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**Size-dependent effects of polystyrene micro- and nanoplastics on
the quality of rice grains and the metabolism mechanism**

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Abstract

Micro- and nanoplastics (MNPs) exhibited size-dependent impacts on crop physiological processes and metabolic functions, ultimately threatening crop quality. The size-specific response patterns warrant close attention to enable a more accurate assessment of their environmental and human health risks. In this study, both soil (10 mg/kg and 100 mg/kg) and hydroponic (1 µg/L and 10 µg/L) exposure systems were employed to investigate the effects of six spherical polystyrene (PS) MNPs particle diameters (80, 100, 200, 500, 800, and 1000 nm) on the grain quality of rice (*Oryza sativa* L.), and to further elucidate the underlying metabolic mechanisms. The results indicated that PS MNPs were absorbed through rice roots and transported upward, with concentrations in rice leaves ranging from 297 to 701 µg/g. When exposure to PS MNPs with diameters ≤ 100 nm (PS ≤ 100 nm), the reactive oxygen species (ROS) levels in the plants increased by 25%, and glutenin content in the grain decreased by up to 29%. Metabolomic analysis revealed that glycolysis/gluconeogenesis and pentose and glucuronate interconversion in rice leaves were closely associated with the particle diameters of PS MNPs. Compared to larger particles, exposure to PS ≤ 100 nm downregulated carbohydrate metabolism and decreased the accumulation of sugars (starch, soluble sugars), thereby weakening the plant's defense response. This study demonstrated that PS ≤ 100 nm triggered weaker defense response in rice, leading to greater biotoxicity, thereby deserving particular attention for governing the MNPs risk and crop safety assessment.

Keywords: Microplastics, Polystyrene, Rice, Particle size, metabolism, quality

1. Introduction

Micro- and nano plastics (MNPs) pollution poses critical environmental and health risks, calling for urgent global action. The extensive production and usage of plastics have resulted in the release of persistent, invisible, yet pervasive MNPs into the environment (Thompson. et al., 2004). Soil, particularly agricultural soil, functions as a major terrestrial sink for MNPs and serves as an important transmission node for MNPs-related risks (Rillig et al., 2024). MNPs entered agricultural ecosystems via sludge application (Weithmann et al., 2018), wastewater irrigation (Hasan & Tarannum, 2025), and plastic mulching (Qi et al., 2020). Recent surveys have shown that the abundance of MNPs in agricultural soils, particularly paddy soils, ranges from 10.3 ± 2.2 to 3805 ± 511 items/kg (Table S1). Most of these particles are smaller than 1 mm, with fibers and microspheres predominating. The presence of MNPs in agricultural soils may detrimentally interfere with the physiochemical functions of crops, impairing their growth, modifying the carbohydrate and protein levels in crop grains, and consequently affecting quality of edible parts (Hasan & Tarannum, 2025; Jin et al., 2022). Therefore, it was necessary to characterize the impact of MNPs on crop growth, development and crop quality, offering important scientific evidence for accurately regulating the risk of MNPs in agricultural soils.

MNPs disturbed crop quality through the entire life cycle of absorption, distribution and metabolism (Chen et al., 2022; Jamil et al., 2025; Shi et al., 2024). They entered crops via roots cracks (L. Li et al., 2020), were transported upward along the vascular cylinder under transpiration pull (R. Li et al., 2020; Liu et al., 2022), and

eventually reaching edible parts (Jiang et al., 2022). MNPs were capable of generating reactive oxygen species (ROS), triggering antioxidant defense response (e.g., antioxidant enzymes such as superoxide dismutase and peroxidase) (Ma et al., 2022; Zhang et al., 2021), and breaking important biomolecules such as carbohydrates, proteins or DNA (Y. Zhang et al., 2023; Zhou et al., 2021; Zhu et al., 2023). Particularly, sugars, including monosaccharides and oligosaccharides (i.e., soluble sugars), and polysaccharides (e.g., starch) are necessary energy materials for crop growth and development, and also vital substance in stress responses (Keunen et al., 2013). Exposure to MNPs detrimentally consumed soluble sugars to adapt oxidative stress. In response, starch was degraded to supplement soluble sugars and reshape carbohydrate metabolism network, thereby improving plant resilience (Asad et al., 2024; Wu et al., 2022). Therefore, MNPs induce oxidative damage in crops, alter the levels of key biomacromolecules such as carbohydrates and proteins, and disturb the overall balance of the metabolic network, thereby exerting adverse effects on crop physiology and metabolism.

Particle diameters impacted the absorption of MNPs into crops, thus may be exhibited as a determinant of the influence on crop growth, metabolism and grain quality (Shi et al., 2024; Yao et al., 2024). Crops tended to absorb nanoscale plastics (diameter < 100 nm) more efficiently than microscale plastics (diameter > 1 μ m), leading to their accumulation within plant tissues (Li et al., 2019; Zhu et al., 2022). Studies indicated that smaller MNPs (e.g., particle diameters of 50 nm and 100 nm) accumulating in plant tissues can obstruct intercellular connections and cell wall pores,

81 disrupt nutrient transport, and cause greater oxidative damage and metabolic
82 disturbance, ultimately may impair crop quality (Jiang et al., 2019; Zhu et al., 2023). In
83 contrast, microscale plastics tend to enhance photosynthesis in crops, while the plants
84 upregulate antioxidant substances (e.g., soluble sugars, soluble proteins and ascorbic
85 acid) to alleviate the biotoxicity of MNPs (He et al., 2024; Z. Li et al., 2020; C. Zhang
86 et al., 2023). Alterations in crop physiological responses and metabolic homeostasis
87 ultimately result in changes in crop quality. Moreover, MNPs particle diameters can
88 differentially affect crop quality by modulating plant physiological traits, metabolic
89 profiles, and enzyme activities. Thus, we hypothesize that MNPs with specific particle
90 diameter can enter crops, differentially modulate oxidative stress responses,
91 carbohydrates and related proteins synthesis, and the global metabolic network,
92 ultimately exerting distinct effects on crop quality.

93 Polystyrene (PS), a synthetic polymer made from monomers of the aromatic
94 hydrocarbon styrene, has been one of the most prevalent and hazardous types of
95 microplastics (Kik et al., 2020). Rice, as a staple crop for more than half of the global
96 population, contributes a significant portion of dietary calories worldwide (Paliwal et
97 al., 2018). Therefore, studying the exposure of rice to PS particles of different diameters
98 is of significant interest. This study aimed to characterize the impact of PS particles of
99 varied particle diameters on the quality of rice grains and reveal the underlying
100 biochemical mechanism, using the combination of soil and hydroponic cultivation,
101 electron microscopy characterization, physiochemical responses and omics analysis.
102 The specific objectives included: (1) to evaluate the impact of PS of different diameters

on rice grain quality; (2) to observe the absorption and accumulation of PS of different diameters in rice; (3) to compare the physicochemical responses of rice to PS of varied diameters; and (4) to elucidate the metabolic mechanisms through which PS different diameters affect rice. These findings will provide valuable insights into understanding the behavior of MNPs in plants, evaluating their potential risks to crop yield and quality, and determining the precise threshold for scientific supervision of MNPs risks in agricultural soil.

2. Materials and methods

2.1 MP characterization

Monodisperse PS microspheres with six diameters (80, 100, 200, 500, 800, and 1000 nm), representative of MNPs, were fluorescently labeled with Nile Blue chloride and purchased from Tianjin Big Goose Scientific Co., Ltd. (Tianjin, China). The morphology of the plastic microspheres was characterized using transmission electron microscopy (TEM) (JEOL, JEM – 1400 Flash, Japan), with detailed procedures provided in Text S1 of the Supporting Information. Fluorescence microscope (Olympus, U-RFL-T, Japan) was used to detect the light emitted by plastic microspheres under excitation. The particle diameters distribution and surface charge of the PS MNPs were determined using a zeta potential analyzer (Malvern, Zetasizer Nano ZS, United Kingdom). The chemical composition and functional groups of plastic microspheres were examined using Raman spectroscopy (WiTec, Alpha300R, Germany), with further details provided in Text S2.

