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# Inflammatory and oxidative responses to PET nanoplastics in the leech *Hirudo verbana*: a comparative analysis of acute and chronic exposure

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Nanoplastics (NPs) are emerging environmental contaminants with the potential to induce cellular stress and immune dysregulation in aquatic organisms. In this study, the freshwater leech *Hirudo verbana* was used as a non-conventional invertebrate model to investigate the effects of acute (24–72 hours) and chronic (1 week–1 month) exposure to polyethylene terephthalate nanoplastics (PET NPs). A multidisciplinary approach combining microscopy, histology, immunocytochemistry, and qPCR was employed to evaluate PET NP uptake and biological responses. PET NPs were internalised in leech tissues and detected in macrophage-like cells. Both exposure regimes triggered a time- and dose-dependent inflammatory response, characterised by macrophage-like cell recruitment, angiogenic remodelling, and upregulation of the pro-inflammatory marker *HmAlf-1*. Endothelial activation was confirmed by increased CD31 expression and neovascularisation. Furthermore, oxidative stress was evidenced by altered expression of glutathione S-transferase (*GST*) and superoxide dismutase (*SOD*) genes. Overall, PET NPs induced conserved immune and stress responses in *H. verbana*, supporting its relevance as an alternative model for nanoplastic ecotoxicology. These findings contribute to our understanding of NP-induced pathophysiology and reinforce the need for further investigation into the ecological impact of plastic pollution on freshwater invertebrates.

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## Environmental significance

Our findings underscore the ecological relevance of PET nanoplastics as emerging contaminants in freshwater systems. The observed immune activation, oxidative stress, and vascular remodelling in *H. verbana* highlight conserved biological responses to nanoparticle exposure and point to a limited capacity for long-term physiological compensation under chronic stress. These results emphasise the potential for PET nanoplastics to disrupt immune and tissue homeostasis in aquatic invertebrates, reinforcing the need for their inclusion in environmental risk assessments and regulation of plastic-derived nanomaterials.

## Introduction

The increasing production and widespread use of plastic materials have led to the pervasive presence of micro- and nanoplastics (MPs and NPs) in aquatic ecosystems, raising growing concern for environmental and public health. Once released, plastic debris undergoes progressive weathering and fragmentation, leading to the generation of these smaller and

more reactive particles whose environmental fate is difficult to predict, as it depends on a combination of physicochemical properties and local ecological conditions.<sup>1,2</sup> Notably, it is worth noting that, beyond their intrinsic persistence, an additional factor that further complicates the environmental fate of these particles is their strict connection with climate change. The production and degradation of plastics contribute significantly to greenhouse gas emissions, while climate change in turn amplifies their dispersion, fragmentation, and ecological impacts through extreme weather events, altered hydrological cycles, and ocean warming.<sup>3</sup> Based on these perspectives, this interaction highlights the interconnected risks for environmental, animal, and human health, broadening the context of nanoplastic pollution beyond the local ecological impacts.

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### Acid phosphatase (ACP) histoenzymatic assay

Cryosections, obtained as described above, were rehydrated in PBS for 10 minutes, incubated for 5 minutes in 0.1 M acetic acid–sodium acetate buffer and then, with the reaction mixture (0.1 M sodium acetate–acetic acid buffer, 0.01% naphthol phosphate, 2% *N,N*-dimethylformamide, 0.06% Fast Red,  $\text{MnCl}_2$  0.5 nM) for 90 minutes, at 37 °C. After several PBS washings, the slides were mounted with Citifluor (Citifluor Ltd., UK) and observed under the light microscope.

### RNA extraction and quantitative PCR (qPCR)

Leech tissues were frozen in liquid nitrogen and crushed with a mortar and pestle. Samples were then resuspended in 1 ml of TRIzol (Life Technologies, Carlsbad, USA), centrifuged at 11 270g for 10 minutes and then incubated for 5 minutes at room temperature. Subsequently, 200  $\mu\text{l}$  of chloroform was added and samples were centrifuged for 15 minutes at 13 700g at 4 °C. Once the different phases were obtained, 500  $\mu\text{l}$  of the supernatant, in which nucleic acids were present, was recovered and gently mixed with 500  $\mu\text{l}$  of isopropanol. After 10 minutes at room temperature, samples were centrifuged for 10 minutes, and the resulting RNA pellets were washed in EtOH 75% and resuspended in DEPC water. Samples were then quantified, and the RNA purity was evaluated with a 1% agarose gel. 2  $\mu\text{g}$  of RNA were retrotranscribed into cDNA using M-MLV reverse transcriptase (Euroclone S.p.a, Pero, Italy) in the presence of oligodT (Invitrogen, Thermo Fisher Scientific, Waltham, USA) of random hexamers in a final volume of 20  $\mu\text{L}$ . Quantitative real-time PCR (qPCR) was carried out in triplicate in a CFX Connect Real Time PCR Detection System (Bio-Rad, Hercules, USA) using the iTaq Universal SYBR™ master mix (BioRad, Hercules, USA) and 0.2  $\mu\text{M}$  each of forward and reverse primer, in a final volume of 15  $\mu\text{L}$ . The primers used for qPCR amplifications were designed with the web-interface software Primer3Plus and are listed in the table below. After initial denaturation, the PCR reaction was performed at 95 °C (10 s), 60 °C (5 s), and 72 °C (10 s) for 39 cycles. Relative gene expressions were calculated using the  $\Delta\Delta\text{Ct}$  method, in which the 18S gene was used as housekeeping (Table 1).

### Intracellular ROS detection

ROS production was measured using the fluorogenic probe 2',7'-dichlorodihydrofluorescein diacetate ( $\text{H}_2\text{DCFDA}$ ) (Molecular Probes, Eugene, OR, USA), which crosses the cell membrane and gets hydrolysed to 2',7'-dichlorofluorescein (DCF). Upon oxidation by ROS, DCF becomes fluorescent. Cryosections of treated and control samples were incubated with 10  $\mu\text{M}$   $\text{H}_2\text{DCFDA}$  for 30 minutes at 37 °C in the dark and nuclei were counterstained with DAPI. Fluorescence was measured at excitation 488 nm/emission 525 nm using a Nikon fluorescence microscope and digital camera system.

**Table 1** Primers used for qPCR analyses

Target genes	Primers	Product size (bp)
<i>Hm</i> AIF-1	Fw: 5'-GACCTCAAAGACAAGCAGGG-3' Rev: 5'-GGCCAATCTTCTCCAGCATC-3'	229
SOD	Fw: 5'-ATCCTCTTGAACCCACCACA-3' Rev: 5'-ATCTGGACGCACATCTTGT-3'	95
GST4A	Fw: 5'-AGACACATCGCCAGGACTAA-3' Rev: 5'-ACGGATACACGACTCCAAC-3'	127
18S	Fw: 5'-GATGGTGACTCTTGGATAACTTC-3' Rev: 5'-CTGCCTTCTTGGATGTG-3'	189

### Statistical analysis

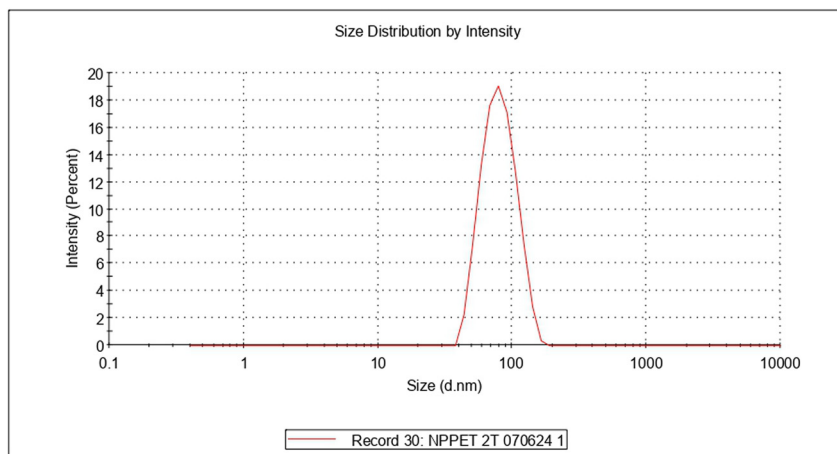
All the experiments were conducted in triplicate and the data show average values  $\pm$  standard deviations (SD). The fluorescence analysis for immunolocalisation approaches was performed using the ImageJ software package (<https://imagej.net/ij/index.html>) while the statistical analysis was conducted using the Graph Prism 8 software (GraphPad Software, La Jolla, USA; <https://www.graphpad.com/features>). Both the normality and the homoscedasticity of the samples were assessed using Shapiro–Wilk's and Levene's tests. The significant differences were calculated by one-way ANOVA variance analysis followed by Dunnett's *post hoc* test, in which a *p*-value <0.05 was considered statistically significant and confidence intervals were reported in the SI (S1–S6). In the graphs, the averages with asterisks represent a significant difference between leeches not exposed to or exposed to NPs PP. For immunofluorescence and vascular counts, five random 45 000  $\mu\text{m}^2$  fields per slide were analysed using ImageJ software.

