

Physicochemical properties, in vitro cytotoxic and genotoxic effects of PM_{1.0} and PM_{2.5} from Shanghai, China

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Abstract Exposure to ambient particulate matter (PM) links with a variety of respiratory diseases. However, compared with coarse particles (PM₁₀) and fine particles (PM_{2.5}), submicrometer particles (PM_{1.0}) may be a more important indicator of human health risks. In this study, the cytotoxic and genotoxic effects of PM_{1.0} samples from Shanghai were examined using A549 cells, and compared with the effects of PM_{2.5}, to better understand the health effects of PM_{1.0} in this area. The PM_{1.0} and PM_{2.5} samples were characterized for morphology, water-soluble inorganic ions, organic and elemental carbon, and metal elements. The cytotoxicity of PMs was measured using cell viability and cell membrane damage assays. The genotoxic effects of PMs were determined using the comet assay, and DNA damage was quantified using olive tail moment (OTM) values. The physicochemical characterization indicated that PM_{1.0} was enriched in carbonaceous elements and hazardous metals (Al, Zn, Pb, Mn, Cu, and V), whereas PM_{2.5} was more abundant in large, irregular mineral particles. The biological results revealed that both PM_{1.0} and PM_{2.5} could induce significant cytotoxicity and genotoxicity in A549 cells, and that exposure to PM_{1.0} caused more extensive toxic effects than exposure to PM_{2.5}. The greater cytotoxic effects of PM_{1.0} can be attributed to the combined effects of size and chemical composition, whereas the genotoxic effects of PM_{1.0} may be mainly associated with chemical species.

Keywords PM_{1.0} · Physicochemical properties · A549 cells · Cytotoxic effects · Genotoxic effects · DNA damage

Introduction

Ambient particulate matter (PM) is a complex air pollutant that originates from both natural sources (e.g., dust, sea salt, and fungal spores) and anthropogenic emissions (e.g., diesel exhaust, coal burning, and cigarette smoke) (Kelly and Fussell 2012). The major components of PM include acid salts, heavy metals, organic chemical components, and biological contaminants (Hueglin et al. 2005). Daily exposure to PM is associated with a variety of cardiovascular and respiratory diseases, and even the increased risk of lung cancer (André 2005). Chronic lung disease is among the major problems that has elicited great public health concern (Vinikoor-Imler et al. 2011). Moreover, increasing researchers have recognized that the production of reactive oxygen species (ROS), immune deficiency, inflammation, and DNA degradation are involved in PM-mediated lung injury (Deng et al. 2013; Senthilkumara et al. 2014).

The relative ability of PMs to provoke adverse effects in target cells and tissues may depend greatly on their chemical constituents (Thomson et al. 2015). A number of studies have demonstrated significant associations between various health outcomes and the chemical components of PM, such as sulfate, ammonium, and nitrate (Bonetta et al. 2009; Pui et al. 2014). Furthermore, PM size is an important factor influencing biological responses (Lee et al. 2006; Velali et al. 2016). According to the size of the aerodynamic diameter (\varnothing), PM can be classified into coarse particles (PM₁₀, \varnothing between 2.5 and 10 μm), fine particles (PM_{2.5}, $\varnothing < 2.5 \mu\text{m}$), and submicrometer particles (PM_{1.0}, $\varnothing < 1.0 \mu\text{m}$) (Huang et al. 2003). The toxic effects of PM₁₀ and PM_{2.5} have been widely

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examined; the coarse fraction was reported to be mainly associated with pro-inflammatory and cytotoxic effects, whereas the fine fraction was mainly related to genotoxic potential (Akhtara et al. 2014; Gualtieri et al. 2010; Osornio-Vargas et al. 2003).

Recently, the health threat of PM_{1.0} has attracted growing interest, because PM_{1.0} is more enriched in toxic and carcinogenic compounds and can penetrate deeper into the respiratory tract than fine and coarse particles (Lim et al. 2012; Longhin et al. 2013). Several studies have been conducted to explore the toxic effects of PM_{1.0} in developed areas. For instance, Huang et al. (2003) reported that PM_{1.0} in Taiwan could result in significantly higher interleukin-8 (IL-8) production and lipid peroxidation than PM_{2.5–10} because of its higher contents of metals (Cr, Mn, and Fe) and carbon compounds. Perrone et al. (2010) found that PM_{1.0} in Milan could induce a more significant reduction in cell viability and increase of IL-8 than the PM_{2.5} fraction. In a more recent study, Longhin et al. (2013) concluded that PM_{1.0} in Milan could elicit high ROS generation, and exhibited considerable DNA-damaging effects.

The mega-city of Shanghai in China suffers from severely deteriorated air pollution. During 2011–2012, the annual PM_{2.5} mass concentration in Shanghai was 54.94 µg/m³, five times higher than the global guidelines (10 µg/m³) that set by the World Health Organization (Qiao et al. 2014). Particularly, PM_{1.0} composing the dominant proportion of fine PM (Gong et al. 2015; Huang et al. 2012; Li et al. 2015) is presumed to play an important role in adverse health effects. However, studies on the toxic effects of PM_{1.0} in Shanghai are scarce.