2.2 Seed cultivation

Rice seeds (*Oryza sativa* L., Lianjing-6188) were purchased from the Lianyungang Academy of Agricultural Science (Jiangsu, China). The seeds were first rinsed using ultrapure water, and then soaked in ultrapure water for 5 min. After that, the seeds were sterilized by soaking in a 3% H₂O₂ solution for 30 min, followed by a 48-hour incubation in ultrapure water under dark conditions. The seeds were then hydroponically cultivated in 11 liters of Hoagland's nutrient solution (pH 5.5–6.0) in a greenhouse (25 ± 0.5 °C, 70 ± 5% humidity, 7000 lx, and 14-hour light/10-hour dark cycle). The components of Hoagland nutrient solution were summarized in Table S2.

2.3 Soil cultivation of rice

To investigate the effects of different PS diameters on rice grain quality, a soil cultivation experiment was conducted. Soil was collected from Huai'an in Jiangsu Province (Latitude: 32.2450°N, Longitude: 118.5415°E), with a pH of 7.2 and an electrical conductivity of 3944 µS/cm (Text S3). The soil was initially screened by hand, dried in the sun and sieved through a 2 mm sieve prior to use. PS particles of each diameters were uniformly incorporated into the soil using a stepwise dilution method as previously described (Chen et al., 2019), to obtain three concentration levels: 0 mg/kg (control group), 10 mg/kg, and 100 mg/kg. After a 14-day nursery period, rice seedlings with a height of 20–24 cm were selected and transplanted into buckets (Height of 240 mm, top diameter of 215 mm, bottom diameter of 185 mm, 8 plants per pot, 3 pots per treatment) which contained 5 kg of contaminated soil. Rice was irrigated every 2–3 days as needed to maintain continuous flooding. The soil cultivation experiment was conducted in a greenhouse (14-h photoperiod, 25/20 °C day/night

temperature, 70% relative humidity). Rice grains were harvested after a cultivation period of 6 months.

2.4 Grain Quality Analysis

After 180 days of soil cultivation, mature rice grains were harvested and dehulled using a rice huller. The dehulled grains were freeze-dried in a freeze dryer (Scientz, Scientz-10 N/A, China) for 48 hours, ground into powder, and sieved through a 0.15-mm mesh sieve for further analysis.

Protein, including albumin, globulin, prolamin, and glutenin, were sequentially extracted based on the solubility differences, and were quantified using the bicinchoninic acid (BCA) assay (Beyotime Biotech Inc, Shanghai, China). The method for extracting protein was described in Text S4. Grain starch and total sugars contents were measured using the Solarbio assay kit (Solarbio, Beijing, China).

2.5 Hydroponic cultivation of rice

A hydroponic experiment was conducted to explore the potential mechanisms underlying rice quality changes induced by PS of different particle diameters. After a 7-day cultivation period, a total of 24 plants of 15–16 cm length were transferred to brown glass vials that contained 480 mL of Hoagland nutrient solution (pH 5.5) for 8–10 days in a greenhouse ($25 \pm 0.5^{\circ}\text{C}$, $70 \pm 5\%$ humidity, 7000 lx, and 14 h light/10 h dark). Three exposure concentrations of PS, 0 $\mu\text{g/mL}$ (control group), 1 $\mu\text{g/mL}$ and 10 $\mu\text{g/mL}$ were selected. The rim of the vials were sealed with sponge stoppers to prevent exogenous interference. After 14 days of cultivation, rice seedling leaves and roots were harvested separately for subsequent analyses.

2.6 Uptake and accumulation of PS MNPs in rice roots and leaves

After 14 days of hydroponic cultivation, fresh roots of control group and 1 $\mu\text{g/mL}$ of PS MNPs exposure were collected. The roots were manually sliced into ultra-thin sections using a blade and placed on the surface water. A coverslip was placed over the sections to prepare manual slices. Cross-sections were observed using a laser scanning confocal microscope (Zeiss, LSM 880, Germany).

After 14 days of hydroponic cultivation, fresh leaves of control group, 1 and 10 $\mu\text{g/mL}$ of PS MNPs exposure were collected, stored at $-80\text{ }^{\circ}\text{C}$ for 24 hours, and then lyophilized using a freeze dryer (Scientz, Scientz-10 N/A) for 48 hours. The dried rice leaves and roots were ground into powder. The pretreatment method and quantification method for PS MNPs in rice samples followed the approach described by Li et al. (C. Li et al., 2021). A Pyrolysis-gas-chromatography/mass spectrometry (Py-GC/MS) system consisting of a microfurnace pyrolyzer (Frontier Laboratories, EGA/Py 3030D, Japan), a GC instrument (ThermoFisher, Trace 1300, USA), and a mass spectrometer (ThermoFisher, ISQ 7000, Singapore) was applied for the analysis of PS MNPs in rice leaves. Quality assurance and quality control (QA/QC) procedures were conducted prior to each analytical batch for mass calibration. Instrumental blanks were analyzed for every six samples. To minimize potential contamination, a piece of quartz wool was placed at the opening of the pyrolysis cup after sample preparation. Experimental details of the Py-GC/MS analysis are provided in Text S5.

2.7 Physiological and Biochemical Analysis

After hydroponic cultivation for 14 days, fresh rice leaves and roots were ground

into powder in liquid nitrogen. Superoxide dismutase (SOD) and peroxidase (POD) activities were determined using the methods reported in previous studies (Charles & Irwin, 1971; Q & Z, 1999). Malondialdehyde (MDA) was determined using the thiobarbituric acid reactive substances (TBARS) method. ROS levels were measured using Meimian assay kits (Jiangsu Meimian industrial CO., Ltd, Jiangsu, China). Soluble sugars content were determined using Solarbio assay kits (Solarbio, Beijing, China).

After 14 days of hydroponic cultivation, fresh rice leaves and roots were harvested, and dried rice samples were prepared using the same method. Proteins were extracted and determined following the method described in Section 2.4.

2.8 Extraction and analysis of metabolites

After 14 days of hydroponic cultivation, 10 mg of ground rice powder from the control group and 1 µg/mL of PS exposure were collected for metabolic analysis, and the metabolite extraction method is provided in Text S6. Metabolites were immediately determined using an ultra-performance liquid chromatography (UPLC) system coupled with an UPLC-QTRAP 5500+ system (SCIEX, Framingham, USA). Analysis was performed using both positive and negative ion scanning modes. Experimental details and instrument parameters for the UPLC analysis are provided in Text S7 and Table S3, respectively. Statistical analysis of metabolomics data was performed using the online MetaboAnalyst 6.0 (<http://www.metaboanalyst.ca/>).

2.9 QA/QC and Statistical analysis

To minimize contamination, dust-proof clothing and latex gloves were worn

throughout the experiment. Plastic materials were avoided, and metal, glass, or quartz wool were used as substitutes. Each treatment group had at least three biological replicates.

The data were analyzed and visualized using R software, and results are presented as means $\pm$ standard deviation. One-way analysis of variance (ANOVA) was used to compare the statistical differences between groups, and Student *t-test* was used to compare the control and treatment groups. A *P*-value less than 0.05 was considered to indicate a statistically significant difference ($P < 0.05$, *; $P < 0.01$, **). To further evaluate the effect size between the treatment and control groups, Cohen's *d* was calculated using Python.

3. Results and discussion

3.1 Characterization of PS MNPs

According to TEM analysis, PS MNPs exhibited a structurally uniform, smooth, spherical morphology with good dispersibility (Fig. 1a–b, Fig. S1). Under the fluorescence microscope, plastic microspheres emitted red fluorescence with a maximum excitation wavelength of 620 nm and a maximum emission wavelength of 680 (Fig. 1c). As the diameters of PS particles increased, their zeta potential increased accordingly (Fig. 1d). The zeta potential of 1000 nm PS MNPs reached 12.13 ± 0.50 mV, whereas PS MNPs diameters smaller than 100 nm ($PS \leq 100$ nm) had the lowest value (-48.23 ± 0.92 mV for 100 nm PS MNPs). PS MNPs of different diameter sizes exhibited relatively narrow hydrodynamic size distributions, with distinct peaks for each particle group and overall uniformity (Fig. 1e). Raman spectra were collected in

the range of 500 to 3500 cm^{-1} , and characteristic bands of PS MNPs were observed at 621, 1001.7, 1031.7, and 1602.7 cm^{-1} (Fig. 1f). The band at 621 cm^{-1} corresponded to the ν_{6b} radial ring-stretching mode, whereas the prominent peak at 1001.7 cm^{-1} was attributed to the ν_{12} C-CC ring bending mode. The characteristic PS MNPs band at 1031.7 cm^{-1} was assigned to the ν_{18a} tangential C-H bending mode, and the peak at 1602.7 cm^{-1} arises from the aromatic ring stretching (ν_{12}) (Lim et al., 2015). Additionally, the bands observed at 2910.2 and 3057.9 cm^{-1} were consistent with aliphatic C-H stretching and aromatic C-H stretching vibrations, respectively (Jiang et al., 2022).

