



Article In Vitro Interactions of TiO₂ Nanoparticles with Earthworm Coelomocytes: Immunotoxicity Assessment

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Abstract: Titanium dioxide nanoparticles (TiO₂ NPs) are manufactured worldwide. Once they arrive in the soil environment, they can endanger living organisms. Hence, monitoring and assessing the effects of these nanoparticles is required. We focus on the *Eisenia andrei* earthworm immune cells exposed to sublethal concentrations of TiO₂ NPs (1, 10, and 100 μ g/mL) for 2, 6, and 24 h. TiO₂ NPs at all concentrations did not affect cell viability. Further, TiO₂ NPs did not cause changes in reactive oxygen species (ROS) production, malondialdehyde (MDA) production, and phagocytic activity. Similarly, they did not elicit DNA damage. Overall, we did not detect any toxic effects of TiO₂ NPs at the cellular level. At the gene expression level, slight changes were detected. Metallothionein, fetidin/lysenin, lumbricin and MEK kinase I were upregulated in coelomocytes after exposure to 10 μ g/mL TiO₂ NPs for 6 h. Antioxidant enzyme expression was similar in exposed and control cells. TiO₂ NPs were detected on coelomocyte membranes. However, our results do not show any strong effects of these nanoparticles on coelomocytes at both the cellular and molecular levels.

Keywords: earthworm; coelomocyte; TiO₂ nanoparticles; reactive oxygen species; innate immunity; lipid peroxidation; alkaline comet assay; phagocytosis; apoptosis; gene expression

1. Introduction

Titanium dioxide nanoparticles (TiO₂ NPs) are commonly used in different industries because of their physico-chemical properties. TiO₂ NPs have photocatalytic properties, protect against UV radiation, are used as semiconductors, etc. These nanoparticles are used, e.g., in cosmetics, food industry, paints, ceramics, devices development, and the agriculture industry [1–3]. In the last decade, TiO₂ NPs have been used in wastewater treatment plants for their ability to degrade some organic pollutants [1]. Thus, TiO₂ NPs reach the soil system from different sources including sludge, nanofertilizers, and nanopesticides. These nanoparticles then interact with the soil biota. It is therefore very important to assess the potential risk of TiO₂ NPs to soil organisms.

Earthworms are dominant soil invertebrate animals. They possess a strong immune system because of their permanent contact with soil bacteria, viruses, and fungi. Defense mechanisms are used in earthworm protection against soil pollutants including nanoparticles. Earthworms *Eisenia andrei* and *E. fetida* are used as model organisms to monitor ecotoxicity according to OECD guidelines [4–6]. TiO₂ NPs do not affect earthworm viability and growth [2,3]. In some cases, reproductive inhibition was observed [7].



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Copyright: © 2021 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Further, these nanoparticles can induce, e.g., oxidative stress, DNA damage, apoptosis, and affect gene expression [3]. Earthworm cellular defense mechanisms are based on coelomocytes present in the coelomic fluid. Coelomocytes can be divided into free chloragogen cells called eleocytes, with a mainly nutritive function, and amoebocytes, which are the immune effector cells [8]. Amoebocytes can be further divided into granular (GA) and hyaline (HA) amoebocytes.

Various nanoparticles were described to impair earthworm defense mechanisms. Hayashi et al. showed that Ag NPs altered the expression of some genes involved in coelomocyte oxidative stress and immune reactions [9]. Further, Ag nanowires detected on coelomocyte membranes increased intracellular esterase activity [10]. ZnO NPs were internalized by coelomocytes, with consequent DNA damage [11]. However, similar mechanisms were not described for TiO₂ NPs in earthworms. TiO₂ NPs cause significant mitochondrial dysfunction by increasing mitochondrial ROS levels and decreasing ATP generation in macrophages. Moreover, TiO₂ NPs exposure activated inflammatory responses and attenuated macrophage phagocytic function [12]. TiO₂ NPs interacted with sea urchin immune cells and increased the antioxidant metabolic pathway in vitro [13]. In earthworms, only increased apoptosis was observed following TiO₂ nanocomposites exposure [7,14–16].

A compromised immune system may result in a decreased reproductive rate and increased mortality of earthworms. Thus, nanoparticle toxicity risk assessment is extremely important, as the adverse health effects remain poorly characterized for many nanomaterials. We aimed to assess the potentially dangerous impact of TiO_2 NPs exposure on earthworms' cellular function, including the immune responses to harmful stimuli.

E. andrei coelomocytes were exposed to 1, 10, and 100 μ g/mL of TiO₂ NPs for 2, 6, and 24 h in vitro. After exposure, viability, oxidative stress (reactive oxygen species and malondialdehyde production), immune functions (phagocytosis), and genotoxicity (DNA damage) were assessed. Further, electron microscopy (transmission and scanning) enabled TiO₂ NPs localization on the cell surface. Gene expression changes were also followed to better understand the underlying cellular mechanisms.

2. Materials and Methods

2.1. Animal Handling, Sample Collection, and Culture Medium Preparation

Clitelate, adult *Eisenia andrei* earthworms were obtained from the laboratory compost breeding. Earthworms were first kept on moist filter paper for 48 h to depurate their guts. Coelomocytes were harvested by applying 2 mL of extrusion buffer (5.37 mM EDTA (Sigma-Aldrich, Steinheim, Germany); 50.4 mM guaiacol glyceryl ether (GGE; Sigma-Aldrich, Steinheim, Germany) in Lumbricus Balanced Salt Solution (LBSS; [17]) per earthworm for 2 min. The cells were then centrifuged and washed twice in LBSS ($200 \times g$, 4 °C, 10 min). Subsequently, cells were counted and diluted to 10⁶ cells/well for scanning electron microscopy (SEM) and lipid peroxidation assessment. 1 × 10⁵ cells/well, 2 × 10⁵ cells/well, and 3 × 10⁵ cells/well were used for the ROS production analysis, apoptosis detection, and phagocytosis assay, respectively.