In the present study, airborne PM_{1.0} and PM_{2.5} from the Minhang District of Shanghai were used as model samples to elucidate the adverse biological effects of PM_{1.0} in Shanghai. The PM samples were characterized in terms of their morphology, water-soluble inorganic ions, organic carbon (OC) and elemental carbon (EC) contents, and metal element contents. To investigate the *in vitro* biological effects of PM_{1.0} and PM_{2.5}, which include cell viability, cell membrane disruption, and DNA damage, the human lung adenocarcinomatous cell line A549 was used. The A549 cell line, which is derived from the neoplastic transformation of type-II alveolar cells, shows high proliferative potential and possesses the typical features of type-I normal epithelial cells (Calabrini et al. 2004). The A549 cell is one of the major *in vitro* cell lines used extensively in the study of airway particle toxicology, although this cell line is sensitive to minor genetic modifications (Alessandria et al. 2014; Berg et al. 2013; Zeng et al. 2016).

Materials and methods

PM sampling and extraction

Sampling was conducted on the roof of a six-floor building in the Minhang campus of Shanghai Jiao Tong University (N 31°

1' 51" E 121° 26' 27"), Shanghai, China. PM_{1.0} and PM_{2.5} samples were collected simultaneously using two mid-volume samplers (Wuhan Tianhong TH III, China) equipped with PM_{1.0} and PM_{2.5} cut-off inlets, respectively. Daily airborne PM samples were collected on quartz fiber filters (Ø 90 mm), with a pump flow rate of 100 L/min. Sampling was carried out over a continuous period of 24 h/day, from September 2 to November 19, 2014. Treatment of the filters before and after sampling was performed following the methods described in our previous work (Zou et al. 2016).

The suspensions of PM_{1.0} and PM_{2.5} used for toxic assays were prepared as follows. A portion of each sampled filters was cut and immersed in ultrapure water with sonication in water bath for 6 × 30 min. The extract was filtered with eight-layer sterile gauze, and transferred into a clean sterile tube (weighed before use). The suspension was freeze-dried in a vacuum, and the resulting pellet was weighed and stored in −80 °C until use.

Morphological characterization of PM_{1.0} and PM_{2.5}

The morphological characteristics of the extracted particles (PM_{1.0} and PM_{2.5}) were investigated using scanning electron microscopy (SEM) and transmission electron microscopy (TEM). Prior to analysis, 1 cm² of the PM_{1.0} and PM_{2.5} filters were cut and sonicated in 1 mL of ethanol (analytical reagent grade) separately. For SEM observation, several drops of each suspension were dropped onto the surface of a double conductive adhesive fixed on an aluminum stub. After drying, each sample was coated with gold and put into the sample compartment of the SEM (FEI, Nano 230, USA) for further analysis. For TEM analysis, a portion of the resuspended particles was placed onto carbon-coated copper grids and air-dried. The grids were analyzed with a FEI TEM (Tecnai G2 spirit Biotwin, USA).

Analysis of chemical species in PM_{1.0} and PM_{2.5}

PM_{1.0} and PM_{2.5} filter samples were analyzed for water-soluble inorganic ions, OC and EC, and metal elements. To determine the contents of water-soluble inorganic ions, one-eighth of the sampled filters were sonicated in 25 mL of ultrapure water for 45 min. The extract was then filtered (with 0.45 µm PTFE filters) to remove the insoluble components. NH₄⁺, F[−], Cl[−], NO₃[−], and SO₄^{2−} ions in the extracts were analyzed via ionic chromatography (IC, Thermo Fisher, ICS-5000⁺, USA), and Na⁺, K⁺, Ca²⁺, and Mg²⁺ were quantified via inductively coupled plasma spectrometry (ICP, Thermofisher, iCAP7600, USA). The OC and EC fractions were analyzed with a thermal/optical carbon analyzer (Atmoslytic Inc., DRI2001A, USA) using the thermal optical transmission (TOT) method, following the EUSAAR2 temperature protocol (Cavalli et al. 2010). A 0.518 cm² punch

area of each filter was used for the measurement of eight carbon fractions. First, the filter was heated to 650 °C in a pure helium atmosphere (OC1, OC2, OC3, and OC4 at 200, 300, 450, and 650 °C, respectively). Subsequently, oxygen was added to the combustion atmosphere, and a pyrolyzed carbon fraction (OP) was determined when reflected laser light attained its original intensity. Lastly, the filter was brought to 850 °C in a 2% oxygen and 98% helium oxidizing atmosphere (EC1, EC2, and EC3 at 550, 700, and 850 °C, respectively). OC is defined as OC1 + OC2 + OC3 + OC4 + OP, and EC is calculated as EC1 + EC2 + EC3 – OP. Metal elements were measured via inductively coupled plasma mass spectrometry (ICP-MS, Thermo Fisher, iCAPQ, USA). Before analysis, one-eighth of the sampled filters were digested using a mixture of HNO₃ and HF in a microwave digestion system (Anton Paar, Multiwave 3000, Austria). After cooling, the mixture was adjusted to a final volume of 25 mL with distilled water. A total of 15 elements (Al, Zn, Fe, Pb, Mn, Cu, V, Cr, As, Se, Ni, Sb, Mo, Cd, and Co) were measured. For all analyses, field blank filters were measured together with each batch of samples to provide information about contaminants that may have been introduced during sample collection, storage, and transport.