3.2 Uptake and accumulation of PS MNPs in rice

The absorption and distribution of PS MNPs in rice roots were observed using a laser scanning confocal microscope and shown in Fig. 2. In the control group, noticeable fluorescence signals were observed in the epidermis along the outer ring of the root cross-section, which were attributed to the natural autofluorescence of rice tissues. No significant fluorescence signals were detected in the stele, including the xylem and phloem (Fig. 2a–c). After exposure to PS MNPs, in addition to the autofluorescence in the rice root epidermis, bright fluorescent aggregates were observed on the outer epidermal layer. Furthermore, distinct single-point fluorescence signals were detected in the xylem and phloem of the stele (Fig. 2d–u). The single-point fluorescence signals in the stele indicated that PS MNPs were absorbed by rice roots and primarily accumulated in the xylem and phloem. Driven by transpiration and supported by cohesion and adhesion, water columns in the xylem may facilitate the

upward movement of PS MNPs to aboveground tissues (Liu et al., 2022; C. Zhang et al., 2023). In addition, leaching experiment indicated that the fluorescent-labeled PS MNPs remained stable after 18 months of storage, without detectable release of fluorescent dye (Fig. S2–S3).

The concentration of PS MNPs in rice leaves was further quantified using Py-GC/MS (Fig. S4). The results indicated that rice has a certain enrichment capacity for PS MNPs, with smaller particle diameters and higher concentrations leading to greater accumulation levels. Meanwhile, no distinct PS MNPs peaks were observed in the control group. Interestingly, the mean accumulation of 100 nm PS MNPs was higher than that of 80 nm particles. Specifically, the accumulation levels of 100 nm PS MNPs reached 445 $\mu\text{g/g}$ at 1 $\mu\text{g/mL}$ and 701.69 $\mu\text{g/g}$ at 10 $\mu\text{g/mL}$. However, no significant difference was observed between the 80 nm and 100 nm treatment groups at the same concentration.

3.3 Size-dependent effects of PS MNPs on rice quality

Protein is the second-largest storage substance in rice grains and plays a critical role in determining the eating and nutritional quality of rice, with its content influenced by environmental factors (Li et al., 2016). Rice proteins are classified into four categories based on solubility, including: water-soluble albumin, salt-soluble globulin, alcohol-soluble prolamin, and alkali-soluble glutenin (Amagliani et al., 2017). Among them, glutenin accounts for over 80% of the total endosperm protein, prolamin comprises 5%–10%, globulin 5%, and albumin 10% (Xinkang et al., 2023). In general, albumin and globulin serve as stress defense proteins (Yan et al., 2024). In this study,

albumin and globulin contents increased only when exposure to PS MNPs with particle diameters ≥ 800 nm (PS ≥ 800 nm), while remaining relatively unchanged in other treatment groups (Fig. 3a–b). The change in albumin content was not statistically significant. Under exposure to PS ≥ 800 nm, globulin content increased by 23.34% at 10 mg/kg and 28.00% at 100 mg/kg (800 nm). Prolamin content exhibited a positive correlation with PS MNPs diameter diameters (Fig. 3c), and under exposure to PS ≥ 800 nm, it increased by 51.06% at 10 mg/kg and 48.34% at 100 mg/kg (1000 nm). PS MNPs exposure significantly reduced glutenin content, and it showed a slight increase with increasing PS MNPs diameters (Fig. 3d). Specifically, glutenin content decreased by 28.50% at 10 mg/kg (100 nm) and 24.07% at 100 mg/kg (80 nm) under PS ≤ 100 , respectively. As the most abundant protein in the endosperm, glutenin is rich in lysine and highly digestible, making it the most nutritionally valuable protein (He et al., 2021).

Rice grains contain 50%-90% starch, which serves as the primary energy storage substance (Razzaq et al., 2020). PS MNPs exposure had a negligible impact on starch content in rice, although a slight increase in grain starch content was observed with larger PS MNPs diameters (Fig. 3e). Total sugars, including soluble sugars such as glucose, fructose, and partially hydrolyzed sugars, directly affects the sweetness of rice grains. Total sugars content gradually increased with PS MNPs diameters, showing a significant rise compared to the control group (Fig. 3f). Under exposed to PS ≥ 800 nm, total sugars content increased by 38.67% at 10 mg/kg and 38.8% at 100 mg/kg (1000 nm).

Based on the grain quality results, the effect of exposure concentrations (10 mg/kg,

100 mg/kg) was smaller than that of the PS particle diameters (80, 100, 200, 500, 800, and 1000 nm) (Table S4). Additionally, Cohen's d was calculated to evaluate the effect size between the PS MNPs treatment groups and the control group (Table S5). Overall, PS MNPs had a minimal effect on stress-response proteins in rice, with their levels increased only under exposed to $PS \geq 800$ nm. Meanwhile, the nutritional quality of rice grains proteins was significantly affected, with the most severe loss observed at $PS \leq 100$ nm. In contrast, PS MNPs exposure increased sugars content in rice grains, with levels gradually rising alongside particle diameters.

3.4 Size-dependent effects of PS MNPs on rice physiological properties.

ROS accumulation leads to a decline in grain quality (Suriyasak et al., 2017). Under non-stress conditions, ROS are continuously generated at basal levels in plants and are effectively scavenged by antioxidant mechanisms (enzymatic and non-enzymatic), thereby avoiding damage (Demidchik, 2015). Under stress conditions, the delicate balance between ROS production and scavenging in plants is disrupted, potentially leading to ROS accumulation in plant tissues (Wu et al., 2021) and oxidative stress (Arora. et al., 2002).

PS MNPs exposure to rice roots caused excessive ROS accumulation in rice (Fig. 4a, Fig. S5a), while ROS content gradually decreased as PS MNPs diameters increased. In leaves, exposure to $PS \leq 100$ nm resulted in a significant increase in ROS levels, reaching 25.22% at 1 $\mu\text{g/mL}$ and 21.18% at 10 $\mu\text{g/mL}$ (80 nm) (Fig. 4a), with other treatment groups showed minor ROS increases. In root, ROS content increased by 22.90% at 1 $\mu\text{g/mL}$ (100 nm) and 22.45% at 10 $\mu\text{g/mL}$ (80 nm) under $PS \leq 100$ (Fig.

S5a). A statistical comparison of groups with the same PS MNPs diameter but different exposure concentrations (1 $\mu\text{g/mL}$ and 10 $\mu\text{g/mL}$) showed that particle diameters had a higher effect on ROS than concentration (Table S6). When excessive ROS accumulated, rice activated a series of antioxidant strategies to sustain normal physiological functions. SOD is the most critical ROS scavenger in organisms and plays a key role in maintaining redox balance. Under exposure to PS MNPs, SOD activity was significantly increased in rice leaves (Fig. 4b) while significantly decreased in roots (Fig. S5b), indicating an impaired antioxidant capacity. In rice leaves, SOD activity initially decreased and then plateaued as PS MNPs diameters increased. Under exposure to $\text{PS} \leq 100$ nm, SOD activity reached a maximum increase of 74.20% at 1 $\mu\text{g/mL}$ and 79.33% at 10 $\mu\text{g/mL}$ (80 nm). In rice root, SOD activity was significantly inhibited, with exposure to $\text{PS} \leq 100$ nm causing the greatest suppression, reaching 23.40% at 1 $\mu\text{g/mL}$ (80 nm) and 26.18% at 10 $\mu\text{g/mL}$ (100 nm). POD is a heme-containing enzyme that uses H_2O_2 as an electron acceptor to catalyze various oxidation reactions (Passardi et al., 2005). POD activity decreased to varying degrees under PS MNPs stress only in rice roots (Fig. S6). MDA, a final product of lipid peroxidation, is an indicator of membrane damage (F. Huang et al., 2023). MDA content remained unchanged under PS MNPs exposure (Fig. 4c, Fig. S5c), with a notable increase observed only under $\text{PS} \leq 100$ nm exposure in leaves, reaching up to 38.90% at 1 $\mu\text{g/mL}$ and 22.98% at 10 $\mu\text{g/mL}$ (100 nm) (Fig. 4c). Compared to larger particle diameters, $\text{PS} \leq 100$ nm may more easily pass through and damage cell walls and membranes, leading to higher levels of oxidative damage.