RPMI 1640 culture medium supplemented with 5% heat-inactivated fetal bovine serum (FBS; Life technologies, Carlsbad, USA), 1 M HEPES (4-(2-hydroxyethyl)-1-piperazineethane sulfonic acid; pH 7.0–7.6, Sigma-Aldrich; Gillingham, UK), 100 mM sodium pyruvate (Sigma-Aldrich, Steinheim, Germany), 100 mg/mL gentamycin (Corning, Manassas, VA, USA), and antibiotic–antimycotic solution (Sigma-Aldrich, Steinheim, Germany) was diluted with autoclaved MilliQ-water to 60% (v/v) to obtain R-RPMI 1640 medium [18]. Subsequently, TiO₂ NPs were dispersed in R-RPMI 1640 medium and incubated with cells in darkness at 20 °C for 2, 6, and 24 h in triplicate.

2.2. TiO₂ NPs Characterization

Aeroxide TiO_2 P25 nanoparticles (irregular and semi-spherical shape; mexoporous NPs, anatase, and rutile 4:1; primary size between 10 and 65 nm) were purchased from

Evonik Degussa (Essen, Germany). TiO₂ NPs were previously characterized in several aqueous solutions, as described by Brunelli et al. [19]. Nanoparticle physico-chemical properties were determined by ZetaSizer Ultra (Panalytical Malvern; Malvern, UK), transmission electron microscope (TEM), and TECAN 200 Pro plate reader. Powder TiO₂ NPs were weighed and dispersed in distilled water. Then, diluted TiO₂ NPs were vortexed thoroughly for 5 min prior to further dilution [20]. TiO₂ NPs were diluted either in R-RPMI 1640 medium or distilled water to a concentration of 1, 10, and 100 μ g/mL, and incubated for 2, 6, and 24 h. Experiments were carried out in triplicate. Culture medium and distilled water without NPs were used as negative controls.

2.3. Electron Microscopy Analyses

2.3.1. Cell Preparation

Coelomocytes were exposed to 1, 10, and 100 μ g/mL TiO₂ NPs for 2, 6, and 24 h. Cell viability was measured by propidium iodide (PI; 1 μ g/mL) staining using flow cytometer. Then, samples were collected and fixation solution (5% glutaraldehyde in PBS) was added in a 1:1 ratio (*v*:*v*). Fixed cells were shaken gently for 15 min and kept overnight at 4 °C.

2.3.2. Scanning Electron Microscopy (SEM)

For SEM, fixed cells were washed with LBSS buffer three times at room temperature for 20 min, and centrifuged at $150 \times g$ for 10 min. Then, they were allowed to adhere onto poly-L-lysine coated round 13 mm Thermanox Plastic Coverslips (Nunc, Thermo Fisher Scientific; Roskilde, Denmark) overnight at 4 °C. The coverslips with attached cells were washed with ddH₂O and fixed with 1% OsO₄ for one hour at room temperature. The coverslips were then washed three times for 20 min, dehydrated through an alcohol series (25, 50, 75, 90, 96, and 100%), and were critical-point dried from liquid CO_2 in a K850 Critical Point Dryer (Quorum Technologies Ltd., Ringmer, UK). The dried coverslips were sputter-coated using a high-resolution Turbo-Pumped Sputter Coater Q150T (Quorum Technologies Ltd., Ringmer, UK) with 3 nm of platinum. Alternatively, for EDS microanalysis, the samples were coated with 10 nm of silver or 5 nm of carbon. The final samples were examined in a FEI Nova NanoSEM scanning electron microscope (FEI, Brno, Czech Republic) at 5 kV using CBS and TLD detectors. An electron beam deceleration [21] mode of the Nova NanoSEM scanning electron microscope performed at a StageBias of 883.845 V and accelerating voltage of 5 kV was used for high-resolution imaging. The EDS microanalysis was performed at 15 kV using an Ametek[®] EDAX Octane Plus SDD detector and TEAMTM EDS Analysis Systems (AMETEK B. V.; Tilburg, The Netherlands).

2.3.3. Transmission Electron Microscopy (TEM)

For TEM, a TiO₂ NPs suspension (500 μ g/mL; 5 μ L) was applied onto glow-dischargeactivated [22] carbon-coated 400-mesh copper grids (G400, SPI Supplies, Structure Probe, Inc., West Chester, PA, USA). Nanoparticles were sedimented for 1 min and the remaining solution was then blotted with filter paper and the grids were air-dried. A Philips CM100 electron microscope (Philips EO, Eindhoven, The Netherlands; Thermo Fisher Scientific) equipped with a Veleta slow-scan CCD camera (EMSIS GmbH, Muenster, Germany) was used to examine the grids. TEM images were processed in the proprietary iTEM software (EMSIS GmbH, Muenster, Germany).