Cell culture

In the current investigation, A549 cells were used to examine the biological endpoints of PM_{1.0} and PM_{2.5}. A549 cells were grown in F-12K (Life Technologies, Thermo Fisher, USA) cell culture medium supplemented with 10% heat-inactivated fetal bovine serum (Life Technologies, Thermo Fisher, USA) and 1% antibiotics penicillin/streptomycin (100 U/mL). The cells were cultivated in plastic cell culture flasks at 37 °C in a humidified atmosphere containing 5% CO₂. After 70–80% cell confluency was reached, the monolayer cells were used for toxicity assays.

Cell viability assay

The MTT assay (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide), which mainly reflects the mitochondrial activity of cells, was employed to evaluate cell viability. For the MTT assay, A549 cells were seeded in a 96-well plate at a density of 8×10^3 cells/well and cultured for 24 h to allow cell adhesion. The cells were then exposed to PM_{1.0} and PM_{2.5} in the concentration ranges of 25, 50, 100, and 200 µg/mL. After 24-h incubation, freshly prepared MTT solutions were added to each well at a final concentration of 200 µg/mL, and incubated with cells for 4 h at 37 °C in darkness. Later, the cell culture medium was discharged, and 200 µL of DMSO was added to each well to dissolve the formazan compound. The optical density (OD) of the solution was recorded at a wavelength of 492 nm using a multi-scan plate reader (BioTek,

Synergy H4, USA). For each concentration of PMs, six replicates were tested, including the untreated control cells. The OD values of PM-exposed cells were compared with those of the control cells to determine the percentage of cell viability.

Cell membrane damage assay

Lactate dehydrogenase (LDH) is a stable cytoplasmic enzyme that is rapidly released into the culture medium when cell membranes are damaged. In this study, LDH leakage in the cell culture medium was assayed to determine the cell membrane disruption of PM-treated A549 cells. For the LDH assay, the cell culture and PM exposure were conducted as for the MTT assay described above. The activity of LDH in the cell-free culture supernatant was detected following the manufacturer's protocol (CytoTox 96, Promega, USA). Briefly, the cell-free culture supernatants were obtained using centrifugal separation (800 rpm, 10 min), and 50 µL of the clean supernatant in each well was transferred to a new 96-wells plate. Then, 50 µL of the pre-prepared substrate mixture was added to each well and incubated with the supernatant for 30 min at room temperature in darkness. Next, 50 µL of stop solution was added to each well, and the OD values of the final solution were recorded on the multi-scan plate reader (BioTek, Synergy H4, USA) at a wavelength of 490 nm. The culture medium without cells served as the background control, the supernatants from lysis solution-treated cells were used as positive controls. Each concentration of PMs was tested in six replicates, including the untreated control cells and the positive control. The results were calculated as fold of LDH release to control.

DNA damage assay

The comet assay, a gel electrophoresis-based method, is widely used for measuring chemical and environmental contaminant-induced DNA damage in individual cells. In this study, the alkaline version of the comet assay, which is sensitive to DNA strand breaks, alkali-labile, and incomplete excision repair sites, was used to detect the DNA damage potential of PM_{1.0} and PM_{2.5}. For the comet assay, A549 cells were seeded in a 12-well plate at a density of 2×10^4 cells/well. After 24 h of seeding, cells were exposed to different concentrations of PM_{1.0} and PM_{2.5} (25, 50, 100, and 200 µg/mL) for 24 h. The cells were then harvested with 0.25% Trypsin-EDTA and washed with phosphate-buffered saline (PBS) three times. Then, 15 µL of the cell suspension (1×10^5 cells/mL in PBS) was mixed with 110 µL of 0.65% low-fusion-point agarose and applied to the prepared slides. The cells were stained with GelRed (Biotium, USA) and lysed under alkaline conditions (2.5 M NaCl, 10 mM Tris, 100 mM Na₂EDTA, 1% sodium sarcosinate, pH = 10), to which 1% Triton X-100 and 10% DMSO were added, for at

least 1 h. Then, the samples were heated with Protease K buffer solution (0.5 mg/L Protease K, 2.5 M NaCl, 5 mM Tris, 0.05% sodium sarcosinate, pH = 7.4) in a water bath at 55 °C for 2 h. The slides were then placed in a single-cell gel electrophoresis chamber (Trevigen; Maryland, USA), positioned close to the anode and covered with an alkaline buffer solution (300 mM NaAC, 100 mM Tris, 100 mM Na₂EDTA, pH = 10). After the DNA was unwound for 30 min, it was electrophoresed at 15 V and 130 mA for 60 min. After the slides were rinsed and soaked in 75% ethanol for 5 min, they were examined under a DMLB microscope (Olympus, Japan) with an adapted charge-coupled device camera (Cohu Inc.; San Diego, USA). Images from 100 random cells were analyzed for each experiment as described by the guidelines. To quantify the level of DNA damage, the extent of DNA migration was determined based on the olive tail moment (OTM), which is the relative amount of DNA in the tail of the comet multiplied by the median migration distance. Statistical analysis was based on the mean OTMs of scored cells.

