glands. (a) Control digestive glands without PS-NH2 exposure, (b-d) pathological digestiveglands exposed with PS-NH2 at 0.02, 0.2, and 2 mg L⁻¹, respectively. (e) Control digestive glands without PS-COOH exposure, (f-h) pathologicaldigestiveglandsexposedwithPS-COOHat0.02,0.2,and2mgL⁻¹,respectively.



Fig. S8. The counts of downregulated and upregulated differentially expressed genes (DEGs) with absolute values of log 2 (fold change) > 1 and false discovery rate (FDR) ≤ 0.05 in the clams exposed to 2 mg L⁻¹ PS-NH₂ and PS-COOH.



Fig. S9. Significantly enriched Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways of differentially expressed genes (DEGs) in the clam digestive glands exposed to 2 mg L^{-1} PS-NH₂ and PS-COOH. The color and number of columns represent the KEGG term and the count of DEGs, respectively.

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