2.4. Flow Cytometry Assays

Coelomocytes were incubated with TiO_2 NPs (1, 10, and 100 µg/mL) for 2, 6, and 24 h. Cells were then treated as described below and analyzed with a laser scanning flow cytometer. Through flow cytometry, coelomocytes were subdivided into eleocytes, granular (GA), and hyaline amoebocytes (HA). The coelomocytes subset detection was based on the cell size (FSC) and the cell inner complexity/granularity (SSC). Cell viability was assessed for every assay. All flow cytometry assays were performed by three independent experiments

with three replicates per each treatment and time interval. The minimum collected events were 1000 per population. Event counts per each gate were calculated by Flowjo (9.9.4 version, BD Biosciences, San Jose, CA, USA). In each flow cytometry assay, coelomocytes were exposed to H_2O_2 as a positive control (Sigma-Aldrich, Steinheim, Germany; 10 mM H_2O_2 for 30 min incubation for apoptosis and phagocytosis, and 1 mM H_2O_2 for ROS production assessment). Controls with and without PI (1 mg/L; Sigma-Aldrich, Steinheim, Germany) were included in each experiment. Further, control analysis of 1, 10, and 100 μ g/mL TiO₂ NPs incubated with or without cells for 2, 6, and 24 h were performed (Figure S1).

For ROS production determination, 20.6 μ M 2',7'-dichlorofluorescin diacetate (DCF-DA; Sigma-Aldrich, Steinheim, Germany) was added to the washed cell suspension (LBSS, 200× *g*, 4 °C, 10 min) for 15 min in darkness. Subsequently, the cell suspension was washed twice with LBSS (200× *g*, 4 °C, 10 min) and stained with PI.

To detect the apoptotic process a cell suspension was washed twice with Annexin V buffer ($200 \times g$, 4 °C, 10 min; 0.01 M HEPES (pH 7.4), 0.14 M NaCl, and 2.5 mM CaCl₂ solution), and subsequently stained with 5 µL of Alexa Fluor 647-Annexin V (15 min in darkness; Thermo Fisher Scientific, Eugene, OR, USA). PI was then added to the cell suspension and measured by flow cytometry. The apoptosis % represented the apoptotic cell number out of each subpopulation. The necrosis % represented the necrotic cell number out of each subpopulation.

The phagocytosis assay was performed using latex beads (Fluoresbrite[®] Plain YG; 1 µm microspheres diameter; Polysciencies Inc., Warrington, PA, USA) added to the incubation plates in a 1:100 ratio (cells:beads) and kept in darkness at 17 °C for 18 h. Then, cell suspensions were washed twice with LBSS ($200 \times g$, 4 °C, 10 min), stained with PI, and analyzed by flow cytometry. The % phagocytic activity was determined by the % of alive cells, which were able to engulf at least one bead out of each subpopulation. Each experiment included samples with NPs dispersed in the medium in order to detect effects exerted by NPs alone.

2.5. Malondialdehyde (MDA) Production and Alkaline Comet Assay

Coelomocytes were incubated with TiO₂ NPs (10 and 100 μ g/mL) or CuSO₄ (100 μ g/mL; positive control) for 2, 6, and 24 h. Afterward, cell suspensions were collected and MDA production was measured. MDA production was detected by high-performance liquid chromatography with fluorescence detection (HPLC/FLD) using derivatized MDA-TBA2 [23]. MDA analysis was performed in three independent experiments with 3 replicates for each treatment and time interval.

For the alkaline comet assay, 1.5×10^4 cells exposed to 1, 10, and 100 µg/mL TiO₂ NPs for 2, 6, and 24 h were mixed with 2% 2-hydroxyethyl agarose (Sigma-Aldrich, Steinheim, Germany) at 37 °C. Glass slides containing agarose with cells were kept at 4 °C for 10 min. Subsequently, samples were incubated for 2 h in lysis buffer (2.5 M NaCl, 10 mM Tris-HCl, 100 mM EDTA, 1% Triton X-100, pH 10). Then, slides were immersed three times in unwinding buffer (0.03 M NaOH, 2 mM EDTA, pH 12.7) for 20 min. Gel electrophoresis was carried out at 24 V, 300 mA for 25 min. Subsequently, slides were rinsed with neutralizing buffer (0.4 M Tris, pH 7.5) and stained with PI (3 µg/mL) for 20 min. The excess dye was removed with distilled water (5 min). Then, samples were stored in humidified chambers until the analysis by LUCIA Comet Assay software. One hundred cells per replicate of each treatment and time interval were analyzed, and the mean of DNA content in 100 comet tails (%) was calculated as a parameter of DNA damage. Positive control (100 mM H₂O₂; 30 min incubation; Sigma-Aldrich, Steinheim, Germany) was included with the assay. The comet assay was repeated in three independent experiments with three replicates for each treatment and time interval.

2.6. mRNA Levels Quantification

Cells were incubated with TiO_2 NPs (1 and 10 μ g/mL) for 2, 6, and 24 h. Cellular RNA was isolated using the RNAqueous[®]-Micro Kit (Invitrogen, Vilnius, Lithuania).

RNA (500 ng) was reverse-transcribed with the Oligo(dT)12–18 primer and Superscript IV Reverse Transcriptase (Life Technologies). Non-RT controls were included to show the elimination of gDNA contamination.