Statistical analysis

Values are given as means ± standard deviations (SD). Data were analyzed using one-way analysis of variance (ANOVA) to determine significance relative to the unexposed controls. In all cases, $P < 0.05$ was considered significant.

Results and discussion

Morphology of PM_{1.0} and PM_{2.5}

The morphological features of PM_{1.0} and PM_{2.5} particles obtained from SEM and TEM observations are shown in Fig. 1. As depicted in Fig. 1a, b, regular and irregular mineral particles were observed in both PM_{1.0} and PM_{2.5}. According to Murillo et al. (2012), irregular mineral particles were related to natural sources, such as feldspar, clay, and limestone. In contrast, regular mineral particles might be generated from high-temperature processes of anthropogenic origin or biomass burning. Compared with PM_{1.0}, there were greater numbers of large irregular mineral particles in PM_{2.5}, which was consistent with previous research that reported that mineral particles of dust tended to be found in PMs of larger size fractions (Wang et al. 2013). Despite the slight difference observed between SEM images of the two kinds of particles, TEM images of the ultrafine particles in PM_{1.0} (Fig. 1c) and PM_{2.5} (Fig. 1d) showed great similarity in particle morphology, which indicated that the majority of ultrafine particles in PM_{1.0} originated from the same sources as PM_{2.5}. These round-shaped nanoparticles, with average sizes of approximately 30–40 nm, were agglomerated in clusters and interconnected with each other in various ways. In addition, the

appearance of these ultrafine particles is similar to the morphology of vehicle-emitted particles (Barone and Zhu 2008), which demonstrates that the ultrafine particles from the sampling site may have been generated from burning of gasoline and diesel fuel.

Mass fractions of main chemical components in PM_{1.0} and PM_{2.5}

The chemical composition of size-fractionated ambient PM aerosols is an important factor affecting their health effects. Therefore, in the present study, the chemical species in PM_{1.0} and PM_{2.5}, samples which include water-soluble inorganic ions, OC, EC, and metal elements, were quantified to better assess their toxicity. The relative mass fractions of these chemical components are depicted in Fig. 2. As shown in Fig. 2a, b, water-soluble inorganic ions (SO₄²⁻, NO₃⁻, and NH₄⁺) were the most abundant constituents in PM_{1.0} and PM_{2.5}, followed by carbonaceous elements (OC and EC). According to Perrone et al. (2010), inorganic ions such as SO₄²⁻, NO₃⁻, and NH₄⁺ that resulted from the transformation of their gaseous precursors SO₂, NO_x, and NH₃ represented secondary pollution. Therefore, the high mass fraction of inorganic ions (SO₄²⁻, NO₃⁻, and NH₄⁺) in PM_{1.0} and PM_{2.5} indicates pronounced secondary contamination in this region. Moreover, Perronea et al. (2014) and Salameh et al. (2015) have demonstrated that an OC/EC ratio exceeding 2.0–2.2 can be used as an indicator of secondary contamination, because EC is mainly derived from primary anthropogenic combustion, whereas OC is produced either from direct emission or gas-to-particle reactions involving atmospheric oxidation or oligomerization. Therefore, in addition to the high contents of water-soluble secondary ions, the relatively high OC/EC ratios of PMs (2.35 for PM_{1.0} and 3.09 for PM_{2.5}) from the sampling site further confirms severe secondary pollution in this area. The present results are broadly consistent with previous findings that secondary aerosols dominated the fine particulate mass in this region (Feng et al. 2009). Regarding to the mass fraction of chemical species in PM_{1.0} and PM_{2.5}, no significant differences were found for water-soluble ions in the two kinds of PMs (Fig. 2a). In contrast, significantly higher proportions of carbonaceous and toxic metal elements (Al, Zn, Pb, Mn, Cu, and V) were observed in PM_{1.0} (Fig. 2b). Carbonaceous aerosols were reported to be major contributors to fine aerosols smaller than 1 μm, and typically constitute significant, sometimes dominant, fractions of the total mass of submicron particles (Huang et al. 2012). Similarly, hazardous metal elements with notable health impacts emitted from anthropogenic sources were found to be mainly accumulated in the smaller particles ($\varnothing < 1 \mu\text{m}$) (Espinosa et al. 2001). The size distribution of the chemical species in PM_{1.0} and PM_{2.5} is consistent with their morphological characters; PM_{2.5} is rich in large, irregular mineral particles from natural sources, whereas