## Results

### Characterisation of PET nanoparticles (NPs) and their internalisation in leech tissues

Prior to *in vivo* experiments, PET NPs were characterised using dynamic light scattering (DLS), transmission electron microscopy (TEM) and fluorescence microscopy (Fig. 1A–C). DLS measurements revealed an average diameter of  $82.48 \pm 24.76$  nm, confirming their nanoscale size (Fig. 1A). TEM analysis supported this finding, showing well-defined spherical particles (Fig. 1B). PET NPs were fluorescently labelled using Nile Red, allowing their visualisation under a Cy3 filter (excitation 550–570 nm) (Fig. 1C). The stability of the nanoparticles was further assessed through DLS analyses performed on PET nanoparticles without the fluorescent probe. The results, reported in Fig. S1, showed highly similar size distributions. Freshly prepared nanoparticles displayed a Z-average of 79.08 nm with a polydispersity index (PDI) of 0.071, while samples stored for nine months exhibited a Z-average of 78.69 nm with a PDI of 0.095. The main intensity peaks were centred at approximately 85–86 nm, with comparable standard deviations (24.5 vs. 26.3 nm). These data indicate that the suspensions remained monodisperse and stable over time, with no evidence of significant agglomeration (S7).





## Results

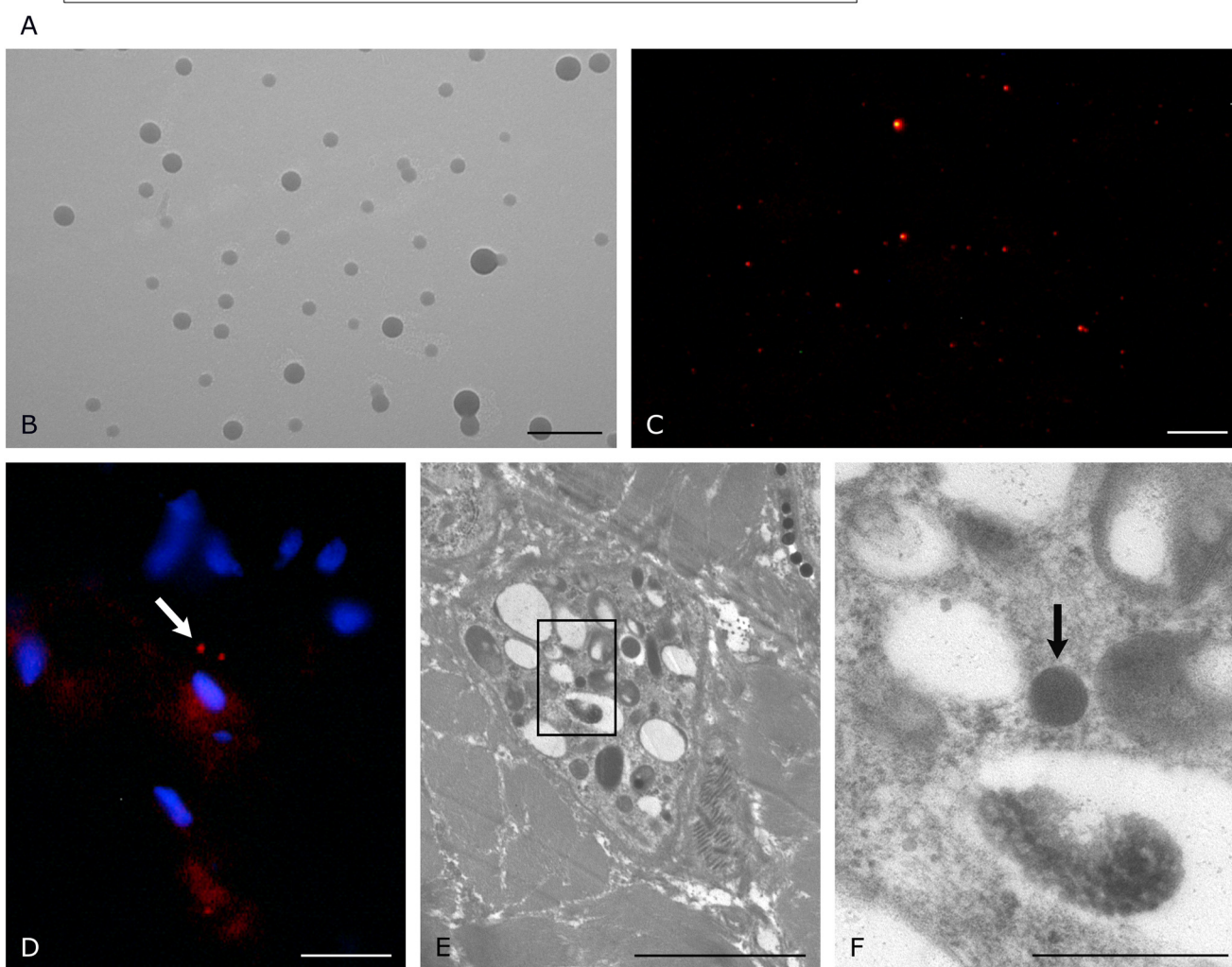
Z-Average (d.nm): 75.94

Pdl: 0.083

Intercept: 0.942

Result quality Good

	Size (d.nm)	% Intensity	St Dev (d.nm)
Peak 1:	82.30	100.0	23.38
Peak 2:	0.000	0.0	0.000
Peak 3:	0.000	0.0	0.000