Quantitative PCR (CFX96 Touch™ Real-Time PCR detection System, Bio-Rad) was performed to detect changes in mRNA levels encoding proteins participating in metal detoxification (metallothionein, phytochelatin), oxidative stress (manganese superoxide dismutase, Mn-SOD; copper-zinc-superoxide dismutase CuZn-SOD; catalase), immunity (endothelial monocyte-activating polypeptide II, EMAPII; fetidin/lysenin, and lumbricin), and signal transduction (MEK kinase I, MEKK I; and protein kinase C I, PKC I). Sequences of primers used in qPCR assays are referred in Table S1. The PCR reactions were performed in a 25μ L volume containing 4 μ L of cDNA (dilution 1:10, except for 1:5 dilution for SODs). The cycling parameters were similar to Roubalova et al., with slight changes [24]: 4 min at 94 °C, 35 cycles of 10 s at 94 °C, 25 s at 60 °C (at 58 °C for MEKK I, PKC I, and catalase), 35 s at 72 °C, and a final extension for 7 min at 72 °C. Gene expression changes were calculated according to the $2-\Delta\Delta$ CT (Livak) method. Two reference genes (RPL13, RPL17) were selected as internal controls for gene expression normalization. Non-template control was included in each experiment. The fold change in the mRNA level was related to the change of the corresponding controls. The results were expressed as the mean \pm SEM of the values. mRNA levels quantification was performed by three independent experiments with duplicates per each treatment and time interval.

2.7. Statistical Analyses

Statistical analyses were performed using GraphPad Prism (8.3.1 version, San Diego, CA, USA). Flow cytometry assays, lipid peroxidation, alkaline comet assay, and gene expression were analyzed by two-way ANOVA with Bonferroni post-test.

3. Results

3.1. TiO₂ NPs Characterization

TiO₂ NPs were dispersed and stabilized in distilled water and in R-RPMI 1640 culture medium evenly. However, differences in NPs characteristics were observed between both mediums along the exposure time (2, 6, and 24 h) (Table 1). In the UV/Vis spectra, NPs exerted a similar wavelength range: 300–370 nm for distilled water; 320–380 nm for R-RPMI 1640 medium. Although NPs absorbed similar UV/Vis wavelengths, differences were observed in the hydrodynamic size distribution. TiO₂ NPs dispersed in R-RPMI 1640 medium were not stabilized and tended to aggregate. The aggregation was detected between 6 and 24 h of incubation. After 6 h, the hydrodynamic size distribution was 35.5 ± 3.94 nm, and it increased to 597 ± 447 nm after 24 h. At 2–6 h, the hydrodynamic size of TiO₂ NPs was stable (31.34 ± 1.55 to 35.5 ± 3.94 nm, respectively). In distilled water, the hydrodynamic size distribution was stable between 2–24 h 581 ± 23.30 and 480 ± 64.3 nm, respectively. Regarding zeta potential, TiO₂ NPs dispersed in both distilled water and R-RPMI 1640 did not change significantly over time (Table 1).

Table 1. Characterization of 100 μg/mL TiO₂ nanoparticles (NPs) suspension in milliQ water and R-RPMI 1640 medium.

	UV/Vis (nm) ^a			Z-Avg. (nm) ^b			ζ (mV) ^c		
	2 h	6 h	24 h	2 h	6 h	24 h	2 h	6 h	24 h
Distilled water	300-370	300-370	300-370	581 ± 23.30	570 ± 2.75	480 ± 64.3	-26.8 ± 2.99	-31.7 ± 0.921	-32.9 ± 2.59
R-RPMI 1640 medium	320-380	320-380	320-380	31.34 ± 1.55	35.5 ± 3.94	597 ± 447	-16.9 ± 0.60	-7.87 ± 0.631	-5.94 ± 0.45

(a) ultraviolet-visible = UV/Vis spectra absorbance (nm), (b) Z-Avg = Hydrodynamic size determined by multi-angle dynamic light scattering (MADLS), and (c) ζ = zeta potential values are expressed as mean of 3 measurements ± SD.

3.2. Electron Microscopy

The TiO₂ NPs size given by the manufacturer was 10–65 nm. However, we were not able to verify this information because of a great aggregation of TiO₂ NPs in concentrations

detectable by TEM. According to our measurements, the nanoparticles ranged between 20 and 100 nm. The TiO_2 NPs were rode/spherical (Figure 1).



Figure 1. Transmission electron microscopy of 500 μ g/mL TiO₂ NPs clustered in distilled water. The scale bar represents 200 nm.

In coelomocytes exposed to 100 μ g/mL TiO₂ NPs, nanoparticles were observed on cell membranes by scanning electron microscopy (Figure 2). Moreover, EDS microanalysis confirmed Ti presence in nanoparticle clusters on the cell surface (Figure 3). At 10 μ g/mL TiO₂ NPs exposure, nanoparticles were also detected on the coelomocyte surface but with lower frequency. EDS microanalysis of non-treated cells is shown in Figure S2.



Figure 2. Scanning electron microscopy of coelomocytes. (**A**) cells exposed to $100 \ \mu\text{g/mL TiO}_2$ NPs for 2 h; (**B**) control cells cultured in the medium. Images recorded with B + C segments of a CBS detector at 3 kV. White arrow indicates a TiO₂ NPs cluster on sample support. Clusters of the same morphology can be seen on the cell surface (white double arrow). The scale bar represents 5 μ m.



Figure 3. EDS microanalysis of coelomocytes incubated with 100 μ g/mL TiO₂ NPs. (**A**) An image showing the area of interest taken with EDX TEAM software at 15 kV using a SED detector. The spectra collection places are marked with EDS labels. Increased charging effects caused by the non-conductive nature of Thermanox coverslips used for sample preparation deteriorated image quality. (**B**) EDS microanalysis confirmed Ti in NPs clusters found on the cell surface (e.g., EDS Spot 2 label) and also in the cluster labeled EDS spot 1. Blue arrow indicates TiO₂ NPs cluster (EDS Spot 2 label), green arrow points to the cell surface without NPs clusters (EDS Spot 4 label). Corresponding EDS spectra in matching colors are shown in B. The scale bar represents 5 μ m.