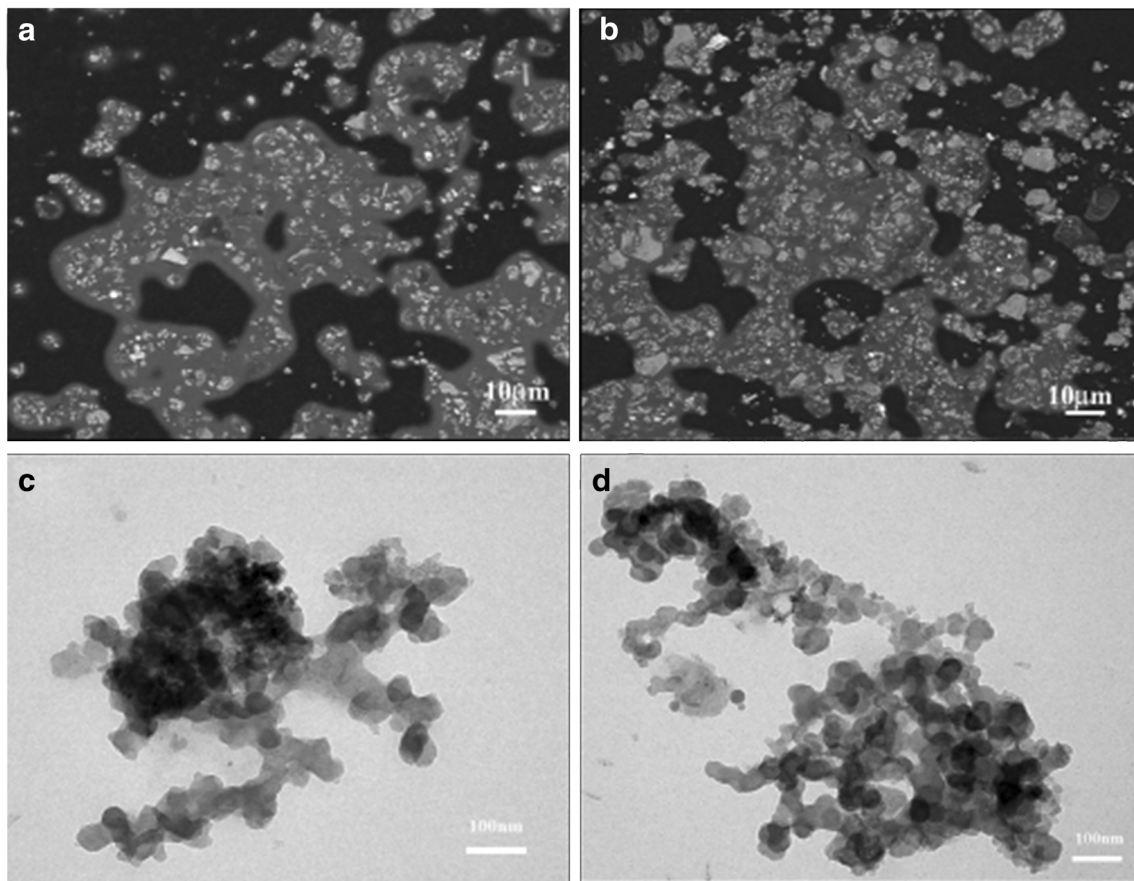


Fig. 1 Morphological characteristics of PM_{1.0} and PM_{2.5}. SEM image of PM_{1.0} (a) and PM_{2.5} (b), scale bar = 10 μm; TEM image of PM_{1.0} (c) and PM_{2.5} (d), scale bar = 100 nm

PM_{1.0} is enriched in chemicals emitted from anthropogenic processes. Consequently, PM_{1.0} may be analyzed to minimize the interference from natural sources, and provide better information for human health than PM_{2.5} (Lee et al. 2006).

Cytotoxic effects of PM_{1.0}- and PM_{2.5}-treated A549 cells

To obtain more reliable results in evaluating the potential cytotoxic effects of PM_{1.0} and PM_{2.5}, both MTT viability assay and LDH release assay were employed to examine the toxic effects of PM_{1.0}- and PM_{2.5}-treated A549 cells; the results are

presented in Fig. 3. As illustrated in Fig. 3a, for the MTT assay, cell viability decreased in PM_{1.0}- and PM_{2.5}-treated A549 cells, and significant decrease in cell viability relative to the control cells occurred at PM mass concentrations of 25 μg/mL. Moreover, LDH release was increased in PM_{1.0}- and PM_{2.5}-exposed A549 cells, with significant increase of LDH release from PM concentrations of 50 μg/mL (Fig. 3b). These results indicate that the amount of LDH release from cells exposed to PMs is related to the decrease of cell viability induced by the particles. The data in our study are consistent with previous observations that the increase of LDH release, as an indicator

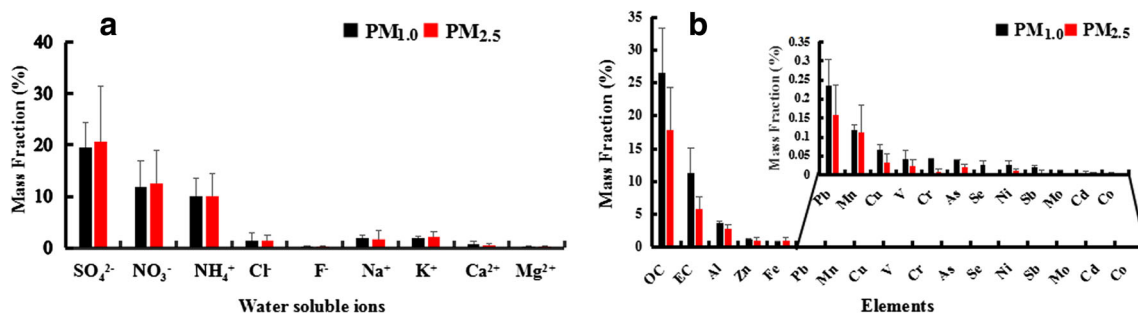


Fig. 2 Mass fractions of chemical components in PM_{1.0} and PM_{2.5} samples. a Water-soluble inorganic ions. b Carbonaceous elements (OC and EC) and metal elements