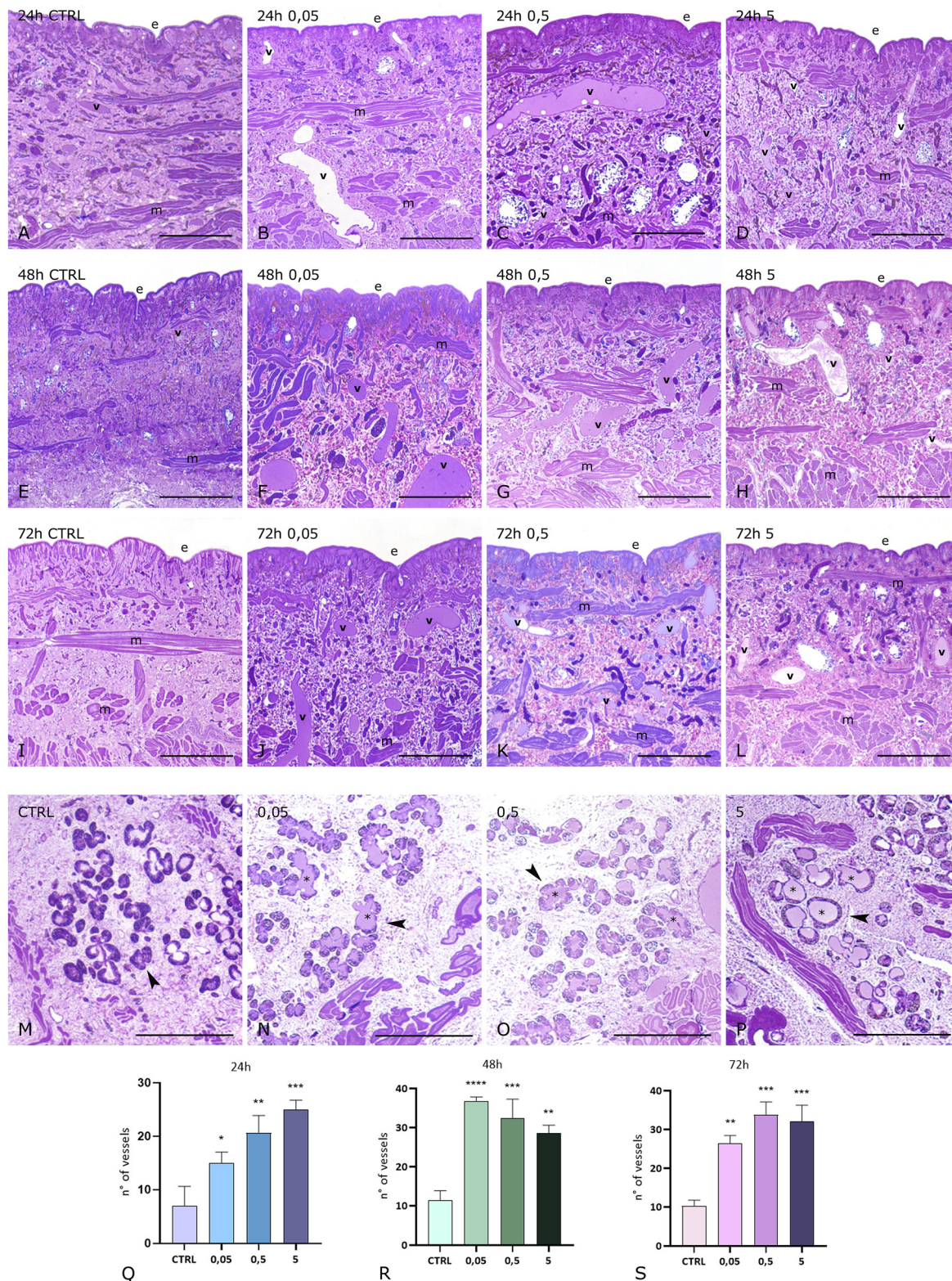
**Fig. 1** Characterisation and localisation of PET nanoparticles (NPs) using DLS, TEM, and fluorescence microscopy. (A) Dynamic light scattering (DLS) analysis showing the size distribution of PET NPs by intensity; (B) TEM image illustrating the morphology and size of PET NPs; (C) fluorescence microscopy image of PET NPs; fluorescent PET NPs (white arrows) detected in a cryosection of the leech body wall; cell nuclei are counterstained with DAPI (blue); (E) TEM image showing intracellular localization of PET NPs (arrows); (F) higher-magnification TEM detail of PET NPs highlighted in figure E. Bars in B and E: 2.5  $\mu\text{m}$ ; in C: 100  $\mu\text{m}$ ; in D: 10  $\mu\text{m}$ ; in F: 0.5  $\mu\text{m}$ .

The ability of PET NPs to penetrate *H. verbana* tissues was evaluated using fluorescence and TEM microscopy. Both techniques confirmed the internalisation of PET NPs within

cells, mainly localised in the subepithelial region (Fig. 1D–F), suggesting that active phagocytosis of these exogenous materials may occur.

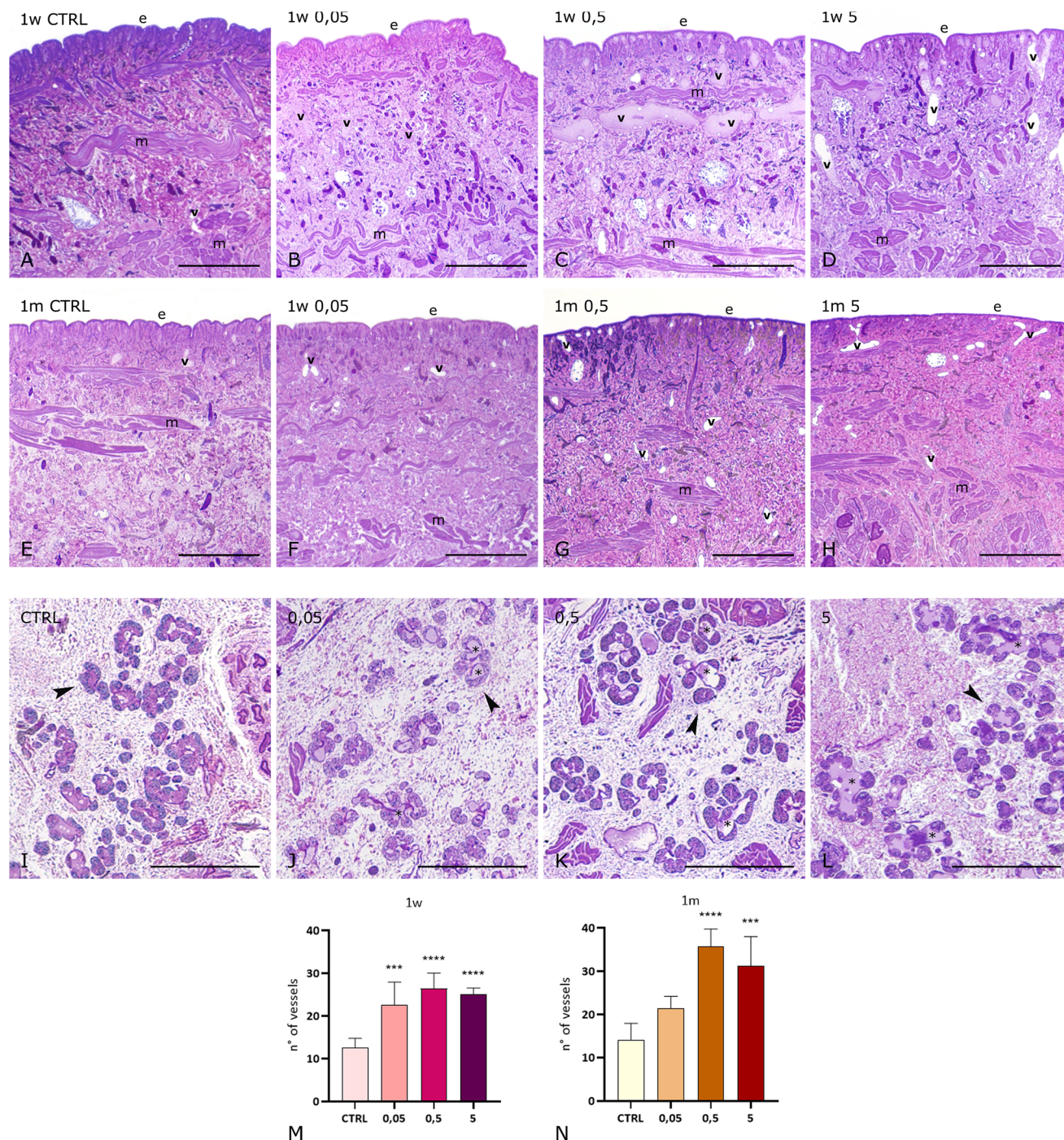






**Fig. 2** PET NPs induce angiogenesis and structural remodelling of botryoidal tissue in *H. verbanda*. Light microscopy images of the leech body wall following exposure to PET NPs at 0.05, 0.5, and 5 mg L<sup>-1</sup> for 24, 48, and 72 hours (A–L). Control tissues (A, E and I) appear largely avascular. Treated samples exhibit increased neovascularisation throughout the musculocutaneous sac (B–D, F–H, J–L). In controls (M), botryoidal tissue (arrowheads) appears as a compact cord-like structure. In PET NP exposed leeches (N–P) pre-vascular phenotype with lumen (asterisks) formation is visible. Graphs (Q–S) show vessel quantification, confirming a significant increase in vessel number at all time points and concentrations compared to controls. A dose-dependent increase is observed after 24 hours, with a peak at 5 mg L<sup>-1</sup>; at 48 hours, the highest vessel count occurs at 0.05 mg L<sup>-1</sup>; and at 72 hours, the peak response is observed at 0.5 mg L<sup>-1</sup>. In the graphs, \* means that  $p < 0.05$ ; \*\* means that  $p < 0.01$ ; \*\*\* means that  $p < 0.001$ . e epithelium, m muscle, v vessels. Bars in A–P: 100  $\mu$ m.