Soluble sugars are dissolved carbohydrates in organisms, primarily including monosaccharides and oligosaccharides such as glucose, fructose, and sucrose. These carbohydrates can act as ROS scavengers (Couee et al., 2006), with $\cdot\text{OH}$ scavenging capacity ranked as sucrose > fructose > glucose (Morelli. et al., 2003). Thus, their increased content can be regarded as a stress resistance mechanism. Soluble sugars content increased with PS MNPs diameters in rice. Under exposure to $\text{PS} \geq 800$ nm, soluble sugars content in leaves increased by 55.57% at 1 $\mu\text{g/mL}$ and 53.07% at 10 $\mu\text{g/mL}$ (800 nm) (Fig. 4d), while in roots, it increased by 54.38% at 1 $\mu\text{g/mL}$ and 42.20% at 10 $\mu\text{g/mL}$ (800 nm) (Fig. S5d). Starch is an insoluble, non-structural carbohydrate composed of α -glucose polymers (Pfister & Zeeman, 2016). When degraded, it releases energy and produces sugars and their derivatives, serving as an indirect indicator of stress resistance (Wu et al., 2022; Yan et al., 2024). Starch content in rice leaves (Fig. 4e) and roots (Fig. S5e) followed the same trend, increasing with PS MNPs diameters.

Albumin and globulin serve as stress defense proteins and can provide energy reserves for plants. Under PS MNPs stress, albumin content in rice leaves (Fig. S7a) and roots (Fig. S7b) gradually decreased with increasing PS MNPs diameters. Under exposure to $\text{PS} \geq 800$ nm, albumin content in leaves declined by up to 14.23% at 1 $\mu\text{g/mL}$ (800 nm) and 11.27% at 10 $\mu\text{g/mL}$ (1000 nm), while in roots, the decrease reached 9.14% at 1 $\mu\text{g/mL}$ and 9.65% at 10 $\mu\text{g/mL}$ (800 nm). In contrast, globulin content increased with PS MNPs diameters. Under exposure to $\text{PS} \geq 800$, globulin levels in rice leaves (Fig. 4f) increased by 19.47% at 1 $\mu\text{g/mL}$ and 25.09% at 10 $\mu\text{g/mL}$ (1000 nm), while in roots (Fig. S5f), the increase reached 14.02% at 1 $\mu\text{g/mL}$ and 10.52% at 10 $\mu\text{g/mL}$ (1000 nm).

The inverse trends in albumin and globulin content with increasing PS MNPs diameters suggested an activation of stress resistance mechanisms in rice to counteract the effects of larger particles. Additionally, the rise in sugars content might have partially compensated for protein loss, contributing to the maintenance of physiological and biochemical balance in rice. Under PS MNPs exposure, prolamin content in rice increased with PS MNPs size, while glutenin content showed no significant changes (Fig. S9). Under exposure to $PS \geq 800$ nm, prolamin content in rice leaves (Fig. S8a) increased by 17.06% at 1 $\mu\text{g/mL}$ and 22.69% at 10 $\mu\text{g/mL}$ (1000 nm), while in roots (Fig. S8b), the increase reached 16.08% and 28.68%, respectively.

Overall, the impact of exposure concentrations (1 $\mu\text{g/mL}$, 10 $\mu\text{g/mL}$) was smaller than that of the PS MNPs particle diameters (80, 100, 200, 500, 800, and 1000 nm) (Table S6). Cohen's d was also calculated to quantify the effect size between the PS MNPs treatment groups and the control group (Table S7-S8). Exposure of rice roots to PS MNPs induced ROS accumulation, and roots experienced greater oxidative damage than the leaves, disrupting intracellular redox balance. In response to excessive ROS, rice modulated the activities of SOD and POD. As PS MNPs diameters increased, rice enhanced the levels of soluble sugars, sucrose, and globulin, alleviating PS MNPs stress while reducing plant damage.

3.5 Impact of PS MNPs on the global metabolic network in rice

To further elucidate the mechanisms by which PS MNPs affected rice physiology, we conducted metabolomics analysis to examine the variation of final products of genome-environment interaction. Since the effects of PS MNPs exposure

concentrations (1 $\mu\text{g/mL}$, 10 $\mu\text{g/mL}$) on rice physiology was minimal, we selected 1 $\mu\text{g/mL}$ PS MNPs treatment group for further metabolic analysis.

Principal component analysis (PCA) was performed to determine metabolic difference among PS-treated and control groups. As illustrated in Fig. 5a–b, PC1 and PC2 cumulatively explained 55.3% and 76.2% of the metabolic variation in leaves and roots, respectively. Totally, 131 metabolites in leaves and 74 in roots were identified using an LC-MS/MS system (Fig. 5c–d). Based on the selection criteria ($P < 0.05$, $-1 < \log_2(\text{fold change}) < 1$), differentially expressed metabolites (DEMs) in leaves and roots under exposure to PS MNPs of different diameters were identified. As shown in Fig. S10–S11, more than 60% of the identified DEMs were upregulated. The DEMs were subjected to KEGG pathway analysis, and the top 15 significantly perturbed pathways were selected according to pathway impact and P -value. Fig. S12 showed that the significantly perturbed pathways in rice leaves across treatments were similar, predominantly involving carbohydrate metabolism and a similar trend was observed in roots (Fig. S13). Nine significantly perturbed metabolic pathways shared among treatment groups were identified. The relative proportion of each significantly disturbed metabolic pathway was assessed, in which carbohydrate metabolism being the most prominent, followed by amino acid metabolism (Fig. S14). The perturbation levels of 12 carbohydrate metabolism pathways were further assessed, with galactose metabolism showing the highest disturbance (Fig. S15).

Six metabolites involved with galactose metabolism pathway were identified in leaves, and three related metabolites were detected in roots. Interestingly, the relative

abundance of the six metabolites detected in leaves was generally upregulated (Table S9), while the abundance of the three metabolites detected in roots was downregulated (Table S10). Most metabolites in plant galactose metabolism, such as galactose, galactitol, act as non-enzymatic antioxidants, scavenging ROS and enhancing plant tolerance to exogenous stress (Bolouri-Moghaddam et al., 2010). Exposure to PS nanoplastics induced the upregulation of galactose metabolism in crops, which not only improved stress tolerance but also met the demands of energy metabolism (Lian et al., 2020). As amino acid metabolism is the second most disturbed metabolic pathway, we further analyzed the amino acids obtained from rice leaves and roots using cluster analysis (Fig. S16–17). Similar to soluble small molecule sugars, amino acids play a significant role in plant development and stress defense (Trovato et al., 2021). Under PS MNPs exposure, most amino acids detected in the leaves were upregulated, whereas the majority in the roots were downregulated. PS MNPs exposure induced a high level of oxidative stress in rice. To maintain normal physiological and biochemical functions, rice upregulated carbohydrate and amino acid metabolism pathway to alleviate oxidative stress.

3.6 The size-dependent effects of PS MNPs on rice metabolism

To further investigate the size-dependent effects of PS MNPs exposure on rice metabolism and uncover the metabolic mechanisms underlying grain quality changes, a weighted gene co-expression network analysis (WGCNA) was performed to identify metabolites highly correlated with exposure diameters based on differentially expressed metabolites. In leaves, the top seven metabolites with the highest interaction degree

were fructose 6-phosphate, glucose 6-phosphate, ribitol, xylitol, arabitol, lysine, and isopentenyl pyrophosphate (Fig. 5f). In roots, the top four metabolites with the highest interaction degree were citrate, lyxose, malate, and mannose 6-phosphate (Fig. 5e).