3.3. Flow Cytometry Assays

Coelomocyte subpopulations were differentiated by flow cytometry (Figure S3). Thus, the viability of HA and GA were analyzed. The eleocyte subpopulation was excluded from the results because of the interaction between their autofluorescence and the fluorescences used in the assays.

HA and GA viability (the percentage of alive cells in each subpopulation) was similar in non-treated cells and TiO_2 NPs-exposed cells. No differences in viability were observed between amoebocyte subpopulations.

We did not observe any significant changes in ROS production in HA or in GA after exposure to any of the TiO₂ NPs concentrations (Figure 4). HA population exerted two times lesser fluorescence intensity in comparison to the GA population. This suggests that HA population is less potent to produce ROS than GA population (Figure 4). Illustrative histograms of ROS production between control samples and positive control (1 mM H₂O₂), indicating a clear shift in sample fluorescence, are shown in Figure S4.



Figure 4. ROS production by hyaline (HA) and granular (GA). ROS production was measured in HA and GA after incubation with 1, 10, and 100 μ g/mL TiO₂ NPs for 2, 6, and 24 h using a cell-permeant tracer 2',7'-dichlorofluorescein diacetate (DCF-DA). Coelomocytes were also exposed to 1 mM H₂O₂ (positive control) for 30 min. The results are shown as the mean of fluorescence intensity (DCF-DA) \pm SEM of three independent experiments with 3 replicates in each. *** *p* < 0.001, and * *p* < 0.05 according to two-way ANOVA and Bonferroni post-test.

Similarly, we did not detect any significant differences in the apoptosis level between TiO_2 NPs exposed and control cells (both in HA and GA; Figures 5 and 6). In both populations, the early apoptosis percent is similar over time, while late apoptosis slightly decreased after 24 h (Figures 5 and 6). Necrosis increased along the exposure time in GA (Figure 6). However, statistically significant differences were not detected between treated and control cells. Representative distributions of the apoptotic/necrotic cell stages in GA and HA cell subpopulations are shown in Figures S5 and S6, respectively.



Figure 5. Early and late apoptosis, viability and necrosis of hyaline amoebocytes (HA). Early and late apoptosis, viability and necrosis of HA of non-treated cells, cells exposed to 1, 10, and 100 μ g/mL TiO₂ NPs after 2, 6, and 24 h. 10 mM H₂O₂ was used as positive control for 30 min exposure. The results are shown as mean (%) ± SEM of three independent experiments with 3 replicates in each. *** *p* < 0.001, and ** *p* < 0.01 according to two-way ANOVA and Bonferroni post-test.



Figure 6. Early and late apoptosis, viability and necrosis of granular amoebocytes (GA). Early and Late apoptosis, viability and necrosis of GA of non-treated cells, cells exposed to 1, 10, and 100 μ g/mL TiO₂ NPs after 2, 6, and 24 h. 10 mM H₂O₂ was used as positive control for 30 min exposure. The results are shown as mean (%) ± SEM of three independent experiments with 3 replicates in each. ** *p* < 0.01, and * *p* < 0.05 according to two-way ANOVA and Bonferroni post-test.

The viable amoebocyte phagocytic activity was measured in both amoebocyte subsets (HA and GA). Representative phagocytic activity density plots of GA and HA cell subpopulations are shown in Figures S7 and S8, respectively. The phagocytic activity was similar in both amoebocyte subpopulations (GA and HA; Figure 7). A decrease in the phagocytic activity of HA control cells and TiO₂ NPs-exposed cells occurred after 24 h (Figure 7). This slight decrease may indicate the greater sensitivity of HA to external conditions. However, phagocytic activity was not significantly affected by NPs treatment or by the incubation time. Phagocytic activity of untreated cells with and without Fluoresbrite[®] YG Plain 1µm microspheres was also compared (Figure S9).



Figure 7. Phagocytic activity of HA and GA. Phagocytic activity was measured after incubation with TiO₂ NPs (1, 10, and 100 μ g/mL) for 2, 6, and 24 h. Coelomocytes were also exposed to 10 mM H₂O₂ (positive control) for 30 min. Results are represented as the mean \pm SEM of three independent experiments with 3 replicates in each. *** *p* < 0.001, ** *p* < 0.01, and * *p* < 0.05 according to two-way ANOVA and Bonferroni post-test.

3.4. MDA and Alkaline Comet Assay

Malondialdehyde (MDA) is a lipid peroxidation subproduct, and it is therefore used as an oxidative stress biomarker in cells. We did not detect any significant increase in MDA production in cells exposed to TiO_2 NPs (10 and 100 μ g/mL) at the tested timepoints (2, 6, and 24 h; Figure 8).



Figure 8. Relative malondialdehyde (MDA) production in coelomocytes exposed to 10, 100 μ g/mL TiO₂ NPs and positive control (100 μ g/mL CuSO₄) for 2, 6, and 24 h. Values are expressed as mean (%) \pm SEM of three independent experiments each with three replicates. *** *p* < 0.001 according to two-way ANOVA and Bonferroni post-test.

The DNA damage in coelomocytes exposed to 1, 10, and 100 μ g/mL TiO₂ NPs for 2, 6, and 24 h was assessed by the alkaline comet assay. DNA damage was evaluated by the mean tail intensity (% DNA in tail) of 100 comets in each incubation. The observed DNA damage was not greater than 40% during exposure with TiO₂ NPs and the non-treated cells (Figure 9).



Figure 9. DNA damage in coelomocytes after their exposure to 1, 10, and 100 μ g/mL TiO₂ NPs for 2, 6, and 24 h. Coelomocytes were also exposed to 100 mM H₂O₂ (positive control) for 30 min. Values are expressed as the mean of DNA content in tail (%) \pm SEM of three experiment with three replicates. *** *p* < 0.001 and * *p* < 0.05 according to two-way ANOVA and Bonferroni post-test.