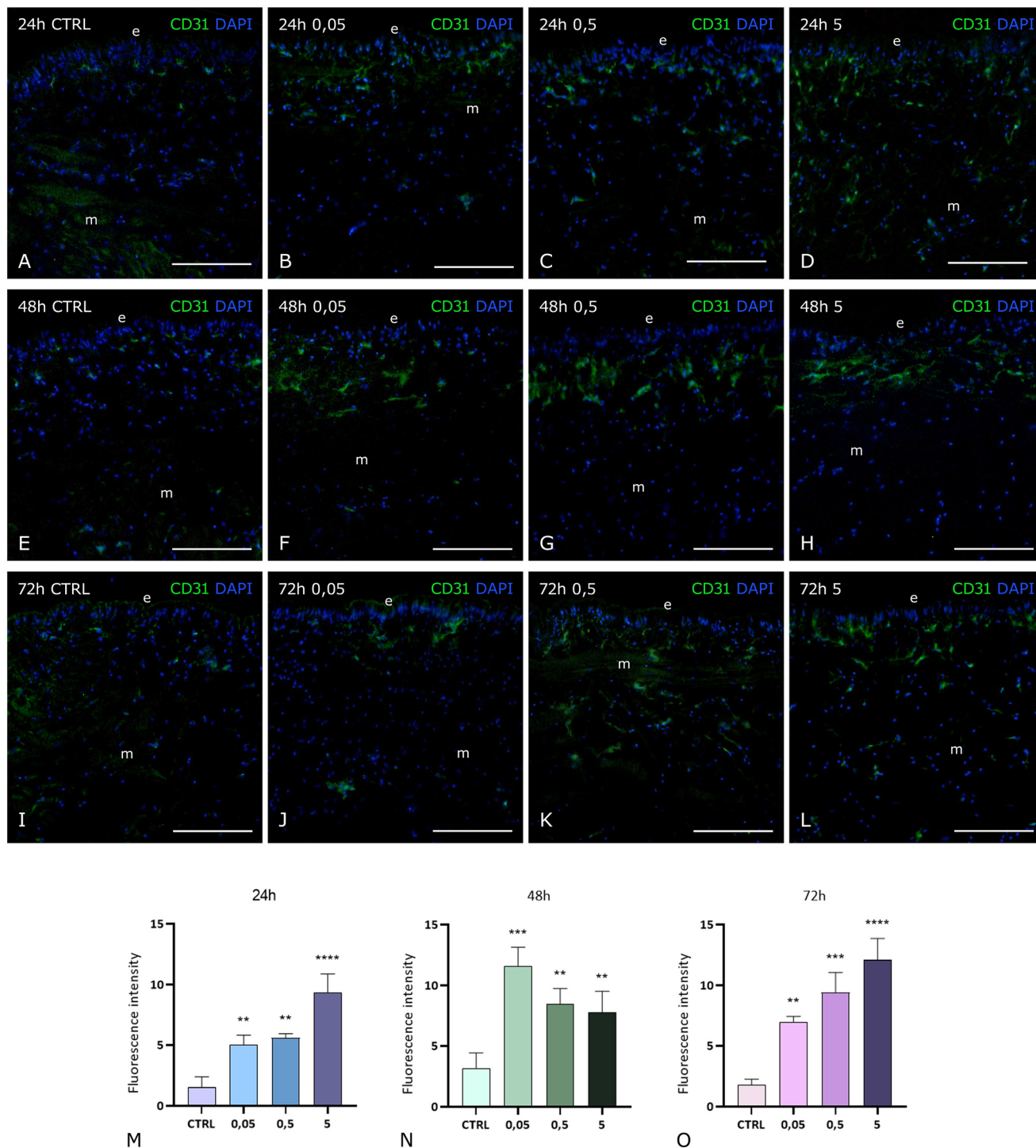
**Fig. 3** Chronic exposure to PET NPs induces persistent angiogenesis and remodelling of botryoidal tissue in *H. verbana*. Light microscopy images of the leech body wall after 1 week (A–D) and 1 month (E–H) of exposure to PET NPs at 0.05, 0.5, and 5 mg L<sup>-1</sup>. Control animals (A and E) display avascular tissues, while treated leeches show increased neovascularisation in a dose-dependent manner (B–D and F–H). In control leeches (I), botryoidal tissue (arrowheads) has a cord-like shape. Upon PET NP exposure (J–L), prevascular structures with open lumens (asterisks) are evident. Graphs (M and N) show quantitative vessel counts, confirming a statistically significant, dose-dependent increase in vascularisation after 1 week at all concentrations, and a persistent angiogenic effect after 1 month, with the highest significance at 0.5 mg L<sup>-1</sup>. In the graphs, \*\*\* means that  $p < 0.001$ ; \*\*\*\* indicates  $p < 0.0001$ . e epithelium, m muscle, v vessels. Bars in A–L: 100  $\mu$ m.

CD31 expression. At 1 month, the signal remained elevated, with significant effects observed at the intermediate (0.5 mg L<sup>-1</sup>) and highest (5 mg L<sup>-1</sup>) concentrations, suggesting a sustained pro-angiogenic effect over time.

#### Macrophage-like cell activation: ACP histoenzymatic assays

**Acute exposure.** In the medicinal leech, macrophage-like cells play a fundamental role in the innate immune response by