Metabolic network diagrams were constructed to visualize the changes in metabolites highly correlated with PS MNPs diameter size (Fig. 6). The metabolites identified in leaves were primarily positively correlated with PS MNPs diameters and were enriched in glycolysis/gluconeogenesis, pentose and glucuronate interconversions (Fig. 6a). With increasing PS MNPs diameters, rice progressively activated the pentose and glucuronate interconversion pathway in leaves, as evidenced by the rising relative abundances of xylitol, D-arabitol, ribitol, and D-glucuronate. However, this upward trend plateaued when PS MNPs diameters exceeded 800 nm. The accumulation of sugars alcohols enhanced plant stress tolerance and played a key role in coping with PS MNPs stress (Ozturk et al., 2021). Under exposure to 100 nm polyamide (PA), the pentose and glucuronate interconversion pathway in rice was upregulated, with increased relative abundance of D-arabinose (Yang et al., 2024). Similarly, Huang et al. (D. Huang et al., 2023) reported that when cucumber (*Cucumis sativus*) leaves were exposed to diameters of 50–100 nm polystyrene (PS) particles three weeks, the pentose and glucuronate interconversion pathway was upregulated, indicating that carbohydrate metabolic reprogramming enhanced the plant's tolerance to PS-induced stress.

The relative abundance of upstream glycolysis/gluconeogenesis metabolites D-glucose-6P and D-fructose-6P were negatively correlated with PS MNPs diameters, while the downstream metabolites glycerate-3P and glycerate-2P showed a positive

correlation. When the PS MNPs diameters exceeded 800 nm, the abundance of D-glucose-6P and D-fructose-6P significantly decreased, while the abundance of glycerate-3P and glycerate-2P also declined, revealing a contradiction between the relative abundances of upstream and downstream metabolites. Fructose-bisphosphate aldolase (FBA), a key enzyme in glycolysis/gluconeogenesis, catalyzes the cleavage of fructose-1,6-bisphosphate into two three-carbon molecules, including glyceraldehyde-3-phosphate (GAP) and dihydroxyacetone phosphate. GAP was further converted into glycerate-3P (3-PGA) and glycerate-2P (2-PGA) through enzymatic reactions, making FBA a critical link between upstream metabolites (D-glucose-6P, D-fructose-6P) and downstream metabolites (glycerate-3P, glycerate-2P). Under PS MNPs exposure, FBA activity gradually decreased with increasing PS MNPs diameters (Fig. S18). Under exposure to PS $\geq$ 800 nm, FBA activity in rice leaves was suppressed by 50.92% at 1 μ g/mL and 45.43% at 10 μ g/mL (1000 nm). The reduced availability of upstream substrates constrained glycolytic flux, and the concomitant decline in FBA activity represented a passive response to substrate scarcity. When FBA activity declined, the levels of 3-phosphoglycerate (3-PGA) and 2-phosphoglycerate (2-PGA) increased, suggesting that their accumulation was primarily due to inhibited downstream utilization rather than impaired production. Prolonged accumulation of 3-PGA and 2-PGA could have exerted feedback inhibition on upstream phosphatase enzymes, further weakening the phosphorylation-dependent breakdown of photosynthetic products. This, in turn, indirectly reduced the efficiency of carbohydrate catabolism and promotes sugars accumulation. In one study, 14-day exposure to 100 nm PS regulated aldolase

activity in wheat leaves and roots, restructuring carbohydrate metabolism to mitigate the adverse effects of PS on wheat (S. Li et al., 2021). In rice leaves, the relative abundance of succinate, a key TCA cycle intermediate, significantly increased under exposure to PS MNPs larger than 100 nm, reflecting heightened energy demands. Lysine is the first limiting essential amino acid in gramineous crops (Galili, 2002). With increasing particle diameters of PS MPs, the lysine content in rice leaves gradually increased, whereas the glutamic acid decreased progressively (Fig. 5, Fig. S16). Glutamic acid is a key precursor for glutelin synthesis and one of its most abundant amino acids. It also played a crucial role in antioxidant defense by participating in the γ -aminobutyric acid (GABA) metabolic pathway, thereby indirectly reducing the accumulation of ROS in rice. Under exposure to $PS \leq 100$ nm, although glutamic acid content in leaves remained relatively high, the glutelin content in grains was comparatively low, suggesting that glutamic acid was more involved in antioxidant processes and less available for protein synthesis. As particle diameters increased, glutamic acid content gradually decreased. However, enhanced defense responses at larger particle diameters might have redirected more glutamic acid toward glutelin synthesis, resulting in a moderate increase in glutelin content.

The metabolites identified in roots were primarily negatively correlated with PS MNPs diameters, and the DEM were mainly enriched in fructose and mannose metabolism (Fig. 6b). Relative abundances of several metabolites in the fructose and mannose metabolism pathway, such as L-sorbose, D-sorbitol, D-mannitol, and D-mannose-6P, gradually decrease with increasing PS MNPs diameters. When PS MNPs

diameters exceeded 100 nm, the decline in fructose and mannose metabolism became more significant in roots, potentially reducing the capacity to resist osmotic stress and disrupting osmotic balance.

4. Conclusion

This study demonstrated that PS MNPs deteriorated the protein nutrition in rice grains and exhibited a size-dependent effect by affecting metabolism. PS MNPs exposure induced a significant decline of glutenin content in rice grains, especially for $PS \leq 100$ nm exposure. As the PS MNPs diameters increased, the extent of glutenin depletion decreased, while starch and sugars accumulation progressively rose. Hydroponic experiment demonstrated that PS MNPs were absorbed by rice roots, and it primarily accumulated in xylem and phloem. Driven by transpiration pull, PS was transported upward, potentially affecting rice physiology, metabolism, and grain quality. The content of starch, soluble sugars, xylitol, D-arabitol, and ribitol were increased with the particle diameter of PS MNPs in the range of 80–800 nm. Simultaneously, the TCA cycle in leaves was progressively activated, and glycolysis/gluconeogenesis was modulated, with the significant inhibition of the key enzyme FBA. Compared to leaves, roots exhibited a more pronounced PS-induced stress response. Notably, $PS \leq 100$ nm imposed the highest biotoxicity and the most detrimental impact on grain quality, whereas $PS \geq 800$ nm triggered an enhanced defense response, improving rice resilience to external stress. Polystyrene exhibited high chemical stability (Ho et al., 2018) and was unlikely to degrade within rice tissues, thus the observed impacts on rice were primarily attributed to the physical properties of MNPs, such as particle diameter, and

MPs of varied composition might have exerted comparable effects on plants. Therefore, special attention should be paid to the potential risks posed by small-diameters (≤ 100 nm) MNPs exposure towards rice.

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Xiangrui Miao: Conceptualization, Data curation, Investigation, Methodology, Software, Visualization, Writing – original draft. **Wei wang:** Writing – review & editing, Supervision, Funding acquisition. **Lizhong Zhu:** Writing – review & editing, Funding acquisition, Project administration, Resources, Supervision.

Declaration of Competing Interest

The authors declare that they have no know competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Data availability

Data will be made available on request.

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Appendix A. Supporting information

Supplementary data associated with this article includes 7 texts, 10 tables and 18 figures.