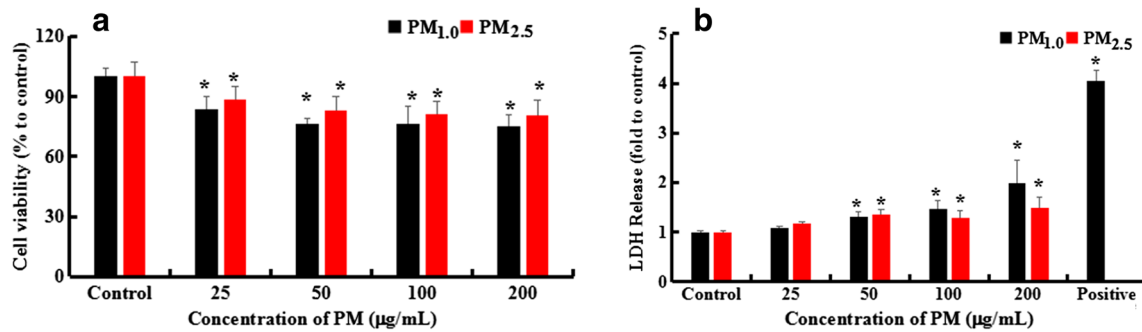


Fig. 3 The cytotoxic effects of A549 cells treated with PM_{1.0} and PM_{2.5} at concentrations of 25, 50, 100, and 200 µg/mL for 24 h. **a** MTT assays of cell viability; data are shown as percentages relative to the control

(untreated A549 cells). **b** Content of LDH release in cell culture medium; data are shown as fold to control. **P* < 0.05 (one-way ANOVA with Dunnett post-test)

of PM exposure, caused cell membrane disturbance, and was highly correlated with decreases of cell viability and other biological effects (Billet et al. 2007; Shang et al. 2014). Although both assay methods showed great similarity in PM_{1.0}- and PM_{2.5}-induced cytotoxic effects, differences were also detected for the two size fractions of PMs. The results clearly revealed that PM_{1.0} was more potent in inducing cytotoxic effects than was PM_{2.5}, as PM_{1.0} could induce significantly lower cell viability at each exposure concentration, as well as significantly higher LDH release, relative to PM_{2.5} at exposure doses of 100 and 200 µg/mL. For the different effects observed for PM_{1.0} and PM_{2.5}, both particle size and source-linked chemical composition are considered to play dominant roles. Massolo et al. (2002) and Velali et al. (2016) have reported a strong correlation between cell cytotoxicity and the finest PM fraction, and Thomson et al. (2015) showed that particle toxicity might be related to the contents of metals and organic compounds. Carbon-containing exhaust and some transition metals were found to be associated with enhanced intracellular oxidants and the resulting cell injuries (Huang et al. 2003). Accordingly, Perrone et al. (2010) found that PM_{1.0} could induce a greater reduction of cell viability in A549 cells than PM_{2.5}, and that the MTT response appeared to be related to As, Zn, Cr, Cu, and Mn contents. More recently, Perrone et al. (2013) reported that the LDH response in PM_{1.0}- and PM_{2.5}-treated cells was associated with metals from anthropogenic sources and organic compounds from biogenic and biomass-burning origins. Therefore, the significantly higher cytotoxicity of PM_{1.0} compared with PM_{2.5} reported in our study can mainly be attributed to the combined effects of its smaller particle size and higher contents of carbonaceous components and metal elements.

DNA damage in PM_{1.0}- and PM_{2.5}-treated A549 cells

In the present study, DNA damage in PM-treated A549 cells was investigated, using the comet assay, to evaluate the genotoxic effects of PM_{1.0} and PM_{2.5}. Images of assayed cells are presented in Fig. 4. As shown in Fig. 4, both PM_{1.0} and

PM_{2.5} demonstrated DNA-damaging potential, which was indicated by relatively smaller comet heads and increased lengths of tails in the exposed cells, compared with control cells. The results of our study are in accordance with general observations demonstrating that inhaled particles can cause the breakage of DNA strands and oxidative damage to DNA (Upadhyay et al. 2003; Zeng et al. 2016). The chemical compositions of PMs have been reported to be the main factor in PM-induced oxidative stress and DNA damage. For instance, Lu et al. (2008) found that the plasmid DNA damage induced by Shanghai PM_{2.5} was highly correlated with heavy metal content. Other studies have shown that PM samples collected in high-traffic locations, which greatly enriched in metal elements, such as Cu, Ni, and Cd, were able to inhibit DNA repair pathways (Bonetta et al. 2009; Wessels et al. 2010). Moreover, Shang et al. (2014) and Velali et al. (2016) have demonstrated that oxidative damage of DNA caused by PMs is mainly associated with the organic fractions. In addition to the considerable importance of chemical composition, the insoluble particles in PMs may also play a critical role in inducing DNA damage. As demonstrated in our previous study (Zou et al. 2016), smaller nanoparticles that were either already present in PM_{2.5} or were generated from decomposition processes were able to penetrate into the nuclear core of A549 cells, and might attack crucial cellular macromolecules including the genomic DNA.