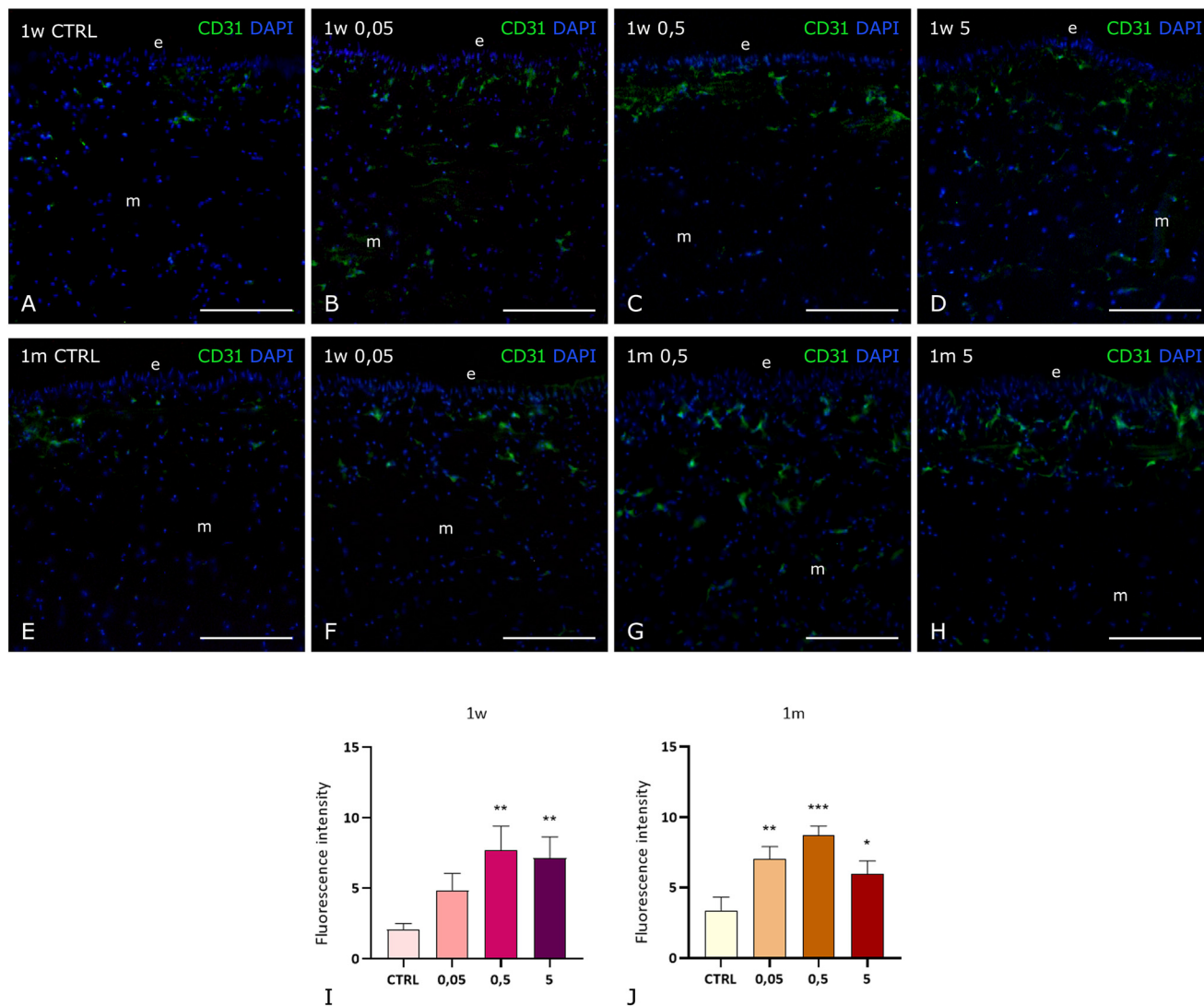


**Fig. 4** PET NPs induce CD31 expression in *H. verbanda*, confirming angiogenic activation. Immunofluorescence staining with anti-CD31 antibody in the leech body wall after exposure to PET NPs at 0.05, 0.5, and 5 mg L<sup>-1</sup> for 24, 48, and 72 hours (A–L). In untreated controls (A, E and I), only a weak CD31 (in green) signal is detected, indicating limited endothelial presence. PET NP-treated leeches show increased CD31 immunoreactivity at all concentrations and time points (B–D, F–H and J–L). At 24 hours, the signal intensity is elevated in all treated groups. At 48 hours, the strongest response is observed at 0.05 mg L<sup>-1</sup>, while at 72 hours the fluorescence intensity increased with dose, peaking at 5 mg L<sup>-1</sup>. Graphs (M–O) show quantification of CD31 fluorescence intensity, confirming a significant increase in endothelial marker expression in all PET NP-exposed groups compared to controls. In the graphs, \*\*\* means that  $p < 0.001$ ; \*\*\*\* indicates  $p < 0.0001$ . e epithelium, m muscle, in blue are nuclei stained with DAPI. Bars in A–L: 100  $\mu$ m.

recognising and eliminating non-self elements, such as pathogens and xenobiotics, through phagocytosis and encapsulation. To confirm the recruitment and activation of these innate immune

cells, previously suggested by TEM analysis, a histochemical acid phosphatase (ACP) assay was performed on leech tissues exposed to all concentrations and time points (Fig. 6).





**Fig. 5** Chronic exposure to PET NPs sustains CD31 expression and endothelial activation in *H. verbana*. Immunofluorescence images of the leech body wall stained with anti-CD31 antibody after 1 week (A–D) and 1 month (E–H) of exposure to PET NPs at 0.05, 0.5, and 5 mg L<sup>-1</sup>. In untreated controls (A and E), a weak CD31 signal is detected, indicating minimal vascular presence. Treated animals show a clear, dose-dependent increase in CD31<sup>+</sup> endothelial cells at both time points (B–D and F–H). Graphs (I and J) show quantification of CD31 fluorescence intensity, confirming a statistically significant increase after 1 week at all concentrations and after 1 month at 0.5 and 5 mg L<sup>-1</sup>. In the graphs, \* means that  $p < 0.05$ ; \*\* means that  $p < 0.01$ ; \*\*\* means that  $p < 0.001$ . e epithelium, m muscle, in blue are nuclei stained with DAPI. Bars in A–H: 100  $\mu$ m.

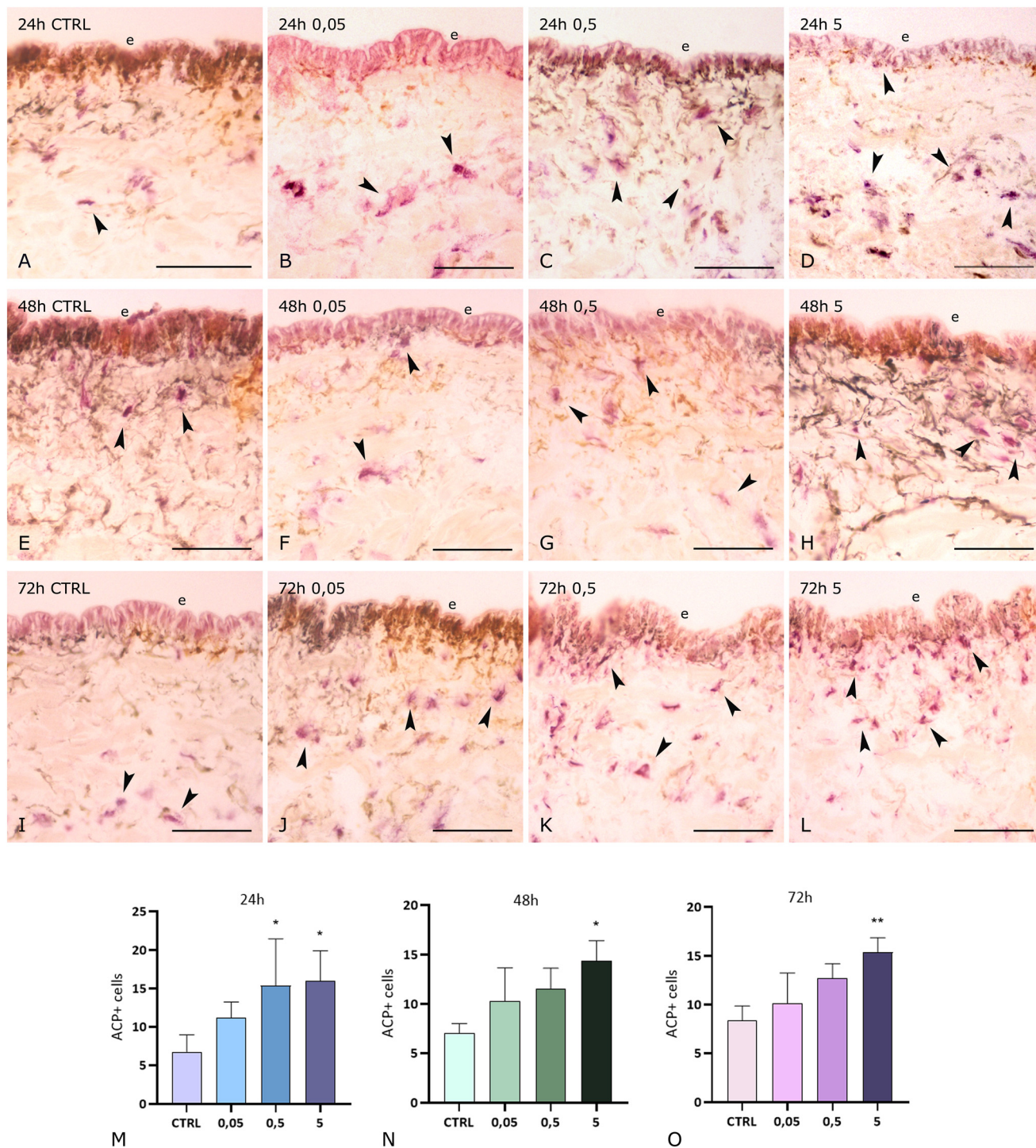
In control specimens not exposed to PET NPs (Fig. 6A, E and I), only a few ACP<sup>+</sup> cells were detectable, as expected under normal physiological conditions. Following PET NP exposure, a dose-dependent increase in ACP<sup>+</sup> cells was observed at all three time points (Fig. 6B–D, F–H and J–L). While tissues exposed to the lowest concentration showed results similar to controls (Fig. 6B and F), a marked accumulation of ACP<sup>+</sup> cells was visible at 0.5 and 5 mg L<sup>-1</sup>, localised within the musculocutaneous sac and underneath the epithelial surface (Fig. 6C, D, G, H, K and L).