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Journal Pre-proof

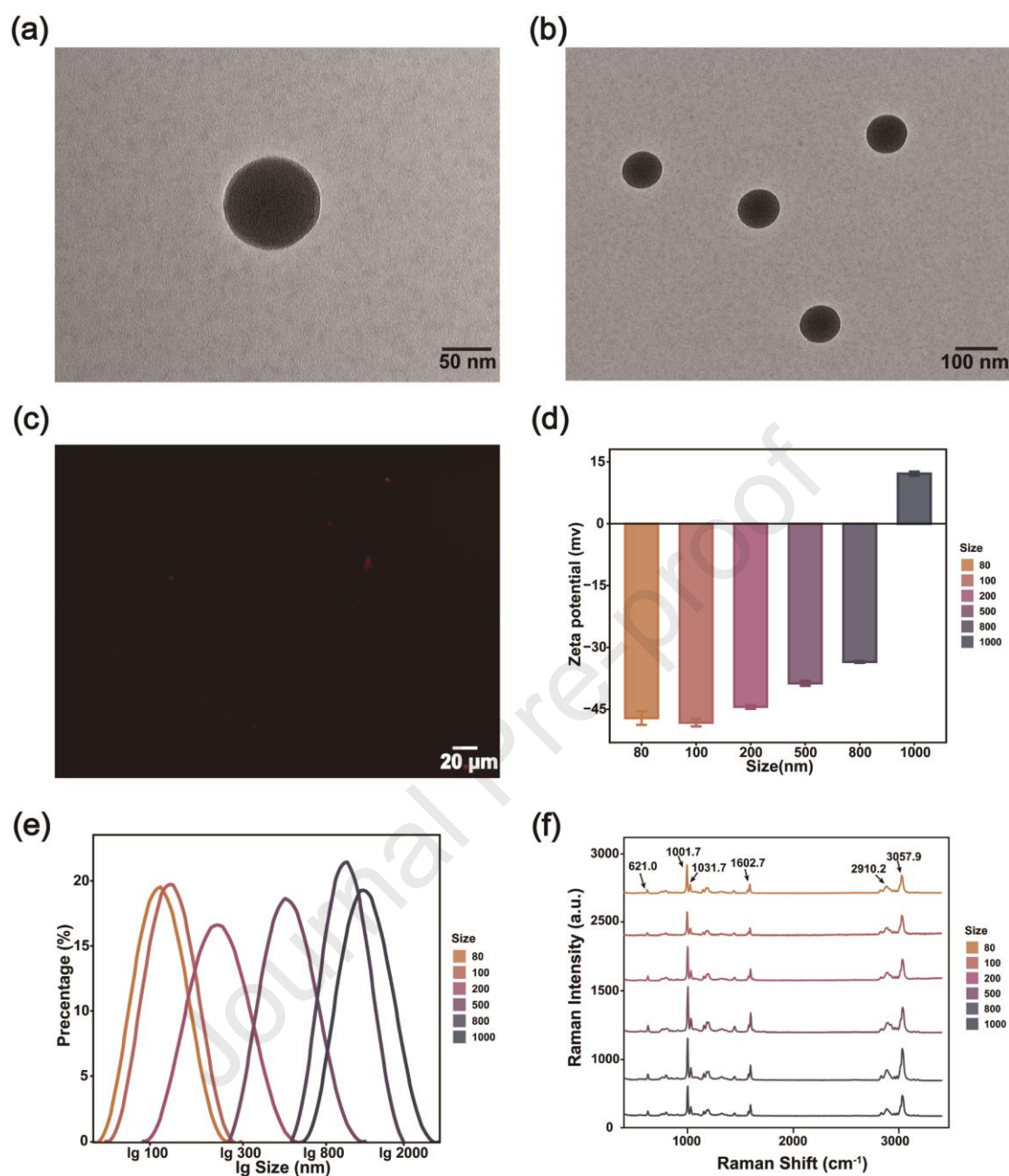


Figure 1. Analytical characterization analysis of polystyrene microplastics, including TEM morphological analysis of 80 nm polystyrene at scale bar of 50 nm (a) and 100 nm (b), fluorescence images at scale bar of 20 μm (c), zeta potential in pure water at 25°C and pH 7.0 (d), the hydrodynamic diameter in pure water at 25°C and pH 7.0 (e), and Raman spectra of polystyrene (f).

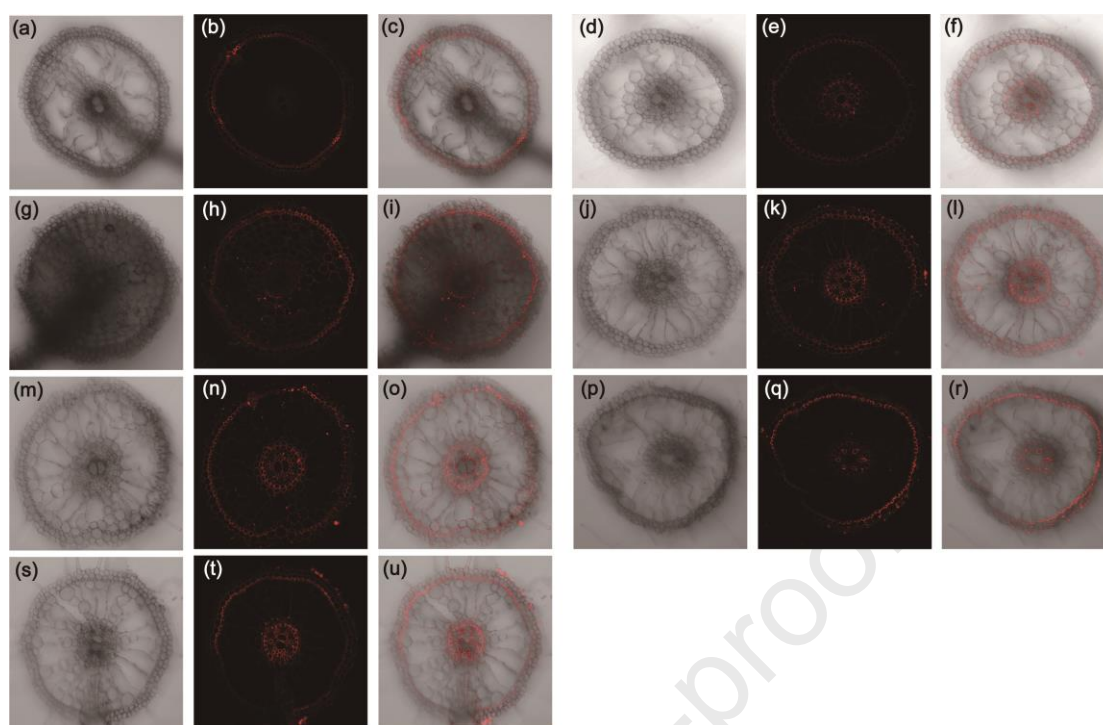


Figure 2. Confocal characterization images of rice root cross-sections after 14 days of hydroponic exposure to 1 $\mu\text{g/mL}$ PS MNPs in the groups of control group (a–c), 80 nm (d–f), 100 nm (g–i), 200 nm (j–l), 500 nm (m–o), 800 nm (p–r), 1000 nm (s–u). Every three images form a group, arranged from left to right as: bright field, fluorescence, and merged.