3.5. mRNA Levels of Detoxification, Immune, Antioxidant, and Signal Transduction Molecules

The change in mRNA levels of appropriate molecules after coelomocyte exposure to TiO_2 NPs was assessed (Table 2). Metallothioneins involved in metal detoxification were significantly upregulated in coelomocytes exposed to 1 µg/mL TiO₂ NPs for 2, 6 and 24 h, and in coelomocytes exposed to 10 µg/mL TiO₂ NPs for 6 h. Further, significant Mn-SOD downregulation was detected in coelomocytes incubated with 10 µg/mL TiO₂ NPs for 6 h. Then, fetidin/lysenin and lumbricin were upregulated upon coelomocyte exposure to 10 µg/mL TiO₂ NPs for 6 h. MEKK I upregulation after 1 µg/mL TiO₂ NPs exposure for 24 h, and PKC I downregulation after 10 µg/mL TiO₂ NPs exposure for 6 and 24 h were detected. Surprisingly, the mRNA levels of catalase and CuZn-SOD (antioxidant enzymes) were not significantly altered.

Turn (fam	Contra	TO NBs (us/mL)	Normalized Gene Expression			
Function	Gene	110_2 MFS (µg/IIIL)	2 h	6 h	24 h	
Matal JatasiGastian	Matallathianain	1	5.16 \pm 1.73 **	2.00 \pm 0.32 *	2.71 \pm 0.20 *	
Metal detoxification	Metallothionein	10	1.11 ± 0.2	1.97 \pm 0.22 **	1.00 ± 0.25	
Heavy motal detexification	Phytocholatin	1	1.38 ± 0.09	1.02 ± 0.04	1.18 ± 0.08	
neavy metal detoxincation	rnytochelatin	10	1.00 ± 0.02	0.82 ± 0.02	0.80 ± 0.01	
	NL COD	1	1.47 ± 0.12	0.85 ± 0.19	0.58 ± 0.05	
	Mn-SOD	10	0.93 ± 0.09	0.53 \pm 0.01 *	0.72 ± 0.01	
		1	0.68 ± 0.05	0.84 ± 0.22	0.98 ± 0.04	
Oxidative stress	CuZn-SOD	10	0.96 ± 0.07	0.71 ± 0.04	0.87 ± 0.01	
	Catalana	1	1.41 ± 0.19	0.87 ± 0.03	0.66 ± 0.03	
	Catalase	10	1.04 ± 0.13	0.71 ± 0.02	0.8 ± 0.2	
		1	0.90 ± 0.07	0.94 ± 0.1	0.86 ± 0.02	
	EMAP II	10	0.84 ± 0.09	1.21 ± 0.01	1.33 ± 0.20	
Immunity	Eatidin /lucanin	1	0.64 ± 0.08	0.62 ± 0.13	0.70 ± 0.04	
Innutity	retiditi/ tysetiiti	10	0.65 ± 0.05	2.20 \pm 0.2 **	0.81 ± 0.19	
	T 1	1	1.33 ± 0.05	0.75 ± 0.10	1.84 ± 0.02	
	Lumbricin	10	0.84 ± 0.10	2.10 \pm 0.43 *	1.92 ± 0.55	
		1	1.40 ± 0.19	1.47 ± 0.44	1.73 \pm 0.04 *	
Signal Transduction	MEKK I	10	1.00 ± 0.15	1.96 \pm 0.11 *	1.33 ± 0.03	
Signal Hansauction	DVCI	1	1.52 ± 0.30	1.08 ± 0.19	1.43 ± 0.06	
	rku i	10	1.10 ± 0.16	0.33 \pm 0.04 **	0.58 \pm 0.11 *	

Table 2. The mRNA levels of distinct molecules in coelomocytes exposed to 1 and $10 \,\mu g/mL \, TiO_2 \, NPs$.

Values were normalized to two reference molecules (RPL13 and RPL17). Fold changes (\pm SEM) in mRNA levels in TiO₂ NPs exposed coelomocytes are relative to the mRNA levels in control cells. Two-way ANOVA and Bonferroni post-test were performed to evaluate data significance (* p < 0.05, ** p < 0.01). mRNA levels quantification was performed by three independent experiments with duplicates per each treatment and time interval. Mn-SOD: manganese superoxide dismutase; CuZN-SOD: copper-zinc-superoxide dismutase; EMAP II: endothelial monocyte-activating polypeptide-II; MEKK I: MEK kinase I; PKC I: protein kinase C I.

4. Discussion

The physico-chemical properties of TiO₂ NPs were analyzed in R-RPMI 1640 medium to understand their behavior in cell cultures. The analyses in distilled water were performed to observe possible changes in nanoparticles behavior in the stock over time. UV/Vis spectra, hydrodynamic size, zeta potential, and TEM were used for the NPs characterization. TiO₂ NPs dispersed in distilled water showed an aggregation behavior at the greatest concentration (100 μ g/mL), and the size remained approximately the same between 2 and 24 h. The zeta potential was also stable at all TiO₂ NPs concentrations (Table 1). However, different NPs behavior was observed when dispersed in the R-RPMI 1640 culture medium. Between 2 and 6 h of incubation, changes were not observed in the size distribution, while zeta potential indicated instability (Table 1). Between 6 and 24 h, we observed a great increase in particle size distribution in comparison with previous intervals. These changes indicate that NPs were dispersed in R-RPMI 1640 medium, and they started to precipitate only after 6 h of incubation. Magdolenova and colleagues assessed the relationship between the cytotoxic effects and the dispersion of TiO₂ NPs [25]. They showed that tested cell culture medium types did not influence TiO₂ NPs dispersion. However, they observed that different dispersion protocols and the use of serum in stock solution affected nanoparticles aggregation and size distribution. Accordingly, Ji et al. showed the improvement in TiO₂ NPs dispersion upon addition of bovine serum albumin (BSA), although the dispersion also depended on cell culture media phosphate concentration [26]. TiO₂ NPs tended to aggregate in R-RPMI 1640 medium (Table 1), which may be related to the low FBS concentration or the effect of phosphate ions in the cell culture medium.

By TEM, the aggregation of $500 \ \mu\text{g/mL TiO}_2$ NPs was also observed in distilled water (Figure 1). Therefore, it was not possible to determine the nanoparticles' size. UV/Vis spectra were similar in exposed and control samples in both distilled water and R-RPMI 1640, as well as during the experiment, indicating that NPs properties did not change. Previously, the addition of HEPES and FBS into RPMI-1640 medium led to NPs re-dispersion [14].

Scanning electron microscopy showed the NPs cluster in contact with the cell membranes (Figure 2). EDS spectra showed TiO₂ NPs that are present on cells at the 100 μ g/mL concentration (Figure 3), but not at the lesser concentration (10 μ g/mL). This may be because of the EDS microanalysis detection limit. TiO₂ NPs are internalized by *E. fetida* coelomocytes. Bigorgne et al. determined their presence in the cell cytoplasm, but not in the nucleus or mitochondria [14]. However, we were unable to detect TiO_2 NPs inside coelomocytes. This could be because TiO₂ NPs aggregates are large. Earthworm coelomocytes are probably unable to engulf large NP clusters via phagocytosis and/or endocytosis, the most probable routes of TiO_2 NPs entry into coelomocytes [1,14]. Phagocytic cells are potentially the most affected because they engulf NPs. Coelomocyte viability was not affected by exposure to 1, 10, and 100 μ g/mL TiO₂ NPs for 2, 6, and 24 h. Similar results were observed in *E. fetida* coelomocytes exposed to TiO₂ NP [14]. Nanoparticles often trigger reactive oxygen species (ROS) production in cells, resulting in biomolecule oxidative damage [27,28]. We did not detect any statistically significant differences in ROS production in TiO₂ NPs-exposed cells in comparison with control cells (Figure 4). Cells exposed to other nanoparticles, such as Ag NPs, nZVI NPs or ZnO NPs release significantly greater ROS amounts. Contrary to TiO_2 NPs, ROS production could be elicited by the metal ions released from these nanoparticles [11,29,30].

We evaluated the apoptotic process in cells treated with TiO_2 NPs, and did not detect any significant differences between exposed and control coelomocytes (Figures 5 and 6). Late apoptosis was similar in GA and HA, with the greatest difference observed after 24 h of incubation. HA population exerted relatively greater early apoptosis than GA. Excess ROS production led to decreased cell viability and apoptosis [31,32]. Homa et al. suggested that coelomocytes are susceptible to bacterial or fungal products that may induce programmed cell death [31]. TiO_2 NPs did not increase ROS production, and simultaneously, apoptosis was not increased as compared to control cells (Figures 4–6). We suggest that TiO_2 NPs do not affect ROS production, and thus do not trigger the apoptotic pathway in amoebocyte subpopulations (HA and GA).

Amoebocytes are earthworm immune effector cells with the ability to phagocytose. At the phagocytic activity level, control cells and cells exposed to TiO_2 NPs (1, 10, and 100 µg/mL) did not show any statistically significant changes (Figure 7). The results are in accordance with Bigorgne et al., who reported that there were no phagocytic activity changes in coelomocytes exposed to 1, 5, 10, and 25 µg/mL of TiO_2 NPs, although TEM images demonstrated that TiO_2 NPs were engulfed by the coelomocytes [14]. Thus, we can confirm that phagocytic activity is not compromised due to TiO_2 NPs exposure.

ROS production initiates harmful radical chain reactions on cellular macromolecules, including DNA mutation, protein denaturation, and lipid peroxidation. At the lipid peroxidation level, MDA production was similar in both exposed and control cells. MDA is a subproduct derived from the reaction of free radical species with fatty acids [33]. We did not observe elevated lipid peroxidation (Figure 8). Ayala et al. explained that MDA is more

stable and has a greater lifespan than ROS, and therefore it is more toxic [33]. Therefore, it could be a better biomarker for cellular oxidative stress detection. Excess ROS leads to MDA production [34]. Two oxidative stress markers, ROS and MDA, were produced at similar levels in control cells and TiO_2 NPs-exposed cells (Figures 4 and 8). The same results were also observed after THP1 human cells and sea urchin cells were exposed to TiO_2 NPs [20,35]. UVA light could also enhance ROS production and increase toxicity several fold [36]. However, in this instance, the cells were mimicking the environmental conditions in the soil ecosystem, where UVA light was not present.

Significant differences between exposed and control cells were not detected regarding DNA damage. The alkaline comet assay results showed that there is no significant DNA damage in coelomocytes exposed to 1, 10, and 100 μ g/mL of TiO₂ NPs for 2, 6, and 24 h (Figure 9). A relationship between ROS, MDA, and DNA damage has been suggested. As mentioned previously, ROS may induce MDA production, which, in turn, affects nucleosides and results in DNA damage [33,34]. This mechanism has been described in coelomocytes exposed to pollutants, antibiotics, or pathogens [34]. Reeves et al. showed that GFSk-S1 cells (primary cell line from goldfish skin) exposed to different doses of TiO₂ NPs (1, 10, and 100 μ g/mL) could result in slight DNA damage, whereas co-exposure with UVA caused a significant increase in toxicity [36]. In vitro analysis described in this study did not reveal substantial changes in cellular physiologic activities, but the long-term exposure experiments can reveal different findings [37]. Zhu et al. described transcriptomic and metabolomic changes in earthworms as a global response to TiO₂ NPs exposure that cannot be observed by conventional toxicity endpoints [38].

Treatment of coelomocytes with TiO_2 NPs induced slight changes in the mRNA levels of distinct molecules. Metallothioneins are proteins protecting against metal-induced oxidative stress [9]. Metallothionein was upregulated in coelomocytes exposed to 10 µg/mL TiO₂ NPs for 6 h, respectively (Table 2). This is in agreement with Bigorgne et al., who immunostimulated coelomocytes with lipopolysaccharides (LPS) (500 ng/mL) for 5 h prior to TiO₂ NPs addition. After 12 h of incubation with 10 and 25 µg/mL TiO₂ NPs, metallothioneins were upregulated [14]. We determined that even 1 µg/mL TiO₂ NPs concentration upregulated metallothionein expression during the whole experiment (Table 2). Interestingly, the highest upregulation was detected in coelomocytes incubated with 1 µg/mL TiO₂ NPs already after 2 h. Further, the induction of metallothionein expression in cells exposed to 10 µg/mL TiO₂ NPs started at 6 h, and afterward decreased after 24 h (Table 2). Bigorgne et al. similarly showed increased metallothioneins expression after 12 h of incubation, with a subsequent decrease after 24 h [14].

Further, the antioxidant enzymes were not affected, except for Mn-SOD, which was downregulated after 6 h of coelomocyte exposure to 10 μ g/mL TiO₂ NPs (Table 2). Mn-SOD is a mitochondrial protein that protects cells against oxidative stress [39]. It seems that macrophages (RAW 264.7 cell line) and coelomocytes can engulf TiO₂ NPs. These nanoparticles affect mitochondria even if they are not located inside the mitochondria [12,14]. Moreover, TiO₂ NPs decreased ATP production in the macrophage RAW 264.7 cell line [12]. Thus, engulfed TiO₂ NPs could target mitochondria and cause mitochondrial malfunction [12]. Mn-SOD downregulation and loss in mitochondrial oxidative phosphorylation function was also reported in primary rat hepatocytes [40].

Elevated levels of the antimicrobial proteins fetidin/lysenin and lumbricin were detected in cells exposed to $10 \ \mu\text{g/mL}$ TiO₂ NPs for 6 h (Table 2). Similarly, Bigorgne et al. observed that fetidin was upregulated in cells exposed to $10 \ \mu\text{g/mL}$ TiO₂ NPs after $12 \ h$ [14]. As previously described in related earthworm species *E. fetida*, lysenin regulation is changed rapidly by environmental stressors and it is suggested as an early biomarker of stress [41]. However, we cannot exclude that the increase in antimicrobial protein mRNA levels could be caused by used TiO₂ NPs that were not LPS-free.

Referring to the signal transduction molecules, PKC I was strongly downregulated after coelomocyte exposure to 10 μ g/mL TiO₂ NPs for 6 and 24 h (Table 2). PKC I is important in cellular homeostasis and is involved in the cell proliferation signaling cas-

cade [42,43]. This downregulation could suggest a coelomocyte homeostasis destabilization upon TiO₂ NPs exposure. Another signal transduction molecule, MEKK, was upregulated in coelomocytes exposed to 1 μ g/mL TiO₂ NPs for 24 h and in coelomocytes exposed to 10 μ g/mL TiO₂ NPs for 6 h (Table 2). This molecule is involved in the MAPK cascade participating in many cellular processes, besides others in stress signaling [9,43]. Generally, coelomocyte exposure to TiO₂ NPs results in slight changes in the mRNA levels of various molecules, however, these changes seem not to be significant enough to affect the observed cellular functions.

5. Conclusions

Coelomocytes exposed to TiO₂ NPs (1, 10, and 100 μ g/mL) did not show any impaired cellular responses as compared to control cells. The oxidative stress pathway and phagocytic activity were not affected as well. Nanoparticles do not cause greater DNA damage in treated cells than in non-treated cells. We also detected some gene expression alterations involved in metal detoxification, oxidative stress, defense reactions, and signal transduction. However, these changes do not seem to affect the observed cellular functions. In summary, we did not determine any detrimental effects of TiO₂ NPs on *E. andrei* coelomocytes.

Supplementary Materials: The following are available online at https://www.mdpi.com/2079-4 991/11/1/250/s1, Figure S1. Illustrative figure of NPs distribution incubated without and with the cells. Figure S2. EDS spectra from standard non-treated cells. Figure S3. Illustrative figure of coelomocytes subpopulations detected by flow cytometry. Figure S4. Illustrative histogram of ROS production between control samples and positive control. Figure S5. Illustrative figure of apoptosis of GA after 24 h of exposure to 100 μ g/mL TiO₂ NPs. Figure S6. Illustrative figure of apoptosis of HA after 24 h of exposure to 100 μ g/mL TiO₂ NPs. Figure S7. Illustrative figure of phagocytic activity of GA after 2 h. Figure S8. Illustrative figure of phagocytic activity of HA after 2 h. Figure S9. Detection of Fluoresbrite[®] YG Plain 1 μ m microsphere. Table S1: Primer sequences used for qPCR.

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