Quantitative analysis of ACP<sup>+</sup> cells confirmed these histochemical results, showing a trend consistent with the microscopic observations (Fig. 6M–O).

**Chronic exposure.** To further investigate the long-term immune response to PET nanoplastics, the ACP

histoenzymatic assay was performed after 1 week and 1 month of exposure (Fig. 7A–H). As observed in the acute phase, the number of ACP<sup>+</sup> cells increases after exposure to PET NPs. After 1 week, the number of phagocytically active cells increased in a dose-dependent manner, with most ACP<sup>+</sup> cells located beneath the epithelium (Fig. 7B–D). At 1 month, numerous ACP<sup>+</sup> cells were still detectable, particularly in leeches treated with 0.5 mg L<sup>-1</sup>, and appeared to migrate from the inner body wall toward the epithelium (Fig. 7F–H), suggesting a sustained and spatially organised immune reaction. The quantification of ACP<sup>+</sup> cells (Fig. 7I and J) confirmed the histological observations, showing a statistically significant increase at all concentrations after 1 week, and a persistent elevation, most marked at 0.5 mg L<sup>-1</sup>, after 1 month of exposure.





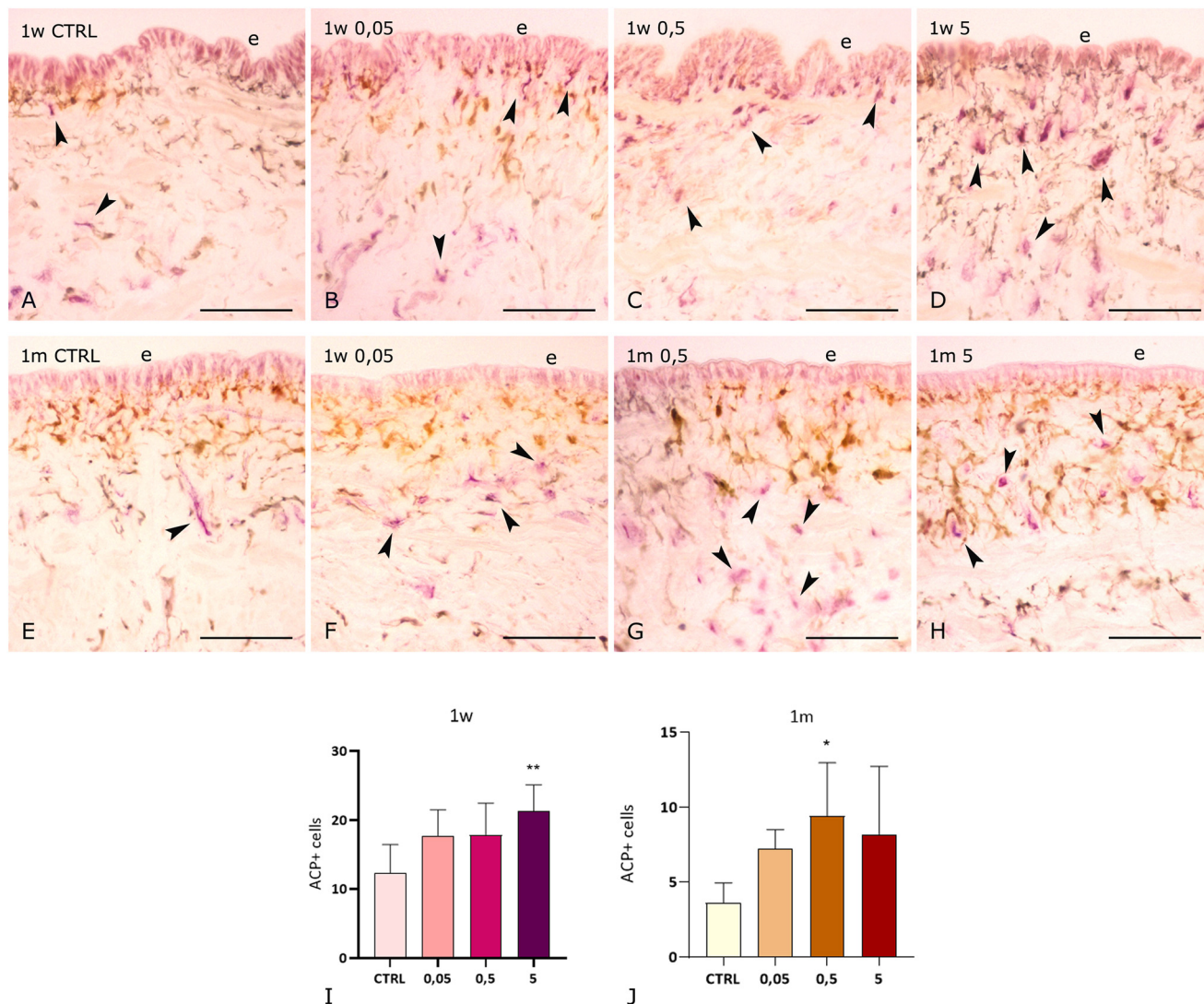
**Fig. 6** PET NPs modulate the recruitment of ACP<sup>+</sup> cells in *H. verbana*. ACP histoenzymatic staining of leech body wall sections following exposure to PET NPs at 0.05, 0.5, and 5 mg L<sup>-1</sup> for 24, 48, and 72 hours. In control samples (A, E and I), only a few ACP<sup>+</sup> cells are visible. In treated samples (B–D, F–H and J–L), a dose-dependent increase in ACP<sup>+</sup> cell number is observed, with the most evident activation at intermediate and high concentrations. Graphs (M–O) show the quantification of ACP<sup>+</sup> cells, confirming the temporal and concentration-dependent dynamics of macrophage activation. In the graphs, \* means that  $p < 0.05$ ; \*\* means that  $p < 0.01$ . e epithelium, arrowheads ACP<sup>+</sup> cells. Bars in A–L: 100  $\mu$ m.

### *Hm*AIF-1 as a marker of inflammation and macrophage-like cells

**Acute exposure and chronic exposure.** To further investigate the activation and recruitment of macrophage-like

cells in response to PET NPs, the expression of *Hm*AIF-1, a macrophage-specific pro-inflammatory cytokine, was analysed through qPCR and immunofluorescence (Fig. 8). After acute exposure, qPCR analysis revealed an increase in *Hm*AIF-1



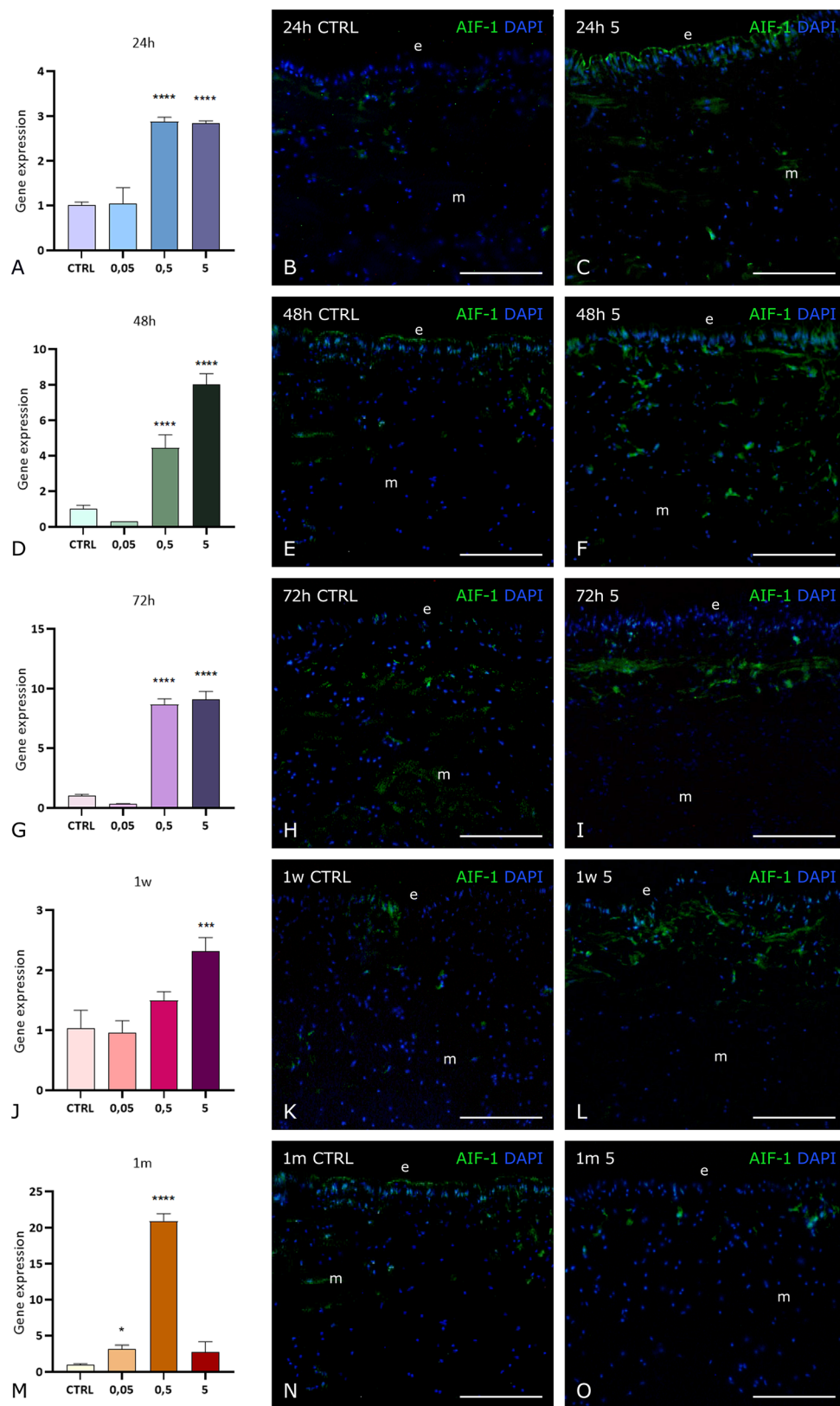


**Fig. 7** Chronic exposure to PET NPs activates ACP<sup>+</sup> cells in the *H. verbana* body wall. ACP histoenzymatic staining reveals the presence of phagocytically active ACP<sup>+</sup> cells after 1 week (A–D) and 1 month (E–H) of exposure to PET nanoplastics at concentrations of 0.05, 0.5, and 5 mg L<sup>-1</sup>. In control samples (A and E), few ACP<sup>+</sup> cells are observed. Treated specimens show a dose-dependent increase in ACP<sup>+</sup> cells after 1 week (B–D), with strong localisation near the epithelium. After 1 month (F–H), numerous activated ACP<sup>+</sup> are visible, particularly at 0.5 mg L<sup>-1</sup>, migrating toward the outer tissue layers. Graphs (I and J) quantify the total number of ACP<sup>+</sup> cells after 1 week (I) and 1 month (J) from treatment, confirming the microscopic results. In the graphs, \* means that  $p < 0.05$ ; \*\* means that  $p < 0.01$ . e epithelium, arrowheads ACP<sup>+</sup> cells. Bars in A–L: 100  $\mu$ m.

mRNA expression at all time points (Fig. 8A, D and G). While no upregulation in gene expression was evident at the lower NP concentration, a significant increase in gene expression was detected at 0.5 and 5 mg L<sup>-1</sup>, confirming a dynamic modulation of the inflammatory response in correlation with nanoparticle concentration and exposure time. Based on these results, immunofluorescence analyses were conducted comparing control tissues with those treated at the highest concentrations. In control leeches, only a few *HmAIF-1*<sup>+</sup> cells were detected beneath the epithelium (Fig. 8B, E and H). In contrast, PET NP-exposed tissues displayed a significant increase in *HmAIF-1* signal, particularly at the higher concentration (Fig. 8C, F and I), supporting the involvement of activated macrophage-like cells during the inflammatory response.

For the chronic exposure, immunolocalisation was performed at 1 week and 1 month (5 mg L<sup>-1</sup>), based on previous ACP data. In untreated animals (Fig. 8K and N), a low signal was again observed, while in treated animals after 1 week, the number of *HmAIF-1*<sup>+</sup> cells clearly increases in the subepithelial region, indicating a persistent immune cell activation. However, after 1 month of exposure to the highest PET NP concentration, a reduced number of positive cells, comparable to that observed in control samples, was detected. qPCR results at 1 week and 1 month (Fig. 8J and M) further confirmed these findings. After 1 week, a dose-dependent increase in *HmAIF-1* gene expression was detected, with peak expression at 5 mg L<sup>-1</sup>. After 1 month, a nonlinear trend was observed: a slight increase at 0.05 mg L<sup>-1</sup>, a marked and statistically significant peak at 0.5 mg L<sup>-1</sup>, followed by a decrease at 5 mg L<sup>-1</sup>. These data supported the





**Fig. 8** *HmAIF-1* expression reveals macrophage-like cell activation following acute and chronic PET NP exposure. Immunofluorescence images showing *HmAIF-1* expression in control (B, E and H) and PET NP-treated leeches at the highest concentration after acute exposure (C, F and I). Relative *HmAIF-1* mRNA levels (qPCR) at 24, 48, and 72 hours (A, D and G), showing dose-dependent expression patterns. *HmAIF-1* immunolocalisation under chronic exposure: control samples (K and N) and samples exposed to 5 mg L<sup>-1</sup> for 1 week and 1 month (L and O). qPCR quantification of *HmAIF-1* gene expression after 1 week and 1 month (J and M), confirming increased expression in treated samples. In the graphs, \* means that  $p < 0.05$ ; \*\*\* means that  $p < 0.001$ ; \*\*\*\* indicates  $p < 0.0001$ . e epithelium, m muscle. Bars in B, C, E, F, H, I, K, L, N and O: 100  $\mu$ m.



immunofluorescence and morphological observations, reinforcing the key role of *Hm*AIF-1 in mediating macrophage-like cell activation during the inflammatory response to PET nanoplastics.

### Assessment of oxidative stress induced by PET NPs

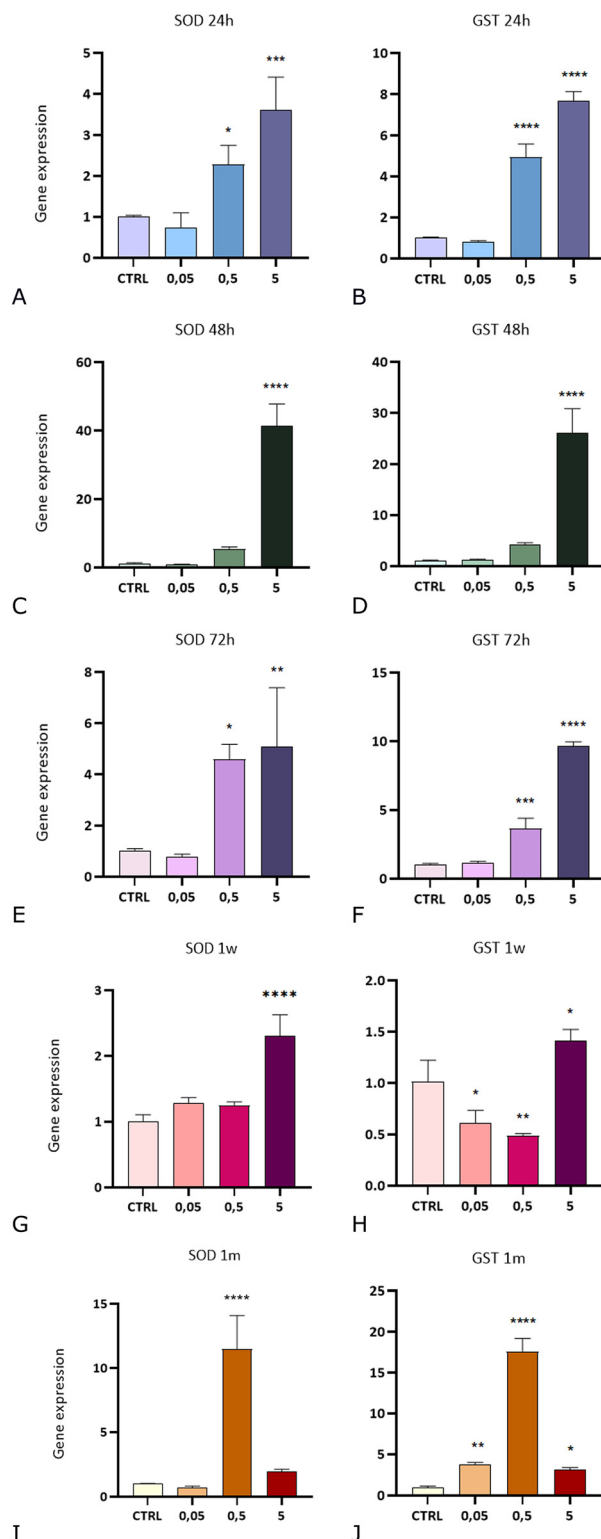
**Acute exposure and chronic exposure.** Plastic nanoparticles are well known to induce oxidative stress, primarily through the generation of reactive oxygen species (ROS) that disrupt cellular redox homeostasis. To investigate the antioxidant response elicited by PET NPs in *H. verbana*, qPCR analyses were performed to quantify the expression levels of two key antioxidant enzymes: glutathione *S*-transferase (*GST4A*) and superoxide dismutase (*SOD*) (Fig. 9). In the acute exposure, both *GST4A* and *SOD* genes showed a dose-dependent increase in expression at all tested time points (24, 48, and 72 hours), with maximum expression levels observed at the highest concentration (5 mg L<sup>-1</sup>) (Fig. 9A–F). These findings indicated that PET NPs rapidly triggered an oxidative stress response, leading to upregulation of protective antioxidant mechanisms.

In chronic exposure, gene expression patterns appeared less linear and more dynamic. Regarding *SOD*, a moderate increase in mRNA expression was noted after 1 week, especially at higher concentrations (Fig. 9G). After 1 month, however, *SOD* expression significantly increased only at 0.5 mg L<sup>-1</sup>, remaining notably low at the lowest and highest concentrations (Fig. 9I). For *GST4A*, after 1 week of exposure, gene expression initially decreased at the lowest concentrations, followed by a significant upregulation at 5 mg L<sup>-1</sup> (Fig. 9H). However, at 1 month, peak expression was recorded at 0.5 mg L<sup>-1</sup>, with a marked decline at 5 mg L<sup>-1</sup> (Fig. 9J), suggesting a time-dependent modulation of *GST4A* activity, possibly due to cellular adaptation or damage. This result was consistent with the *SOD* pattern and further supported the idea that the intermediate concentration may trigger a more effective or sustained antioxidant response, whereas extreme concentrations may lead to cellular exhaustion or toxicity. Comparable results were obtained by evaluating ROS (reactive oxygen species) production through the 2',7'-dichlorodihydrofluorescein diacetate (H<sub>2</sub>DCFDA) assay, as shown in SI (S9).

These findings clearly demonstrated that PET NP exposure induces oxidative stress, as evidenced by the transcriptional activation of *GST* and *SOD*, and that this response is both time- and dose-dependent. Furthermore, the production of ROS was significantly modulated following exposure, confirming that PET NPs trigger an imbalance between ROS generation and cellular antioxidant defences. This imbalance leads to oxidative damage and contributes to the observed cellular stress response, consistent with previous reports on nanoparticle-induced oxidative mechanisms.

## Discussion

The widespread environmental distribution of nanoplastics (NPs) and their accumulation in freshwater ecosystems pose a growing threat to aquatic organisms. Among these



**Fig. 9** Expression of antioxidant enzymes *SOD* and *GST4A* in response to PET NP exposure. Relative mRNA levels of *SOD* and *GST4A* after 24, 48, and 72 hours of PET NP exposure (A–F), showing a dose-dependent upregulation, with peak expression at 5 mg L<sup>-1</sup>. Gene expression of *SOD* and *GST4A* after 1 week and 1 month of chronic exposure (G–J). A significant increase at 0.5 mg L<sup>-1</sup> was observed for both enzymes at 1 month, while expression at 5 mg L<sup>-1</sup> declined. In the graphs, \* means that  $p < 0.05$ ; \*\* means that  $p < 0.01$ ; \*\*\* means that  $p < 0.001$ ; \*\*\*\* indicates  $p < 0.0001$ .







prey and host–parasite interactions, altering community composition and impairing many ecosystem functions, including nutrient cycling and organic matter decomposition. Over time, such disruptions may reduce biodiversity and compromise the resilience of freshwater ecosystems to additional stressors. These findings thus emphasise the importance of integrating nanoplastic contamination into ecological risk assessments and highlight the broader environmental significance of the responses observed in this model organism.

Future studies will focus on the impact of PET nanoplastics on regenerative processes in *H. verbana*, with particular attention to how chronic exposure may impair tissue renewal and post-injury repair, thus providing a more comprehensive understanding of the long-term ecological risks associated with nanoparticle contamination.

## Conclusions

This study provides a comprehensive overview of the inflammatory and immunological responses triggered by PET NPs in *H. verbana*. The findings highlight how this pollutant is able to induce the activation of well-conserved stress and defence pathways. Inflammatory and oxidative stress activation were shown to be strictly linked to increased concentration of NPs in a dose and time-dependent manner, despite showing limitations in long-term exposure. The observed biphasic pattern is indeed representative of the disruptive and unbalancing effect that PET NPs may exert on living organisms when constitutively exposed to high concentrations of pollutants, drawing more focus and the possible threatening consequences of plastic contamination. Overall, these results confirm *H. verbana* as a sensible model for ecotoxicological studies in freshwater environments, establishing a basis for future studies investigating nanoparticles effects on physiological mechanisms involved in immune response and regeneration events.

## Author contributions

C. B.: formal analysis, investigation, writing – review and editing. L. P.: formal analysis, investigation, validation, writing – review and editing. E. B.: formal analysis, investigation, writing – review and editing. S. A.: formal analysis, investigation, writing – review and editing. M. B.: formal analysis, investigation, writing – review and editing. D. T.: formal analysis, methodology, supervision; writing – review and editing. N. B.: formal analysis, funding acquisition, supervision; validation; writing – original draft. A. G.: conceptualization, funding acquisition, project administration, supervision, validation, writing – original draft.

## Conflicts of interest

The authors declare no conflicts of interest.

## Data availability

All data generated or analyzed during this study are included in this article and its supplementary information (SI).

Supplementary information: content of SI: 1. List of mean values and 95% confidence intervals for: vessel counts, CD31 expression, ACP-positive cells, and gene expression of *HmAIF-1*, *SOD*, and *GST*. 2. DLS analyses on non-fluorescent PET NPs. 3. Immunofluorescence negative controls. 4. ROS detection using fluorescence dye 2',7'-dichlorodihydrofluorescein diacetate ( $H_2DCFDA$ ). See DOI: <https://doi.org/10.1039/d5en00733j>.

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