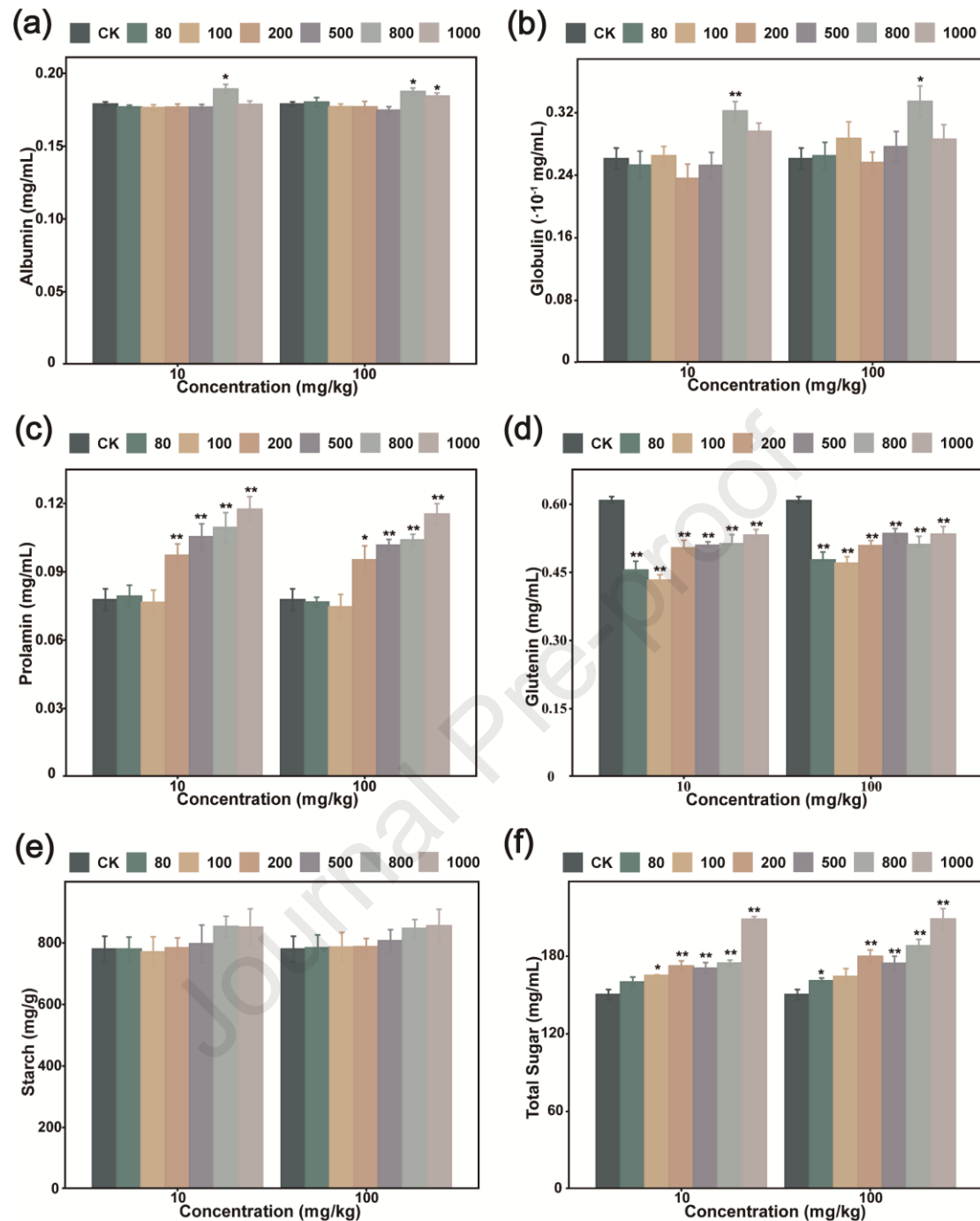


Figure 3. Content of albumin (a), globulin (b), prolamin (c), glutelin (d), starch (e) and total sugar(f) in rice grains after 180 days of exposure to polystyrene microplastics. For a given PS particle diameter and concentration, significant differences between the control and treatment groups (intergroup comparison) are marked with * ($P < 0.05$) and ** ($P < 0.01$).

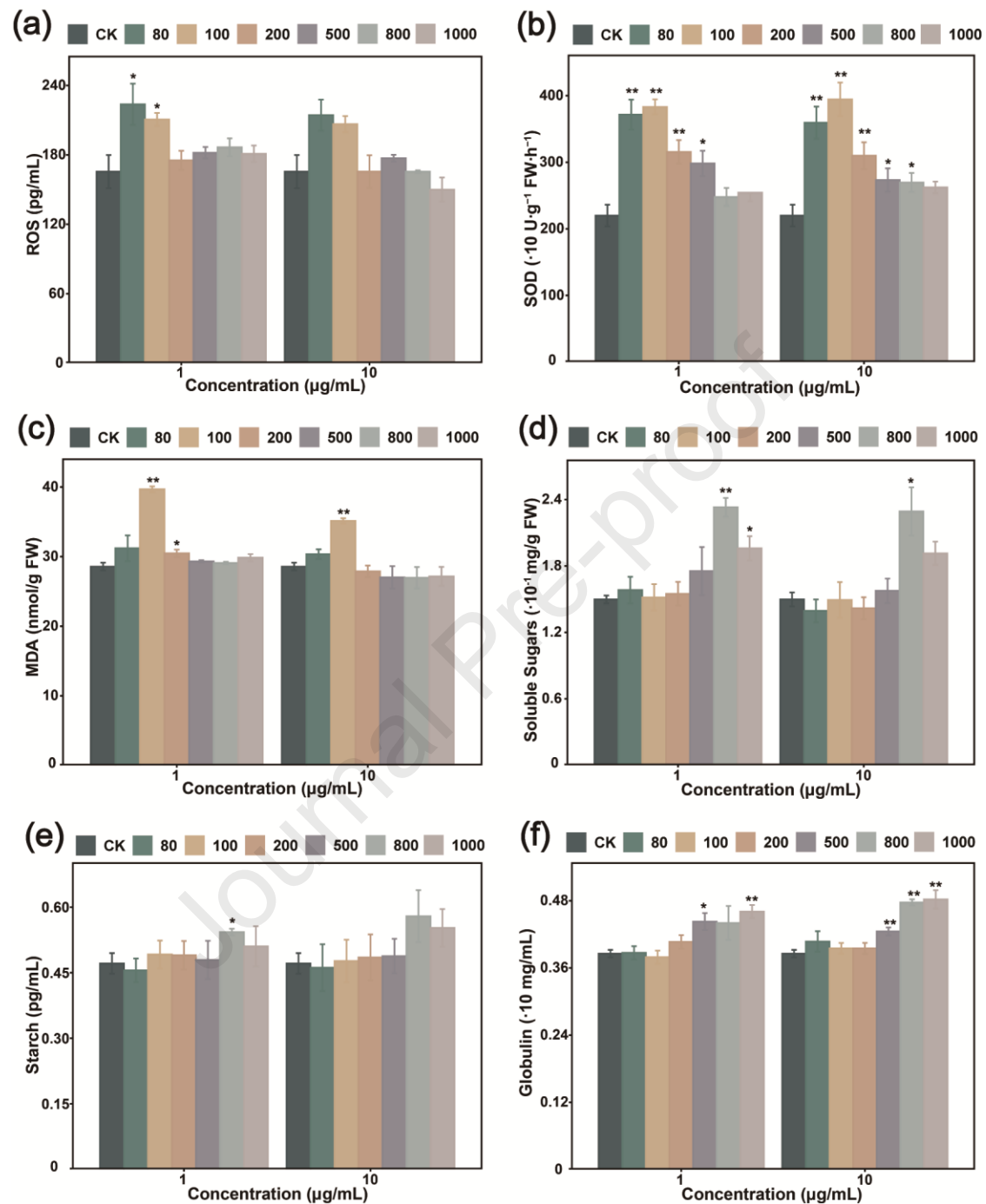


Figure 4. Physicochemical parameters and components of rice leaves after 14 days of exposure to PS MNPs of different particle diameters (80, 100, 200, 500, 800, and 1000 nm) at concentrations of 1 µg/mL and 10 µg/mL, including ROS content (a), SOD enzyme activity (b), MDA content (c), soluble sugars content (d), starch content (e),

globulin content (f). For a given PS MNPs diameter and concentration, significant differences between the control and treatment groups (intergroup comparison) are marked with * ($P < 0.05$) and ** ($P < 0.01$).