In addition to the above information, there are several specific characteristics of PM_{1.0}- and PM_{2.5}-induced DNA damage (Fig. 4). First, the analysis revealed significant cellular DNA damage even in the presence of low concentrations of PM_{1.0} and PM_{2.5} (25 µg/mL), indicated by the presence of DNA tails in Fig. 4b, f. This finding can mainly be attributed to the fact that the comet assay can provide sensitive and reliable results for DNA damage detection. In contrast to the traditional chromosome aberration and micronuclei tests, the comet assay allows rapid detection of DNA damage shortly after the injury, before the DNA is repaired, and without any need to wait for progression into mitosis (Liao et al. 2009). Secondly, PM_{1.0} and PM_{2.5} may induce more extensive DNA

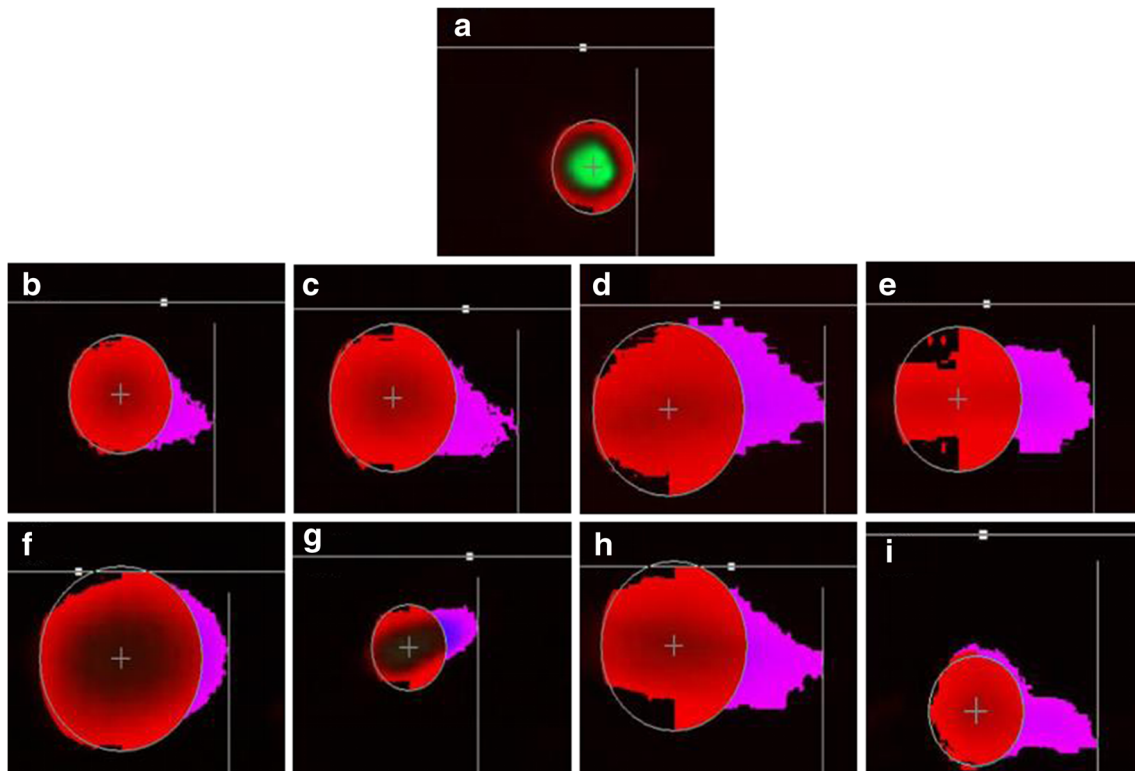


Fig. 4 Comet assay of A549 cells exposed to PM_{1.0} and PM_{2.5} for 24 h. **a** Control A549 cells. **b–e** PM_{1.0}-treated A549 cells in the concentration ranges of 25, 50, 100, and 200 µg/mL, respectively. **f–i** PM_{2.5}-treated A549 cells in the concentration ranges of 25, 50, 100, and 200 µg/mL

damage in cells treated with a higher concentration of PMs, as demonstrated by DNA tails with greater areas and lengths (distance from DNA head to the end of DNA tail) in cells exposed to higher doses of PMs. These findings are in good agreement with previously reported data, which indicated that particle exposure can cause increased DNA damage in a dose-dependent manner (Bonetta et al. 2009; Shang et al. 2014). Thirdly, PM_{2.5} was found to be less potent than PM_{1.0} in inducing oxidative DNA damage in A549 cells. Because there are no agreeable comments about the outcomes of DNA damage outcomes and particle size fractions, the greater DNA damage induced by PM_{1.0} can be attributed to the abundant

chemicals species, such as metals and organic compounds. For example, some studies have showed that PM_{2.5} has the ability to induce stronger DNA damage than PM_{1.0} in the comet assay (Perrone et al. 2010), whereas Osornio-Vargas et al. (2003) reported that PM₁₀ samples could induce higher DNA degradation than PM_{2.5}. Moreover, Wessels et al. (2010) have emphasized that the genotoxic effects of PMs depend more on the chemical composition of the PMs than on the size fractions.

To better compare the genotoxic effects induced by PM_{1.0} and PM_{2.5}, quantitative data from comet assays were analyzed; the results are shown in Fig. 5. Tail length, percentage

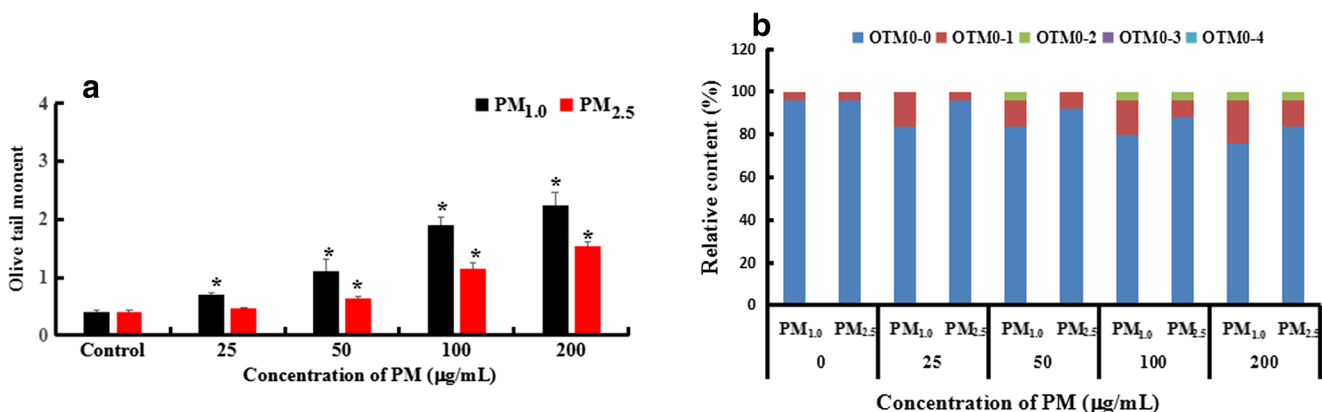


Fig. 5 The genotoxic effects of PM_{1.0} and PM_{2.5}. A549 cells were treated with PM_{1.0} and PM_{2.5} at concentrations of 25, 50, 100, and 200 µg/mL for 24 h. **a** OTM values of assayed A549 cells. **b** Percentages of OTM damage classes in the exposed cells. * $P < 0.05$ (one-way ANOVA with Dunnett post-test)

of DNA in the tail, and OTM values were used as the main indicators for analysis of the comet assays (Collins 2004). Of these parameters, OTM as the fraction of DNA in the tail multiplied by the distance between the means of the head and tail distributions more comprehensively reflects the degree of DNA damage than do the tail length and percentage of DNA indicators (Olive and Banáth 2006). In the present study, DNA migration in assayed cells was quantified based on the OTM values, and five classes of OTM damage, ranging from 0 (no tail) to 4 (almost all DNA in tail) (Liao et al. 2009), were then used to provide high quantitative resolution of PM-induced DNA damage. As illustrated in Fig. 5a, exposure to PM_{1.0} and PM_{2.5} caused statistically significant dose-dependent increases of the OTM in A549 cells, and PM_{1.0} was associated with higher OTM values than PM_{2.5} at each exposure concentration, which is in qualitative agreement with the trends observed in Fig. 4. Furthermore, as shown in Fig. 5b, a shift from a low DNA damage category (OTM-1) to a high DNA damage category (OTM-2) was observed in PM_{1.0}- and PM_{2.5}-treated A549 cells, with increasing concentration of PMs. Moreover, PM_{1.0} induced a higher portion of each OTM damage class than did PM_{2.5} at each exposure concentration. These quantitative data provide a clear explanation for the qualitative observations of the comet assays shown in Fig. 4, which further confirms that both PM_{1.0} and PM_{2.5} can induce DNA damage in A549 cells in a dose-dependent manner, and that PM_{1.0} is a more potent genotoxic inducer.

Conclusions

These results indicate that PM_{1.0} and PM_{2.5}, collected from the same sampling site and in a specific season, had similar morphological features and chemical properties. PM_{1.0} was enriched in chemicals emitted from anthropogenic processes, whereas PM_{2.5} consisted more of large, irregular mineral particles from natural sources. The biological results revealed that both PM_{1.0} and PM_{2.5} could induce significant cytotoxicity and genotoxicity in A549 cells, and that PM_{1.0} induced more extensive toxic effects. The higher cytotoxic effects of PM_{1.0} may be attributed to the combined effects of size and chemical composition, whereas the genotoxic effect of PM_{1.0} may be mainly associated with chemical species. To better understand how sources and specific constituents impact fine PM toxicity, long-term investigations of the chemical composition and in vitro biological activity of PM_{1.0} are needed.

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