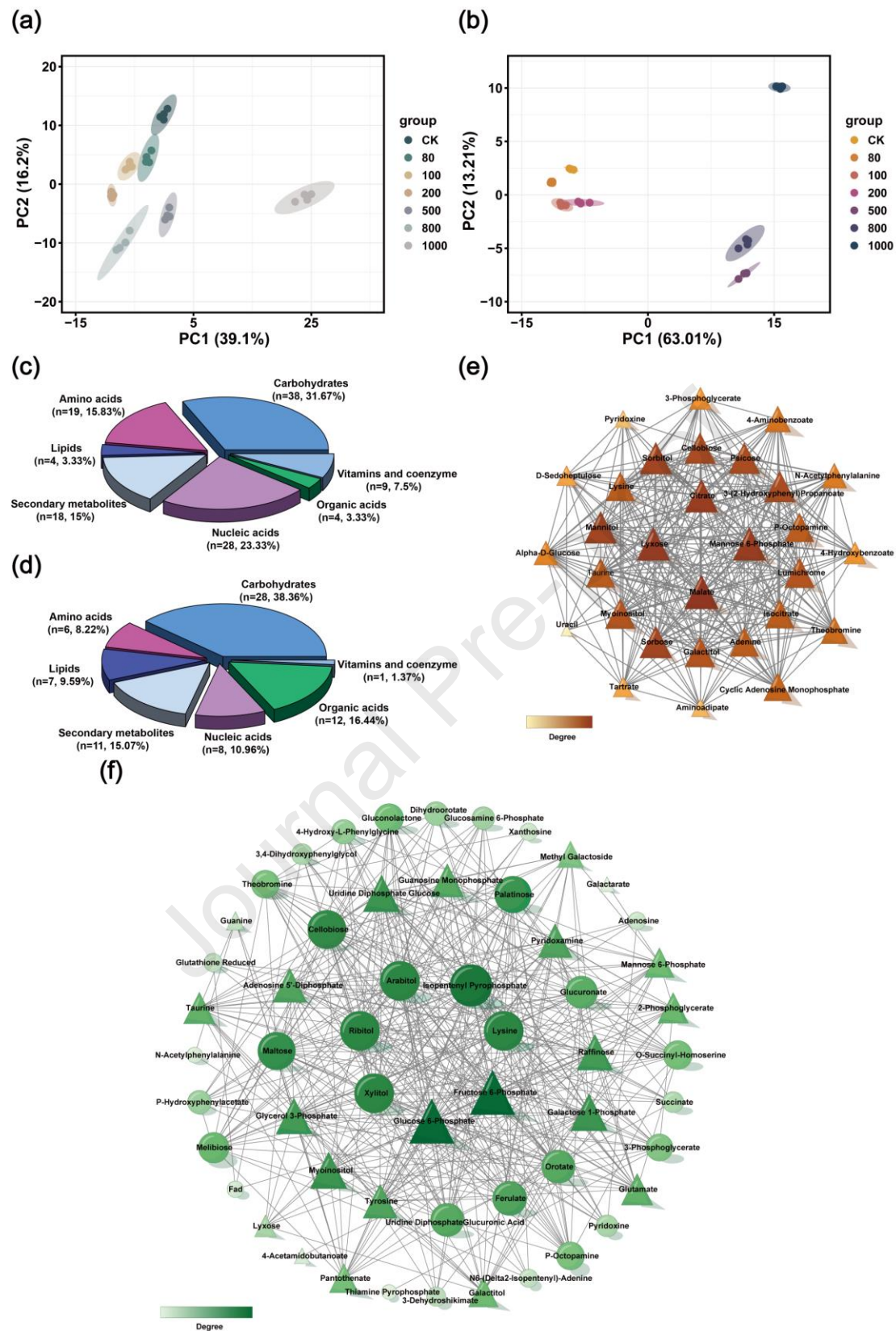
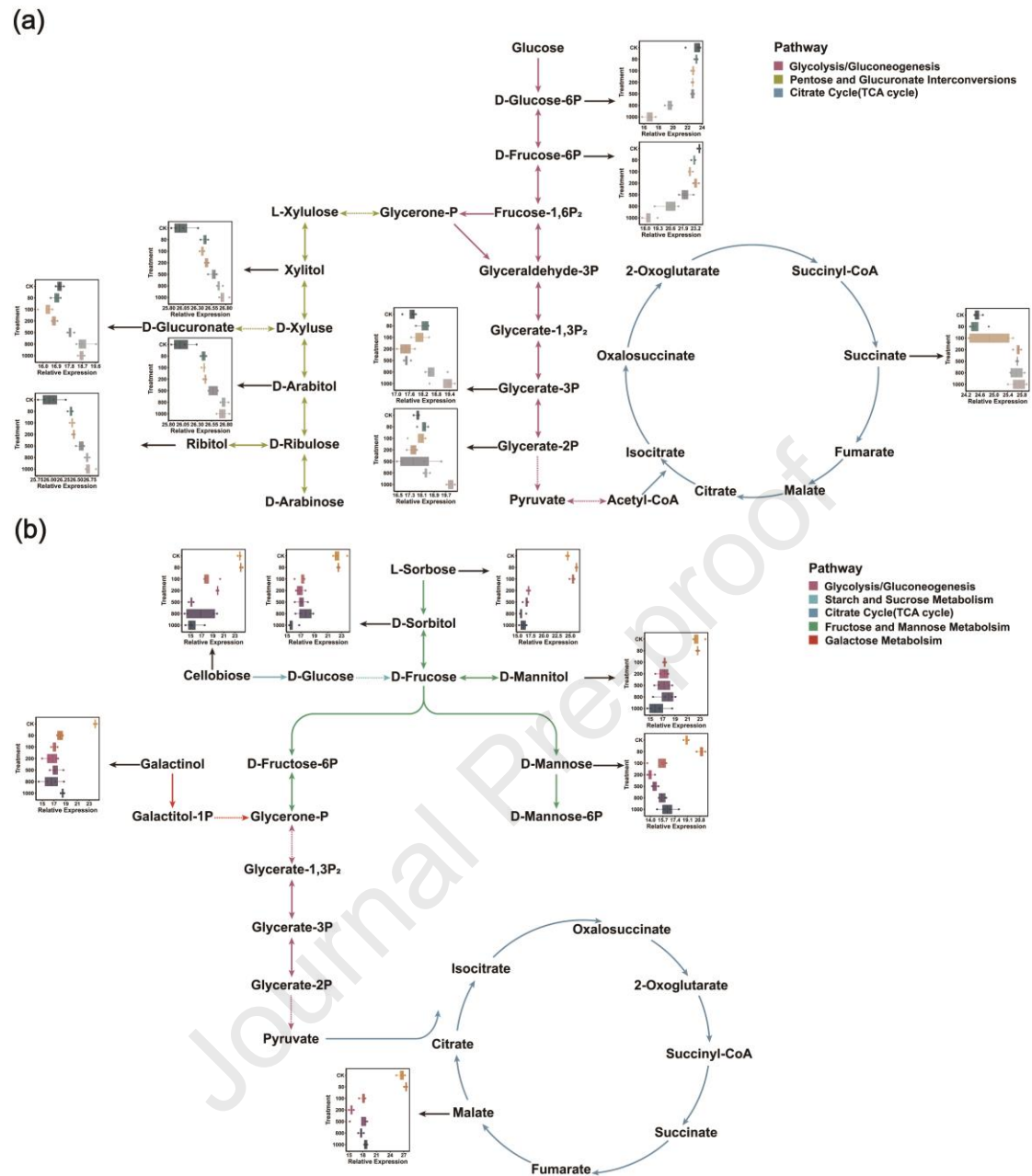
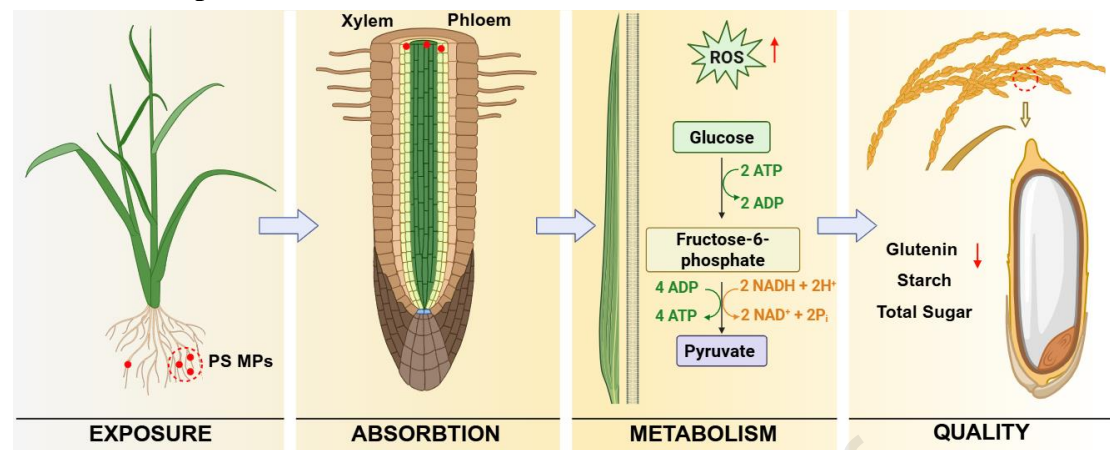


Figure 5. Metabolic analysis in rice after 14 days of exposure to PS of different particle diameters (80, 100, 200, 500, 800, and 1000 nm) at concentrations of 1 $\mu\text{g/mL}$,

including PCA analysis of metabolic differences between PS-exposed and control groups in leaves (a), and roots (b), categorization of differentially expressed metabolites in rice leaves (c) and (d) roots, and metabolites in leaves (f) and roots (e) associated with PS exposure size were identified using weighted gene co-expression network analysis (WGCNA). Darker colors and larger nodes indicate stronger interaction degree; circles represent positive correlations, and triangles represent negative correlations with PS MNPs exposure diameter. The closer to the center of the circle, the higher the degree of interaction with other metabolites.



Abstract Graphics



(Created with bioRender.com)

Highlights:

PS MPs size ≤ 100 nm induce ROS accumulation in rice and lead to reduced glutenin content in the grains.

The glycolysis/gluconeogenesis, as well as pentose and glucuronate interconversion pathway, is strongly associated with the particle size of PS MPs exposure.

Rice upregulates carbohydrate metabolism, promoting sugar accumulation to counter PS MPs stress.

Declaration of interests

☒ The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

☐ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: