

Toxicity of fine and quasi-ultrafine particles: Focus on the effects of organic extractable and non-extractable matter fractions

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1	Toxicity of fine and quasi-ultrafine particles: focus on the effects of organic extractable
2	and non-extractable matter fractions
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23

- 24 Abstract
- 25

26 To date no study has been able to clearly attribute the observed toxicological effects of 27 atmospheric particles (PM) to a specific class of components. The toxicity of both the organic 28 extractable matter (OEM_{2.5-0.3}) and non-extractable matter (NEM_{2.5-0.3}) of fine particles 29 $(PM_{2.5-0.3})$ was compared to that of $PM_{2.5-0.3}$ in its entirety on normal human epithelial 30 bronchial BEAS-2B cells in culture. The specific effect of the quasi-ultrafine fraction $(PM_{0.3})$ 31 was assessed, by comparing the responses of cells exposed to the PM_{2.5-0.3} and PM_{0.3} organic 32 extractable matter, OEM_{2.5-0.3} and OEM_{0.3} respectively. Chemically, PAH, O-PAH, and 33 N-PAH were respectively 43, 17, and 4 times more concentrated in PM_{0.3} than in PM_{2.5-0.3}, 34 suggesting thereby a predominant influence of anthropogenic activities and combustion 35 sources. BEAS-2B cells exposed to PM2.5-0.3, NEM2.5-0.3, EOM2.5-0.3 and OEM0.3 lead to 36 different profiles of expression of selected genes and proteins involved in the metabolic 37 activation of PAH, O-PAH, and N-PAH, and in the genotoxicity pathways. Specifically, 38 $OEM_{0.3}$ was the most inducer for phase I and phase II enzymes implicated in the metabolic 39 activation of PAH (AHR, AHRR, ARNT, CYP1A1, CYP1B1, EPHX-1, GSTA-4) thereby 40 producing the highest DNA damage, felt by ATR and, thereafter, a cascade of protein 41 phosphorylation (CHK1/CHK2/MDM2) closely related to the cell cycle arrest (P21 and P53 42 induction). This study underlined the crucial role played by the organic chemicals present in 43 $PM_{0,3}$. These results should be considered in any future study looking for the main chemical 44 determinants responsible for the toxicity of ambient fine PM.

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Keywords: fine and quasi-ultrafine particles, organic extractable matter, physicochemical
characteristics, toxicity, metabolic activation, genotoxicity.

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49 List of abbreviations:

50 1-NPyr: 1-nitropyrene; 9-FluO: 9-Fluorenone; Ace: Acenaphthalene; Acy: Acenaphthylene; 51 Ant: Anthracene: BaA: Benz[a]anthracene; BaP: Benzo[a]pyrene; BbF: 52 Benzo[b]fluoranthene; BghiP: Benzo[g,h,i]perylene; BkF: Benzo[k]fluoranthene; Chr: 53 DahA: Dibenz[a,h]anthracene; Fla: Fluoranthene; Flu: Fluorene; Chrysene; InPy: 54 Indeno[1,2,3-c,d]pyrene; N-PAH: nitrated-PAH; Nap: Naphthalene; OEM: organic 55 extractable matter; NEM: non-extractable matter; O-PAH: oxygenated-PAH; PAH: polycyclic 56 aromatic hydrocarbons; Phe: Phenanthrene; PM_{2.5-0.3}: particles with an equivalent 57 aerodynamic diameter between 0.3 and 2.5 µm; PM_{0.3}: particles with an equivalent

- 58 aerodynamic diameter below 0.3 μm (quasi-ultrafine particles); Pyr: Pyrene; ROS: reactive
- 59 oxygen species.
- 60

61 1. Introduction

Nowadays, air pollution represents one of the major risk factors for human health: 62 91% of the world's population live in places where air quality exceeds world health 63 organization (WHO) guideline limits and 7 million premature deaths are related to air 64 65 pollution each year (WHO, 2018). An important part of this threat is due to the specific presence in the atmosphere of particles having an equivalent aerodynamic diameter (EAD) 66 67 lower than 2.5 µm, also called fine fraction (PM_{2.5}). This latter is classified as carcinogenic 68 group I for humans by the international agency for research on cancer (IARC) (Loomis et al., 69 2013). $PM_{2.5}$ includes the ultrafine fraction ($PM_{0.1}$) which has the ability, in addition to the 70 high deposition potential in the lungs, to translocate to other parts of the body (Stone et al., 71 2017). Many studies have assessed the toxicity of ambient PM_{2.5}, but much less were 72 interested specifically in ambient PM_{0.1}, mainly due to the technical limitations regarding its 73 sampling and handling. Available studies have generally focused on the toxicity of ultrafine 74 particles related to a well-defined source. Baldauf et al. (2016) have evaluated the toxicity of 75 diesel exhaust particles, which include not only ultrafine particles, but also particles of other 76 diameters, which puts in place an uncertainty to discriminate the respective roles played by 77 the different size fractions. Loxham et al. (2015) were interested in studying the toxicity of 78 size-fractionated underground railway PM. Bendtsen et al. (2018) evaluated the *in-vivo* effects 79 of ultrafine particles related to airport emissions and some other authors such as Farina et al. 80 (2019) reported the effects of biomass burning ultrafine particles. Few studies were dealing 81 with airborne ultrafine PM originated from several source as it is encountered in the ambient 82 air. Between them, we can cite the works done by Carter et al. (2018) considering ambient 83 ultrafine particles retrieved from a filter using water, sonication and lyophilization, as well as 84 the study performed by Gualtieri et al. (2018) who used an air liquid interface exposure 85 system and a cut-off cyclone to expose cells to PM with a diameter estimated in the 8.9 nm to 86 1.5 µm size range.

87 Ambient PM_{2.5} is a mixture of particles emitted by natural and anthropogenic sources, 88 and in most of the cases during combustion processes related to engine exhausts and biomass 89 burning (Cheng et al., 2015). It forms a heterogeneous complex of inorganic components (e.g. metals, salts), organic ones (e.g. polycyclic aromatic hydrocarbons: PAH, polychlorinated 90 91 dibenzo-p-dioxins (PCDD), polychlorinated dibenzofurans (PCDF), polychlorobiphenyls 92 (PCB)), and biological materials (e.g. pollen, bacteria, fungi). The assumption that its toxicity 93 depends mainly on its composition is a widely accepted hypothesis (Kelly and Fussell, 2012). 94 However, both the chemical composition and the particle size are widely dependent of the

95 geographic location, meteorological conditions and emission sources, rendering thereby highly complex the identification and the prevision of their toxic effects (Loomis et al. 96 97 2013). To date, a lot of researches have been devoted to the toxicological responses induced 98 by airborne PM exposure. However, a lack of understanding remains about the specific role of 99 PM chemical components and/or fractions involved in the PM toxicity. As well, there is a 100 growing body of evidence that several parameters should be considered together, such as 101 composition, origin or size of PM, to explain the response a cell models exposed to 102 atmospheric PM (Abbas et al., 2018). It was also intended to gain information on the 103 toxicological effects induced by PM_{2.5} collected under rural, industrial or urban influences, in 104 various seasons and having various physicochemical characteristics (Billet et al., 2018; 105 Dergham et al., 2012, 2015; Melki et al., 2017).

106 Indeed, many studies have confirmed that heavy metals within PM were harmful to 107 human health (de Kok et al., 2006). Among them, transition metals, such as Cr, As, Cd and Ni 108 have been listed as carcinogenic to human (IARC, 2012). They can also induce cells to 109 overproduce reactive oxygen species (ROS), which contribute to the lipid peroxidation of the 110 cell membrane, DNA and protein damage, and even lead the cells to the death (Chen and 111 Lippmann, 2009; Fortoul et al., 2015; Zhang et al., 2018). Among the organic constituents of 112 PM, PAH are of great interest. They are mainly introduced in the atmosphere during 113 combustion process and can be subject to atmospheric physicochemical reactions resulting in 114 the formation of PAH derivatives such as nitrated PAH (N-PAH) and oxygenated PAH (O-115 PAH) (Abbas et al., 2018). Concentrations of the latter in airborne PM can be very low, but 116 their toxic effects are often considered as higher than those of their parent compounds. 117 Several works have shown that the PM organic extractable matter (OEM) contains toxic and 118 carcinogenic compounds playing a role in the overall PM toxicity without being clearly able 119 to identify the underlying mechanisms (Alessandria et al., 2014; Xiang et al., 2018). 120 Accordingly, PM organic constituents could directly contribute to the induction of an 121 oxidative stress and an inflammatory response in pulmonary cell models as well indirectly 122 after their metabolization (Abbas et al., 2019; Longhin et al., 2013, 2016; Zhou et al., 2014). 123 Even so, complementary works focusing on the critical role of the different PM constituent 124 appears necessary to better understand the PM toxicity. Hence, in the present work, airborne 125 PM with an EAD between 0.3 and 2.5 μ m (PM_{2.5-0.3}) and quasi-ultrafine (PM with EAD < 0.3 126 µm, PM_{0.3}) particles were considered. This study aimed at comparing the *in vitro* biological 127 effects and, especially, the relative genotoxic effects of both OEM of PM_{2.5-0.3}, (OEM_{2.5-0.3}), 128 non-extractable matter (NEM) of PM_{2.5-0.3} (NEM_{2.5-0.3}) and PM_{2.5-0.3} considered in its entirety

- using a relevant normal human epithelial bronchial BEAS-2B cell model. The specific effect
- 130 of the quasi-ultrafine fraction was also assessed, by exposing cells to the $PM_{2.5-0.3}$ and $PM_{0.3}$
- 131 organic extractable matter, also called OEM_{2.5-0.3} and OEM_{0.3}, respectively.
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2. Materials and Methods

136 2.1. PM Sampling

137 The sampling campaign was carried out in the southern suburb of Beirut, Lebanon 138 (Figure S1). This region is characterized by huge road traffic, electricity generation by 139 diesel-powered generators, and suffers from a problem of uncivilized incineration of 140 household waste. The southern suburb of Beirut, belonging to the district of Baabda, has the 141 largest population, reaching one million inhabitants as a result of successive migrations from 142 villages towards the capital. The sampling device was placed on the roof of the Lebanese 143 atomic energy commission (33°50'59" N; 35°29'57"E), at approximately 12 meters from the 144 street level to prevent direct influence of soil particles resuspension. This site, located on the 145 main road between the city center and Beirut airport, is a major axis of travel between Beirut and the southern part of the country. The PM sampling was done continuously from the 1st of 146 January to the 3rd of March 2017 using a high volume five stages plus back-up cascade 147 impactor (model 235, 68 m³/h, Tisch, USA), allowing the separation of collected particles, 148 149 according to their equivalent aerodynamic diameter. PM with EAD > $2.5 \mu m$ was retained on 150 a filter placed on the first stage and was not considered in this study. PM_{2.5-0.3} was impacted 151 on the different impaction plates (stage 2 to 5), while the quasi-ultrafine fraction ($PM_{0.3}$) was 152 retained on a pure quartz fiber filter (Pallflex® Tissuquartz 2500 QAT-UP, Pall, USA) placed 153 after the impaction system. The system was changed every 5 days. After sampling, impaction 154 plates were dried in a laminar flow hood during 48 h. PM_{2.5-0.3} were recovered from stages 2 155 to 5, weighted, and finally stored at -20 °C until further handling. Back-up filters were placed 156 under laminar flow hood for 48 h as well and frozen at -20°C till further use. At the end of the 157 sampling campaign, individual PM_{2.5-0.3} samples were gathered and carefully homogenized to 158 form an average PM_{2.5-0.3} sample. In a similar manner, average PM_{0.3} samples were 159 constituted by considering a proportional mix of pieces of the individual back-up filters. The 160 methodology for sampling, recovery, and homogenization of PM has been published 161 elsewhere (Borgie et al., 2016). The meteorological conditions encountered during this 162 sampling period were normal for the January and February months. As well, no dust storm 163 was observed during the sampling period as it could suddenly occur in the region.

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2.2. Physicochemical characterization of collected PM

166 Chemical characterization of PM was carried out using several methods. For both 167 PM_{2.5-0.3} and PM_{0.3} samples, total carbon was determined using CHNS/O analyzer (FLASH

168 2000, Fisher Scientific, UK), water-soluble ions were quantified by ion chromatography (IC, 169 model ICS 5000, ThermoElectron, UK) and inorganic composition (major and trace elements) 170 was quantified using inductively coupled plasma- atomic emission spectrometry (ICP-AES, 171 iCAP 6000 Series, ThermoElectron, UK) as published elsewhere (Borgie et al., 2015; Melki 172 et al., 2017). Quantification of the 16 United-States Environmental Protection Agency 173 (USEPA) priority PAH and n-alkane was carried out by gas chromatography-mass 174 spectrometry (GC-MS, model 1200 TQ, Varian, USA), after ultrasonic extraction with 175 dichloromethane (DCM) for 20 min at 20 °C. The obtained solution was concentrated under nitrogen flux until a volume of 1 mL for analysis. PCDD, PCDF, and PCB concentrations 176 177 were analyzed by Micropolluants Technology SA (Saint Julien Les Metz, France), using the 178 USEPA methods 1613 and 1668 using high resolution gas chromatography and high-179 resolution mass spectrometry (HRGC/HRMS). The concentration of 8 O-PAH and 15 N-PAH 180 was investigated by gas chromatography-mass spectrometry with negative chemical 181 ionization (GC-NCI-MS). This analysis was performed using a 7890B GC system coupled to 182 a 7000D MS detector (Agilent technologies, USA) in Negative Chemical Ionization mode 183 with a DB-5MS-UI capillary column (Agilent technologies). For PM_{2.5-0.3} only, morphology 184 and elemental composition at the particle scale was also studied by Scanning Electron 185 Microscopy (438 VP; LEO Electron Microscopy Ltd, UK) coupled with energy dispersive X-186 ray analysis (IXRF, Oxford Instruments, UK), SEM-EDX. As well, Identification of PM 187 crystalline phases was done by X-ray diffraction at room temperature (XRD AXS D8 188 Advance, Brüker, Germany) as published elsewhere (Melki et al., 2017).

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2.3. Sample preparation for toxicological studies

191 Toxicological approaches were performed on PM_{2.5-0.3}, OEM_{2.5-0.3} and NEM_{2.5-0.3} as 192 well as on the OEM_{0.3}. PM_{0.3} was collected on filters. As PM_{0.3} was trapped within the quartz 193 fibers, it was not possible to retrieve it to test the quasi-ultrafine particles (PM_{0.3}) in its 194 entirety (without filter). Consequently, only an organic extract (OEM_{0.3}) was considered for 195 studying the PM_{0,3} effects. Briefly, to prepare OEM, known masses of PM_{2,5-0,3}, and quasi-196 ultrafine particles (PM_{0.3}) collected on filters were placed in a glass vial, with 10 mL 197 dichloromethane (DCM), and subjected to a 20 min sonication assisted extraction step at 198 20 °C. After the ultrasonic treatment, the obtained solution was centrifuged (1000 g) for 199 20 min, 20 °C. Then, the supernatant solution containing extracted organics was carefully 200 removed and transferred into a second vial. Remaining PM and filters were then rinsed two 201 times using fresh DCM followed by centrifugation, and supernatant was retrieved again.

202 These extraction and rinse steps were repeated three times on the remaining PM and filters to 203 ensure the recovery of all extractable matter. The remaining PM after the extraction procedure 204 was considered as the non-extractable matter (NEM_{2.5-0.3}) and is assumed as the PM skeleton 205 on which the OEM was bound. Finally, the obtained extracts were gathered, and DCM was 206 evaporated and replaced by dimethylsulfoxide (DMSO). Hence, two organic extracts from 207 $PM_{2,5-0,3}$ (OEM_{2,5-0,3}) and $PM_{0,3}$ (OEM_{0,3}) in DMSO were obtained together with the 208 non-extractable matter of PM_{2.5-0.3} (NEM_{2.5-0.3}). Concentrations of final OEM_{2.5-0.3} and OEM_{0.3} solution were respectively 98.1 and 96.4 mg PM-equivalent/mL (PM-eq/mL). The PM-eq/mL 209 210 unit is in relation with the mass of PM extracted and the final volume of DMSO. Define 211 volumes of these OEM stock solutions were diluted in the LHC-9 culture medium to expose 212 BEAS-2B cells to the desired concentrations of OEM (expressed in µg PM-eq/cm²).

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214 **2.4.** Evaluation of the toxicity

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2.4.1. Cell culture and exposure

216 BEAS-2B cells (ATCC® CRL-9609TM, ECACC, Wiltshire, UK) were used for the 217 cytotoxicity experiments. BEAS-2B cell line originally derives from normal human bronchial 218 epithelial cells (non-cancerous individuals). BEAS-2B cells used in all the presented tests 219 derived from the same cell culture and were exposed before reaching the confluence (< 80%). 220 Cell culture method has been detailed elsewhere (Leclercq et al. 2017a). Briefly, BEAS-2B cells were grown in LHC-9 medium at 37 °C in a humidified air (relative humidity 95%, 5% 221 CO2) in 75 cm² CellBIND® surface plastic flasks (Corning, USA). Medium was refreshed 222 every 2-3 days. BEAS cells were exposed to control (LHC-9 and DMSO 0.5% in LHC-9) or 223 224 to one of the different samples (PM_{2.5-0.3}, OEM_{2.5-0.3}, NEM_{2.5-0.3} or OEM_{0.3}) suspended in 225 LHC-9 at different concentrations ranging from 3 to 96 µg PM-eq/cm². Non-exposed 226 BEAS-2B cells were used as a negative control for all experiments. In addition, 227 benzo[a]pyrene (BaP) (25 µM), 1-nitropyrene (1-NPyr) (25 µM), 9 fluorenone (9-FluO) 228 (25 µM) were used as positive controls of PAH, N-PAH and O-PAH effects respectively. 229 When exposing to OEM, the DMSO concentration in the culture medium was below 0.5%230 rendering its toxic effect insignificant (Billet et al., 2018). After 6, 24, 48 and/or 72 h of 231 exposure, aliquots of cell-free culture supernatants media were sampled and quickly frozen (-232 80 °C) till cytotoxicity assessment. Adherent BEAS-2B cells were washed twice with 1 mL 233 cold sterile PBS, then quickly frozen (-80°C) for further analyses (gene expression and genotoxicity). 234

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2.4.2. Evaluation of cytotoxicity

237 Cytotoxicity of $PM_{2.5-0.3}$, $NEM_{2.5-0.3}$, $OEM_{2.5-0.3}$, and $OEM_{0.3}$ was evaluated, after 238 exposure of cells to 3, 6, 12, 24, 48 and 96 µg PM-eq/cm² for 6, 24 and 72 h, by the 239 quantification of intracellular ATP using the Cell-Titer-Glo Luminescent cell viability Assay 240 (Promega).

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2.4.3. Metabolic activation of PAH

243 Evaluation of gene expression was carried out for several genes implicated in the PAH 244 metabolic pathway: cytochrome P4501A1: CYP1A1, cytochrome P4501B1: CYP1B1, Epoxide 245 hydrolase 1: EPHX, and Glutathione S-Transferase Alpha 4: GSTA4), in addition to aryl 246 hydrocarbon receptor: AHR, Aryl-Hydrocarbon Receptor Repressor: AHRR, and aryl 247 hydrocarbon receptor nuclear translocator: ARNT, using quantitative RT-PCR, TagmanTM (ARN18S: Hs99999901, AHR: Hs00169233_m1, ARNT: 248 gene expression assays 249 CYP1A1: Hs01121918 m1, AhRR: Hs01005075 m1, Hs01054797 g1, *CYP1B1*: 250 Hs00164383_m1, EPHX1: Hs01116806_m1, GST-A4: Hs01119249_m1), as published 251 elsewhere (Abbas et al. 2019). The fold changes (FC) in the mRNA expression levels of the 252 target genes between exposed and control BEAS-2B cells, corrected by the level of RNA18S, 253 were determined using the following equation: FC: $2-\Delta(\Delta Ct)$, where $\Delta Ct = Ct(target gene)$ -254 Ct(RNA18S gene), and $\Delta(\Delta Ct) = \Delta Ct(exposed group) - \Delta Ct(control group)$. Only FC>1.5 255 were considered as biologically significant.

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2.4.4. Genotoxicity induced by atmospheric PM

258 The genotoxicity was evaluated using the "Milliplex Map Kit- 7-plex DNA 259 damage/Genotoxicity Magnetic Bead Kit 96-well Plate" based on the Luminex xMAP 260 technology. This Milliplex Kit is used to detect changes in phosphorylated checkpoint kinase 261 1 (CHK-1) (Ser345), phosphorylated checkpoint kinase 2 (CHK-2) (Thr68), phosphorylated 262 H2A histone family member X (H2AX) (Ser139), and phosphorylated protein 53 (pP53) 263 (Ser15) as well as total protein levels of ataxia telangiectasia and Rad3-related protein (ATR), 264 mouse double minute 2 homolog (MDM2), and protein 21 (P21) in cell lysates, according to 265 the manufacturer's recommendations.

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2.4.5. Statistical analysis

Results were normalized to control, and presented as mean values and standard deviations. For each incubation time, results from BEAS-2B cells exposed to PM_{2.5-0.3},

- OEM_{2.5-0.3}, NEM_{2.5-0.3} and OEM_{2.5-0.3} or to positive controls BaP, 1-NPyr, and 9-FluO were compared to those obtained in non-exposed cells using the Mann–Whitney U-test (SPSS software). Significant differences were reported with p values <0.05.
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275 3. Results and discussion

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3.1. PM_{2.5} atmospheric concentrations

277 Concentrations of atmospheric PM_{2.5-0.3}, PM_{0.3} and PM_{2.5} were calculated taking into 278 consideration the mass of collected PM during the two months sampling period, as well as the 279 volume of air that passed through the impactor. In these conditions, the average daily PM_{2.5} 280 atmospheric concentrations, in the southern suburb of Beirut, was 54 μ g/m³. This 281 concentration largely exceeds the daily guideline value recommended by the WHO 282 $(25 \ \mu\text{g/m}^3)$ for PM_{2.5}. It was also higher than those reported by Abbas et al. (2019) at the same site in 2016 (48 μ g/m³), by Borgie et al. (2015) in the Northern suburb of Beirut in 2011 (41 283 284 $\mu g/m^3$), and by Massoud et al. (2011) in three different locations in this city in 2010 (close to 285 $20 \,\mu g/m^3$).

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3.2.1. Element and water soluble ion content

3.2. PM_{2.5-0.3} and PM_{0.3} physico-chemical characteristics

289 Table 1 represents the average concentration of major and trace elements, water 290 soluble ions and total carbon in fine and quasi-ultrafine particles. Results showed that Ca 291 $(13.4\% \text{ and } 10\% \text{ for PM}_{2.5-0.3} \text{ and PM}_{0.3} \text{ respectively}), NO_3^-$ (5.6% and 2.9%), Fe (2.6% and 292 1.2%), Al (2.1% and 1.5%), Cl⁻ (1.7% and 2.1%), Mg (1.2% and 1.8%) and Na (1.1% and 293 4.1%), were the major constituents, in addition to SO_4^{2-} (2.7%) in PM_{2.5-0.3} and NH₄⁺ (1.7%) in PM_{0,3}. Then, concentrations of K (0.5% and 0.7%), Cu (0.2% and 0.3%), Ti (0.2% and 294 295 0.1%) and P (0.1% and 0.1%) were between 0.1% and 1%, in addition to NH_4^+ (0.1%) and Ba (0.1%) in PM_{2.5-0.3} and SO₄²⁻(0.7%) and Zn (0.1%) in PM_{0.3}. Finally, the other species, Zn, 296 297 Mn, Sr, Pb, Cr, V, Sn, Ni, Mo, Co, As, and Cd showed concentrations below 0.1% for both 298 PM_{2.5-0.3} and PM_{0.3}, in addition to Ba in PM_{0.3}. The presence of elements such as Al, Ca, Fe, 299 Mg, Ca, and Ti within the atmospheric particles can be related to the resuspension of soil dust 300 (Wedepohl, 1995). Particularly, the predominance of Ca element could be related to the soil 301 type in Lebanon, which is typically calcareous (Verheye, 1973). It was also confirmed by 302 XRD analysis that evidenced the presence of calcite (Figure S2). Calcium was also found in 303 the form of gypsum, as confirmed by individual particle analysis done by SEM-EDX analysis, 304 evidencing Ca and S containing particles (Figure S3). Using XRD, Al was also detected in the 305 kaolinite form, which is one of the minerals commonly identified in Lebanese soils (Al-306 Zubaidi et al., 2011).

307 Minerals and metals are generally considered as the main components of PM collected 308 from roadsides as a result of the re-suspension by traffic of road dust and particles generated by the wear of mobile parts of vehicles, such as tires, brake or clutch disk (Putaud et al.,
2004). Indeed, trace elements showing the influence of vehicular emissions were also detected
(Ba, Ni, and Zn) (Gietl et al., 2010).

NO₃⁻ and SO₄²⁻, the major ions detected in PM, can be originated from gas to particle conversion of NO_x and SO₂ precursor gases as well as from long range transport (Borgie et al., 2016; Luria et al., 1989). Indeed, in Lebanon, it was estimated that road transport contributes to 52% of the NO_x emissions. In addition, SO₂ emissions are mainly related to industrial activities and electricity production from the combustion of heavy fuel oil, with sulfur content between 25,000 and 30,000 ppm. The latter are also used in domestic generator sets and diesel heavy vehicles (buses and trucks) (Waked et al., 2012).

The concentration of total carbon was about 7-fold higher in $PM_{0.3}$ compared to PM_{2.5-0.3} (126 vs 17.5 mg/g). This highest concentration in the quasi-ultrafine particles may be linked to the contribution of soot known to belong to this size fraction in relation with the combustion process largely influenced by the huge road traffic close to the sampling site.

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3.2.2. Organic composition

325 The determination of the 16 USEPA priority PAH showed that PM_{2.5-0.3} and PM_{0.3} had 326 substantially similar profiles with an important representation of the higher molecular weight 327 PAH (i.e., 4 aromatic rings or more) (Table 2). The concentration of PAH found in PM_{0.3} was 328 about 43 times higher than in PM_{2.5-0.3}, proving thereby a major influence of anthropogenic 329 activities and combustion sources (industries, road traffic and electric generators) on the 330 emission of quasi-ultrafine particles. Among them, dibenz[a,h]anthracene, 331 benzo[b]fluoranthene, indeno[1,2,3-cd]pyrene, benzo[a]pyrene, and chrysene were 332 predominant. They are usually emitted by combustion sources including gasoline motor 333 vehicles, biomass burning, and industrial process (INERIS, 2005). To investigate the PAH 334 emission sources, PAH diagnostic ratios were calculated and compared with those available in 335 the current literature (Table 3). The Fla/(Fla+Pyr) ratio value was ranging from 0.4 to 0.5 336 indicating fossil fuel combustion as a source of PAH. The (Fla/Pyr) ratio higher than 0.6 (0.92 337 and 0.79 respectively for PM_{2.5-0.3} and PM_{0.3} as well as the BaP/(BaP+Chry) higher than 0.35 338 (0.43 and 0.59) suggested the influence of traffic emissions. Finally, the Flu/(Flu+Pyr) ratio 339 values, 0.44 and 0.48 respectively, were slightly lower than 0.5 indicating a major influence 340 of gasoline emissions versus diesel ones. The use of BaP-toxic equivalent factor (BaP-TEQ) 341 to PAH concentrations can provide an indicator of the carcinogenic risk related to PAH 342 environmental exposure. PAH concentrations were converted in BaP-TEQ concentration using the factors published by (Kumar et al., 2018) and (Nisbet and LaGoy, 1992). This calculation showed that the considered $PM_{2.5}$ ($PM_{2.5-0.3} + PM_{0.3}$) had a huge carcinogenic potential with 8.2 ng/m³ BaP-TEQ. This value is clearly higher than the BaP WHO guideline (0.12 ng/m³; cancer risk 1/100 000). The highest BaP-TEQ values were for dibenz[a]anthracene (3.9 ng/m³ BaP-TEQ) followed by BaP (2.5 ng/m³ BaP-TEQ) and indeno[1,2,3-cd]pyrene (0.7 ng/m³ BaP-TEQ).

349 Moreover, six O-PAH and nine N-PAH were detected in PM (Table 2). Total concentration of O-PAH was higher than those of N-PAH as already reported in other studies 350 351 (Abbas et al., 2018; Albinet et al., 2007). In addition, concentrations of O-PAH and N-PAH 352 were respectively 17 and 4-fold higher in PM_{0.3} compared to PM_{2.5-0.3}. Among O-PAH 353 congeners, 1,8-naphtamic anhydre, 7H-benzo(de)anthracen-7-one, and 9,10-anthraquinone 354 were the most abundant in PM_{2.5} (45, 28 and 21 μ g/g, respectively). The 2-nitroanthracene 355 and 7-nitrobenz(a)anthracene were the most abundant nitro-PAH (4.9 μ g/g and 3.7 μ g/g 356 respectively). The emission sources of these compounds are still not well explored in the 357 literature; O-PAH and N-PAH also originate as direct or indirect products of incomplete 358 combustion (Abbas et al., 2018). Indeed, 1 nitropyrene is the most abundant pyrene in diesel 359 exhaust emissions (Bamford et al., 2003). Some N-PAH were used in industries such as the 360 5-nitroacenaphthene, which is a an intermediate in the synthesis of naphthalimide dyes that 361 are used in biomedical applications or as fluorescent whitening agents and photochemical 362 agents (WHO, 2003; Zhang et al., 2016). Concentrations and profiles of n-alkanes are 363 presented in Figure S4. As observed for PAH and PAH derivatives, quasi-ultrafine particles 364 were as well more concentrated in n-alkanes (10-fold higher) than fine ones. Two different 365 distribution profiles were evidenced with a main difference for C29 and C31. Odd n-alkanes 366 are known to mainly derive from natural origins whereas even ones can be related to 367 anthropogenic origins (Waples, 1985). The carbon maximum number (C_{max}) has been used to 368 evaluate the sources of n-alkanes (Chen et al., 2014). Higher value of Cmax indicated that n-369 alkanes primary originated from terrestrial plants, while lower C_{max} was associated with anthropogenic sources. For fine particles, Cmax was 31, while for quasi-ultrafine particles, it 370 371 was 24, showing the influence of anthropogenic emissions in the latter's composition. The 372 carbon preference index (CPI), can be used to determine the sources of n-alkanes and was 373 calculated as published in Melki et al. (2017). Values of overall CPI₂₁₋₃₁ and high CPI were 374 respectively 1.35 and 1.65 for the fine particles and 1.06 and 1.41 for the quasi-ultrafine 375 particles. These ratio value suggested that alkanes were originated from both natural and 376 anthropogenic sources (1.5 < high CPI < 3 and overall CPI > 1) in fine particles, while they

mainly derived from anthropogenic sources (high CPI < 1.5 and overall CPI close to 1) in
quasi-ultrafine ones (Figure S4).

379 The total concentrations of PCDD, PCDF and PCB were respectively 30.7 ng/g (1100 380 fg/m³), 40.4 ng/g (1170 fg/m³), and 27.9 ng/g (770 fg/m³) in fine particles. It should be 381 noticed that concentrations of these compounds were higher in the quasi-ultrafine fraction 382 than in the fine one (PCB: 15.7 vs 12.2 ng/g, PCDD: 25.2 vs 5.5 ng/g, and PCDF: 26.4 vs 383 14 ng/g respectively) (Table 2). By comparing the values found in this study with previous 384 published ones, the concentrations of PCDD and PCDF were almost twice higher than those 385 determined at the same site in 2015 by Abbas et al (2019): 1100 vs 449 fg/m³ and 1170 vs 560 386 fg/m³ respectively. Furthermore, the concentration of dioxins and furans were respectively 387 about 4.4 (770 vs 214 fg/m³) and 13-fold (1170 vs 87 fg/m³) higher than that found in 2015 in 388 the northern suburb of Beirut by Borgie et al. (2015). These higher concentrations could be 389 related to the worst solid waste management crisis in Lebanon, since 2015, when open air 390 waste burning has become a common practice in Beirut (Baalbaki et al., 2018).

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3.3. Evaluation of PM toxicity

The comparison of the effects of $OEM_{2.5-0.3}$ and $NEM_{2.5-0.3}$ *vs* $PM_{2.5-0.3}$ on the one hand, and $OEM_{0.3}$ *vs* $OEM_{2.5-0.3}$ on the other hand, were thereafter carried out to better investigate the critical role of chemicals related to a specific size fractions and/or to the extractable/nonextractable fractions involved in the cellular and molecular damages induced by PM. Cytotoxicity, pathways of metabolic activation of the PAH and other related compounds, as well as pathways of genotoxicity-DNA damage were also the critical toxicological endpoints under study.

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3.3.1. Cytotoxicity

402 Evaluation of cytotoxicity was carried out using the quantification of intracellular ATP 403 (Figure S5). The exposure of BEAS-2B cells to PM_{2.5-0.3} and NEM_{2.5-0.3} showed a significant decrease in ATP after 6 and 24 h of exposure to 6, 12, 24, 48 and 96 µg/cm² (dose-404 405 dependent). For the time corresponding to 72 hours of exposure, ATP concentrations slightly 406 increased when BEAS-2B cells were exposed to low and moderate concentrations (3, 6, 12, 407 24 and 48 μ g PM-eq/cm²) and decreased significantly when exposed to the highest concentration of PM_{2.5-0.3} (96 µg/cm²) and NEM_{2.5-0.3} (48 and 96 µg/cm²). The increase of 408 409 ATP concentrations in exposed cells for longest time (> 24 hours which is the minimum time 410 required for cell division) is related to the proliferation of cells which is naturally

411 accompanied by an increase of ATP concentration. Decrease of ATP concentration indicated 412 a mitochondrial dysfunction and proved the ability of $PM_{2.5-0.3}$ to induce toxicity, even at the 413 lowest doses and the shortest times. A more pronounced effect was observed for the highest 414 doses. The toxicity of the OEM $_{0,3}$, after 72 h of exposure to the two highest concentrations (48 415 and 96 μ g PM-eq/cm²) was the most remarkable. The ATP concentration decrease was 416 significantly higher for cells exposed to $OEM_{0,3}$ in comparison with cells exposed to 417 $OEM_{2.5-0.3}$ (p<0.020 and p<0.034 for the two doses respectively). This could be explained by 418 the highest concentration of PAH and organic compounds in PM_{0.3} compared to PM_{2.5-0.3}.

Based on the above-detailed results, two concentrations of exposure ensuring sufficient doses to observe some harmful effects without a massive cell death were selected for the study of the other toxicological endpoints. Indeed, the two considered doses, 3 and $12 \mu g PM$ -eq/cm² were also among the lowest reported in the literature to give harmful effects whilst limiting a massive cell death (Boublil et al. 2013, Leclercq et al. 2016, 2017b, Gualtieri et al., 2010, Longhin et al. 2013).

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3.3.2. Metabolic activation

427 PAH are known to enter into the cell through cytoplasmic membrane before binding to 428 the aryl hydrocarbon receptor (AHR). Following ligand binding, the PAH-AHR complex 429 translocates to the nucleus and forms a heterodimer with its nuclear translocator (aryl receptor 430 nuclear translocator, ARNT). This latter then binds to a specific region of the DNA called 431 xenobiotic response element (XRE), recruits the RNA polymerase II and induces the 432 transcription of several genes, included in the AhR gene battery. It is also implicated in the 433 metabolic activation of PAH into more easily eliminable metabolites (Puga et al., 2009). 434 Among them, phase I enzymes (CYP1A1 and CYP1B1) and phase II enzymes (EPHX-1 and 435 GSTA-4) are closely involved (Dietrich, 2016; Rogers et al., 2017). The main function of the 436 AHR signaling pathway is to activate parent PAH and thereafter to conjugate their highly 437 electrophilic intermediate metabolites with highly hydrophilic groups in order to easily 438 eliminate them from cells and organism thereby avoiding severe toxic effects.

In this study, the expression of some selected genes involved in the AhR signaling
pathway was evaluated after exposure of BEAS-2B cells to PM, NEM and OEM at 3 and 12
µg PM-eq/cm² for 6 and 24 h.

Exposure of BEAS-2B cells to PM_{2.5-0.3}, NEM_{2.5-0.3}, EOM_{2.5-0.3} and EOM_{0.3} revealed different gene expression profiles of some key proteins involved in the metabolic activation of PAH (Table 4). PM_{2.5-0.3} in its entirety was able to induce significant gene expressions of 445 AHR, AHRR, ARNT, CYP1A1, CYP1B1 and GSTA-4, 6 and 24 h after cell exposure to the two 446 doses. Results obtained for positive controls (BaP, 1-Npyr and 9-FluO) confirmed the 447 activation of the AHR signaling pathway by exogenous chemicals in BEAS-2B cells. In 448 general, metabolic activation helps to decrease the toxicity of pollutants. However, this 449 metabolization can formed metabolites with higher toxicity than initial molecules (Saint-450 Georges et al., 2008, 2009). Gualtieri et al. (2011) also reported that the AHR signaling 451 pathway was activated in BEAS-2B cells exposed to PM_{2.5} and PM₁₀ collected in urban background area in relation with their high PAH contents. In contrast, no significant induction 452 453 of these genes was observed in BEAS-2B cells exposed to NEM_{2.5}, whatever the tested dose 454 and time. In comparison, Billet et al. (2006, 2008) have used PM having undergone a thermal 455 desorption under vacuum (dPM) before studying the metabolic activation. The technique used 456 preserve the PM inorganic structure while organic chemicals were desorbed. Somewhat 457 surprising result was the effective transcription of CYP1A1 gene cells derived from a 458 type II-like human alveolar epithelial carcinoma (A549 cell line) exposed to dPM, thereby 459 supporting that the 400°C thermal desorption under vacuum was not enough sufficient for 460 removing totally the organic fraction. In contrast, as NEM_{2.5-0.3} did not significantly induce 461 metabolic activation, the method used here seemed to be more efficient to recover both OEM 462 and organics almost free NEM.

463 For OEM_{2.5-0.3}, a significant increase of the gene expression of CYP1A1 and CYP1B1 464 was observed only 6 h after the exposure and only for the highest dose. Accordingly, Oh et al. 465 (2011) indicated that OEM which contain aliphatic chlorinated hydrocarbons, PAH/their alkyl 466 derivatives and nitro-AH/ketones/quinones, induced CYP1A1-dependent EROD activity, 467 through AHR signaling pathway activation. After 24 h of exposure to OEM_{2.5-0.3}, no 468 significant gene expression was observed. In comparison, PM_{2.5-0.3} was able to induce 469 CYP1A1 and CYP1B1 gene expression after 6 h of exposure at the lowest dose and after 24h 470 of exposure as well. One explanation could be related to both the kinetics of metabolic 471 activation, the difference regarding the bio-accessibility of PAHs in PM_{2.5-0.3} and OEM_{2.5-0.3} 472 (these points will be further discussed below) and the contribution of some metallic species 473 only present in PM_{2.5-0.3}. For this latter, Garçon et al. (2004a, 2004b) reported that the 474 concurrent exposure to Fe₂O₃ particles and BaP favors dramatically the metabolic activation 475 of PAH by phase I enzymes (i.e., CYP1A1) to the detriment of the inactivation of PAH 476 metabolites by phase II enzymes (i.e., GST-YA). They finally concluded that the kinetics of 477 the adverse effects might be different depending on the presence or absence of BaP on Fe₂O₃ 478 particles. Korashy et al. (2005) also reported that *CYP1A1* gene expression could be induced by heavy metals (Korashy and El-Kadi, 2004). Nevertheless, the NEM_{2.5-0.3} fraction was not
able to induce metabolic activation allowing to reject this hypothesis.

481 Regarding the quasi-ultrafine fraction, OEM_{0.3} was able to induce a significant 482 expression of all tested genes involved in the AHR signaling pathway 6 and 24 h after 483 exposure, mainly to the highest tested dose (Table 4). Compared with OEM_{2.5-0.3}, OEM_{0.3} led 484 to a higher expression of the proteins involved in the AHR signaling pathway. This could be 485 closely related to the much higher contents of PAH and other related compounds in OEM_{0.3} compared with OEM_{2.5-0.3} (till 43 times higher for PAH). This result is in agreement with a 486 487 recent study showing the significant induction of the two AHR-related CYP, CYP1A1 and 488 CYP1B1, in BEAS-2B cells 6 and 24h after their exposure to increasing concentrations of 489 PM_{2.5} OEM, ranging from 3 to 30 µg PM-eq/cm² (Abbas et al., 2019). Using the same 490 concentrations as those used in this study, Borgie et al. (2015) reported also the effective up-491 regulation of the gene transcription of CYP1A1 and CYP1B1, 24, 48, and 72 h after the 492 exposure of BEAS-2B cells to $OEM_{0.3}$.

493 Taken together, these results supported not only the significant activation of the AHR 494 signaling pathway by the PAH but also put into question both the bioavailability of PM-495 linked-PAH vs PAH in an organic extract, the kinetics of the metabolic activation and the 496 dose parameter. PM_{2.5-0.3} and OEM_{2.5-0.3} have the same PAH composition. Nevertheless, 497 CYP1A1 and CYP1B1 expression were not the same for both samples. Especially, no 498 expression was observed for 6 h of exposure at the lowest dose for OEM_{2.5-0.3}, as well at 24 h, 499 for all the doses. In OEM_{2.5-0.3}, PAH are highly bio-accessible compared to PAH linked to 500 PM_{2.5-0.3}. This higher bio-accessibility could be responsible for a quicker metabolic activation 501 of PAH. It can be suggested that this one took less than 6 hours to be completed especially 502 when doses (and PAH concentrations) are low. This hypothesis is reinforced by the fact that 503 no metabolic activation was observed at 24 hours as well. The early metabolic activation of 504 PAH was also observed by Totlandsdal et al. (2010) who have studied the CYP1A1 expression 505 of BEAS-2B cells exposed to Diesel Exhaust Particles (DEP) for 2, 4 and 8 h. They found that 506 the CYP1A1 gene expression occurred at early time with a pronounced increase after 2 hours 507 of exposure and a levelling off at 4 and 8 h. Then, after 24 h, the level of this gene was lower 508 than the one observed at 8 h. Also testing DEP but considering HBEC3 cells, Rynning et al. 509 (2018) measured a CYP1A1 gene expression after 24 hours. It was the shortest time 510 considered in their study. However, Brinchmann et al. (2018) have considered DEP exposure 511 using a 3D tri-culture (A549/THP-1/EA.hy926) model for 2 h and 20 h. They have also 512 studied the DEP-OEM exposure using HMEC-1 cells for 2 and 24 h. They observed a

CYP1B1 significant expression in the epithelial side of the 3D model after only 2 h of 513 514 exposure at DEP. This gene expression was also much higher after 20 h of exposure 515 compared to 2 h. DEP-OEM exposed HMEC-1 cells also show a significant CYP1A1 and 516 CYP1B1 gene expression after 2 h, staying quite constant after 24 h of exposure. Taken 517 together these results are in agreement with an early metabolic activation of PAH that can be 518 spread out over time especially when PAH are linked to PM (reduced accessibility and slow 519 release). The fact that the metabolization of PAH in PM_{2.5-0.3} appears to be time-shifted and 520 spread-out compared to PAH in OEM_{2.5-0.3} is also consistent with this statement. For higher 521 PAH content such as that in OEM_{0.3}, the metabolization of PAH seems to be extended 522 compared to $OEM_{2.5-0.3}$.

523 Finally and in agreement with previous works done by Garçon et al. (2004a, 2004b), 524 these results involve also two hypotheses based on the bioavailability and bioaccessibility of 525 organic compounds, whether those linked to PM, or found in the organic extract. The first one 526 support the idea that particles represent a vector which facilitates penetration of organic 527 compounds into the cell by endocytosis (Chen et al., 2016). The second hypothesis is based 528 on the lipophilic properties of organic compounds. They can easily and even quickly penetrate 529 the cell through the cytoplasmic membrane to be freely metabolized within the cell. 530 Accordingly, Brinchmann et al (2018) reported that the fast induction of AhR-regulated genes 531 in the basolateral endothelial cells of their alveolar 3D tri-culture model exposed to relevant concentrations of DEP (0.12 μ g/cm²) seemed mainly to be due to the soluble lipophilic 532 533 constituents rather than particle translocation. Indeed, DEP-EOM, next fractionated with 534 increasing polarity, chemically characterized, and examined for direct effects on 535 AhR-regulated genes in human microvascular endothelial cells and primary human 536 endothelial cells, supported that the most lipophilic EOM, that contained the majority of AhR 537 agonists such as PAH, had the clearest low-concentration effects. The different kinetics 538 observed for PM and OEM could also be related to the fact that PAH linked to PM could be 539 only progressively released in cell cytoplasm after particle internalization, and subsequently 540 metabolized, while free PAH in the organic extract can be metabolized directly once in the 541 cell.

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3.3.3. Evaluation of genotoxicity biomarkers

544 Given the complexity of atmospheric particles, the identification of the chemical 545 compounds responsible for the genotoxic effect is still not clear. Several studies suggested the 546 role of PAH, and therefore of the organic fraction, while other studies suggested the role of 547 water soluble (Prahalad et al., 2001) and non-water soluble metals (Ghio, 1999; Knaapen et 548 al., 2002) in the initiation of oxidative stress and DNA damage. The different DNA lesions 549 can be caused by a wide variety of agents. If they are still present during replication, tolerance 550 mechanisms can be put in place. For each type of lesion, specific repair mechanisms exist. 551 Thus, to maintain the integrity of the genome, eukaryotic cells have established signaling 552 pathways that are activated in response to DNA damage. In case of DNA damage, the ATR 553 and ATM sensors phosphorylate a number of downstream proteins, such as CHK1, CHK2 and 554 P53, in addition to the activation of MDM2 and P21, allowing the cell cycle to be stopped, in 555 order to provide the cell to repair its DNA with time, but also to induce apoptosis in the case 556 of cells with serious lesions (Asimuddin and Jamil, 2012).

557 Proteins involved in the genotoxicity pathway including phosphorylated ones, 558 p-CHK1, p-CHK2, p-H2AX, p-P53, and total proteins, ATR, MDM2, and P21, have been 559 tested after exposure of BEAS-2B cells to the different fraction of PM_{2.5-0.3} and OEM_{0.3} for 24 560 and 72 h. A very weak induction of the proteins involved in this pathway following the 561 exposure of BEAS-2B cells to the different fractions of PM_{2.5-0.3}, including NEM_{2.5-0.3} (Table 562 5) was observed. This indicates that most of the biologically active metabolites resulting from 563 the metabolic activation of PM, their organic extract and the non-extractable fraction, are all 564 conjugated during the phase II of this process and removed from the cell. However, different 565 results were reported with OEM_{0.3}. The highest dose of OEM_{0.3} was able to induce a 566 significant increase of ATR, p-CHK1 and p-CHK2, MDM2, P21, and p-P53, almost only 567 after 72 h (Table 5). EOM_{0.3} contains relatively high proportion of organic compounds 568 compared to OEM_{2.5-0.3}. It is clearly observed that EOM_{0.3} induced DNA damage felt by ATR 569 and followed by a cascade of protein phosphorylation, thereby contributing to the cell cycle 570 arrest (i.e., P21 and P53 induction). Accordingly, higher levels of phosphorylated P53, CHK1 571 and CHK2 were generally related to the cell cycle arrest (Fragkos et al., 2009).

572 In addition, this study showed a significant decrease in phosphorylated H2AX in 573 BEAS-2B exposed for 72 h to the two doses of OEM_{2.5-0.3}. In addition, the same decrease was 574 observed after 72 hours of exposure to NEM_{2.5-0.3} at 3 and 12 μ g PM-eq/cm², and to 575 $3 \mu g PM$ -eq/cm² of OEM_{0.3}. These results are in agreement with those published by Abbas et 576 al. (2019) and Bouquet et al. (2006) showing that, at low doses of exposure, the DNA repair 577 mechanism was highly active leading to the degradation of phosphorylated H2AX. They also 578 observed that, at high doses of exposure, the DNA repair mechanism seemed to be less active 579 leading to phosphorylated H2AX accumulation, which is also the case in this study as shown 580 for OEM_{0.3}, 72 h and 12 μ g PM-eq/cm².

With regards to the different chemical compounds tested as positive controls, BaP was able to activate the DNA damage pathway by inducing the expression of several studied proteins including p-CHK2, p-H2AX, and p-P53, after 24 h, and p-CHK1 and p-CHK2 after 72 h. Moreover, 1-NPyr significantly induced expression of p-CHK2 and p-P53, after 24 h, and of p-P53 after 72 h. In contrast, cells exposed to the 9-FluO didn't show any significant expression of the studied proteins.

587 Taken together, these results showed that the exposure to OEM_{0.3} was able to induce 588 the expression of most of the studied proteins related to cell cycle arrest and DNA damages. 589 In addition, these results showed that BaP and 1-NPyr were able to induce DNA damages, 590 while 9-FluO, the positive control used for O-PAH toxicity evaluation was not.

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592 4. Conclusion

593 In the present work, PM was collected at the southern suburbs of Beirut. Chemical 594 analysis clearly showed the influence of soil resuspension and also traffic emissions. Organic 595 fraction analysis showed that the contents of PAH and n-alkanes were far higher in PM_{0.3} than 596 PM_{2.5-0.3}, which supports a strongly relation between the combustion process emission and the 597 ultrafine particle composition. Toxicological studies suggested that the metabolic activation of 598 PAH contained in PM_{2.5-0.3} in its entirety was delayed in comparison with PAH in OEM. The 599 NEM was not able to significantly express genes involved in the metabolic activation pathway 600 of organic compounds. All the tested PM fractions, including the NEM were able to increase 601 the expression of the studied proteins involved in the genotoxicity pathway. In addition, the 602 PM-induced genotoxicity appeared to be strongly related to the organic chemical contents. 603 The quasi-ultrafine particles were the most genotoxic due the predominance of organic 604 compounds such as PAH in this size fraction. The obtained results confirmed that considering 605 only OEM or NEM might lead to results regarding mechanisms and kinetics of toxicity that 606 would be different of those obtained from cells exposed to PM considered in its entirety. This 607 work evidenced that the co-presence of organic compounds and other inorganic elements 608 (such as metals and trace elements) may determine the biological effects of fine PM as a 609 whole. It also shows that organic compounds associated to the quasi-ultrafine fraction are the 610 main drivers of the fine PM genotoxicity and phase I and phase II enzymes implicated in the 611 metabolic activation of PAH. Finally, this study underlined the importance of evaluating the 612 biological impact of ultrafine and quasi-ultrafine particles for a better understanding of the 613 health effects related to exposure to the fine PM.

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Table captions

Table 1: Elemental, water-soluble ions and total carbon concentrations in $PM_{2.5-0.3}$ and $PM_{0.3}$ collected in the suburb of Beirut (Lebanon).

Table 2: Concentrations of PAH, N-PAH, O-PAH (in $\mu g/g$) and PCB, PCDD and PCDF (in ng/g) detected in PM_{2.5-0.3} and PM_{0.3} collected in the southern suburbs of Beirut. The relative percentage of each compound in the corresponding class of organic compounds is also indicated.

Table 3: PAH diagnostic ratios calculated for $PM_{2.5-0.3}$ and $PM_{0.3}$ collected at the southern suburb of Beirut and compared to the bibliography, in order to determine the possible emission sources of PAH.

Table 4: Metabolic activation: gene expression of aryl hydrocarbon receptor (AHR), Aryl-Hydrocarbon Receptor Repressor (AHRR) and aryl hydrocarbon receptor nuclear translocator (ARNT), cytochrome P4501A1 (CYP1A1), cytochrome P4501B1(CYP1B1), epoxide hydrolase 1 (EPHX), glutathione S-transferase alpha 4 (GSTA4) enzymes in BEAS-2B cells exposed 6 and 24 h to PM_{2.5-0.3}, organic extract matter (OEM_{2.5-0.3}), non-extractable matter fraction (NEM_{2.5-0.3}) and organic extractable matter of the quasi-ultrafine fraction (OEM_{0.3}) at 3 and 12 µg PM-eq/cm² and positive controls (BaP, 1-NPyr and 9-FluO). These values are depicted as mean values and standard deviations of 3 replicates for controls and 3 replicates for exposed cells (Mann–Whitney U-test; vs. Controls, *: P<0.05).

Table 5: Genotoxicity: Protein expression of ataxia telangiectasia and Rad3-related protein (ATR), phosphorylated checkpoint kinase 1 (p-CHK-1), phosphorylated checkpoint kinase 2 (p-CHK-2), phosphorylated H2A histone family member X (p-H2AX), total proteins of mouse double minute 2 homolog (MDM2), total protein 21 (P21), phosphorylated protein 53 (p-P53), in BEAS-2B cells exposed 6 and 24 h to PM_{2.5-0.3}, organic extractable matter (OEM_{2.5-0.3}), non-extractable matter fraction (NEM_{2.5-0.3}) and organic extractable matter of the quasi-ultrafine fraction (OEM_{0.3}) at 3 and 12 µg PM-eq/cm² and positive controls (BaP, 1-NPyr and 9-FluO). These values are depicted as mean values and standard deviations of 3 replicates for controls and 3 replicates for exposed cells (Mann–Whitney U-test; vs. Controls, *: P<0.05).

Table 1

Species	PM2.5-0.3 (mg/g)	PM _{0.3} (mg/g)
Ca	134	100
NO_3^-	56	28.8
SO 4 ²⁻	27.1	7.11
Fe	25.8	12.2
Al	21.4	14.9
Cl	16.5	21.2
Mg	12.3	17.6
Na	10.7	40.5
Κ	5.27	6.67
Cu	2.38	2.73
Ti	2.01	1.21
Ba	1.02	0.40
NH_4^+	1.01	17.4
Р	0.58	1.39
Zn	0.40	1.12
Mn	0.30	0.19
Sr	0.23	0.13
Pb	0.15	0.32
Cr	0.14	0.08
V	0.11	0.32
Sn	0.11	0.06
Ni	0.05	0.16
Мо	0.04	0.34
Со	0.14	0.01
As	<dl<sup>a</dl<sup>	0.01
Cd	<dl<sup>a</dl<sup>	0.01
Total carbon	17.5	126

^aDL : Detection limit

	Abbreviation	PM _{2.5-0.3}	Relative	PM _{0.3}	Relative
	Abbieviation	1 112.5-0.5	%	1 110.5	%
PAH (µg/g)					
Naphthalene	Nap	0.3	0.9	1.1	0.1
Acenaphthylene	Acy	0.1	0.2	0.7	0.1
Acenaphthene	Ace	0.01	0.0	0.2	0.0
Fluorene	Flu	0.1	0.2	0.5	0.1
Phenanthrene	Phe	1.2	3.3	8.8	0.9
Anthracene	Ant	1.7	4.6	1.4	0.1
Fluoranthene	Fla	2.5	6.7	19	1.8
Pyrene	Pyr	2.7	7.3	23	2.3
Benzo[a]Anthracene	BaA	3.4	9.2	64	6.6
Chrysene	Chr	3.6	9.7	51	5.2
Benzo[b]Fluoranthene	BbF	5.6	15.2	173	17.6
Benzo[k]Fluoranthene	BkF	1.5	4.0	38	3.8
Benzo[a]Pyrene	BaP	2.7	7.4	73	7.5
Indeno[1.2.3-c.d]Pyrene	InPy	4.7	12.8	206	21.0
Dibenzo[a.h]Anthracene	DahA	0.4	1.1	22.9	2.3
Benzo[g.h.i]Perylene	BghiP	6.4	17.4	299	30.5
ΣPAH	DSIII	37	100	<u>980</u>	100
O-PAH ^a (µg/g)					
1.8-naphtalic anhydre		2.3	32.9	43	40.6
7H-benz(de)anthracen-7-one	19-BAntO	0.8	11.4	27	25.5
9.10-anthraquinone	9.10-AntQ	2.5	35.7	19	17.9
9-fluorenone	9-FluO	0.7	10.0	3.3	3.1
Benzo(a)fluorenone	BaFluO	0.8	11.4	6.8	6.4
Benz(a)anthracen-7.12-dione	7.12-BaAQ	-	-	6.7	6.3
Σ О-РАН		7.0	100	106	100
N-PAH ^b (µg/g)					
2-nitroanthracene	2-Nant	0.2	4.9	4.8	31.2
7-nitrobenz(a)anthracene	7-NBaA	1.1	26.8	2.7	17.5
9-nitroanthracene	9-NAnt	0.05	1.2	1.5	9.7
6-nitrochrysene	6-NChry	0.3	7.3	1.1	7.1
1-nitropyrene	1-NPyr	0.6	14.6	1.0	6.5
1.3-dinitronaphthalene	1.3-DNNap	0.6	14.6	1.2	7.8
5-nitroacenaphthene	5-NAce	0.14	3.4	1.1	7.1
3-nitrophenanthrene	3-NPhe	0.6	14.6	1.1	7.1
2-nitrofluorene	2-NFlu	0.4	9.8	0.9	5.8
Σ Ν-ΡΑΗ		4.1	100	15.4	100
Σ PCDD (ng/g)		5.5		25.2	
Σ PCDF (ng/g)		14.0		26.4	
Σ PCB (ng/g)		12.2		15.7	

^a1-naphtaldehyde and phenanthrene-9-carboxyaldehyde were not detected.

^b1-nitronaphtalene; 2-nitrobiphenyl; 1.5-dinitronaphtalene and 1.8-dinitropyrene were not detected.

PAH diagnostic ratiosPM2.5-0.3PM0.3Characteristic ratio values and sources								
InPy /(InPy+ BghiP) ^a	0.42	0.41	0.18	0.62	0.35-0.70			
			Cars	Wood burning	Diesel burning			
Flu/(Flu+Pyr) ^a	0.48	0.44	<0.5		>0.5			
			Gasoline		Diesel			
BaP/(BaP+Chry) ^b	0.43	0.59	> 0.35	(0.2-0.35			
			Vehicular emission	Coal	combustion			
BaP/ BghiP ^a	0.42	0,25	0.5-0.6		>1.25			
			Traffic emission	Brown	coal or lignite			
Anth/(Anth+Phe) ^b	0.59	0.14	>0.1	<0.1				
			pyrogenic	petrogenic				
InPy / BghiP ^a	0.74	0.69	<0.4	~1				
			Gasoline	Diesel				
Fla/(Fla+Pyr) ^b	0.48	0.44	<0.4	0.4-0.5	>0.5			
			petrogenic	Fossil fuel	Grass, wood, coal			
				combustion	combustion			
Fla/Pyr ^b	0.92	0.79	< 0.6	> 0.6				
			Non traffic	traffic emission				
			emission					
CPAH/TPAH ^a	0.98	0.98		~1				
				combustion				

^a Ravindra et al., 2008

^b Tobiszewski and Namiesnik, 2012

Table	4:
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			PM _{2.5-0.3}		NEM _{2.5-0.3} OEM _{2.5-0.3}		1 _{2.5-0.3}	OEM _{0.3}		BaP	1-NPyr	9-FluO	
		Control	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	25 μΜ	25 μΜ	25 μΜ
	AHR	1 ± 0.02	1.96* ± 0.2	2.93* ± 0.31	0.99 ± 0.06	1.06 ± 0.15	1 ± 0.03	1 ± 0.08	1.18 ± 0.25	2.79* ± 0.38	0.9 ± 0.13	4.18* ± 0.41	277.7* ± 63.9
	AhRR	1 ± 0.08	2.10* ± 0.14	2.86* ± 0.88	0.99 ± 0.11	1.43 ± 0.33	$1.2 \\ \pm 0.30$	1.44 ± 0.46	1.19 ± 0.50	2.21* ± 0.60	0.94 ± 0.22	3.15* ± 1.05	61.3* ± 12.9
	ARNT	1 ± 0.02	1.41 ± 0.33	1.91* ± 0.53	1.13 ± 0.15	1.21 ± 0.26	$\begin{array}{c} 0.9 \\ \pm \ 0.06 \end{array}$	0.9 ± 0.13	$\begin{array}{c} 1.45 \\ \pm \ 0.44 \end{array}$	1.66* ± 0.14	$\begin{array}{c} 1.05 \\ \pm \ 0.2 \end{array}$	2.36* ± 0.54	134.8* ± 39.0
6 h	CYP1A1	1 ± 0.07	1.78* ± 0.46	3.65* ± 0.82	1 ± 0.10	1 ± 0.12	$\begin{array}{c} 0.97 \\ \pm \ 0.01 \end{array}$	1.55* ± 0.10	2.18* ± 0.75	1.48 ± 0.33	$\begin{array}{c} 1.03 \\ \pm 0.2 \end{array}$	2.28* ± 0.35	37.3* ± 3.71
	CYP1B1	1 ± 0.04	2.26* ± 0.49	4.51* ± 0.83	1.14 ± 0.38	$1.5 \\ \pm 0.50$	$\begin{array}{c} 0.98 \\ \pm \ 0.04 \end{array}$	2.37* ± 0.91	1.59* ± 0.31	2.06* ± 0.15	0.69 ± 0.26	2.55* ± 0.67	46.4* ± 16.1
	EPHX-1	1 ± 0.12	0.99 ± 0.26	$\begin{array}{c} 1.68 \\ \pm \ 0.58 \end{array}$	1 ± 0.28	1.01 ± 0.26	$\begin{array}{c} 0.87 \\ \pm \ 0.05 \end{array}$	$\begin{array}{c} 0.87 \\ \pm 0.06 \end{array}$	1.80* ± 0.13	1.75 ± 0.38	0.82 ± 0.06	2.44* ± 0.45	114.5* ± 19.5
	GSTA-4	1 ± 0.09	1.30 ± 0.18	2.59* ± 0.75	0.84 ± 0.09	$\begin{array}{c} 1.32 \\ \pm 0.30 \end{array}$	1.21 ± 0.24	1.18 ± 0.10	2.30 ± 0.87	1.71* ± 0.15	0.81 ± 0.20	3.04* ±1	202.7* ± 34.5
	AhR	1 ± 0.01	1.31 ± 0.34	4.04* ± 0.8	0.95 ± 0.06	1.03 ± 0.16	0.94 ± 0.03	0.92 ± 0.10	0.43 ± 0.08	1.67* ± 0.22	1.50* ± 0.16	2.48* ± 0.72	152.1* ± 25.4
	AhRR	1 ± 0.04	1.45 ± 0.12	4.13* ± 0.79	0.93 ± 0.14	1 ± 0.20	$\begin{array}{c} 0.86 \\ \pm 0.20 \end{array}$	1 ± 0.02	1 ± 0.07	1.64* ± 0.36	2.50* ± 0.93	1.64* ± 0.11	30.3* ± 5.23
	ARNT	1 ± 0.05	1.30 ± 0.39	3.56* ± 1.16	$\begin{array}{c} 0.92 \\ \pm \ 0.14 \end{array}$	1.2 ± 0.36	$\begin{array}{c} 1.17 \\ \pm \ 0.51 \end{array}$	1.11 ± 0.34	0.89 ± 0.14	1.98* ± 0.22	1.63* ± 0.10	$\begin{array}{c} 1.34 \\ \pm \ 0.60 \end{array}$	117.0* ± 27.7
24 h	CYP1A1	1 ± 0.04	2.59* ± 0.87	7.46* ± 2.70	$\begin{array}{c} 0.9 \\ \pm \ 0.10 \end{array}$	1.49 ± 0.42	$\begin{array}{c} 1.01 \\ \pm \ 0.12 \end{array}$	1.40 ± 0.28	2.45* ± 0.26	4.3* ± 0.97	7.70* ± 1.64	1.86* ± 0.46	18.1* ± 3.14
	CYP1B1	1 ± 0.13	1.97* ± 0.29	4.07* ± 0.68	1.19 ± 0.50	1.26 ± 0.41	1.26 ± 0.33	1.23 ± 0.41	1.09 ± 0.16	3.21* ±1	6.02* ± 1.57	1.67* ± 0.17	17.1* ± 3.16
	EPHX-1	1 ± 0.02	1.38 ± 0.32	4.04* ± 1.02	1.05 ± 0.45	1.38 ± 0.38	$\begin{array}{c} 1.000 \\ \pm 0.02 \end{array}$	1.24 ± 0.45	1.15 ± 0.43	1.69* ± 0.14	1.70* ± 0.50	1.75* ± 0.26	58.1* ± 9.65
	GSTA-4	1 ± 0.04	$\begin{array}{c} 1.03 \\ \pm 0.02 \end{array}$	3.62 ± 1.06	$\begin{array}{c} 0.84 \\ \pm \ 0.18 \end{array}$	1.17 ± 0.34	1 ± 0.03	1 ± 0.048	1.16 ± 0.27	1.65* ± 0.19	1.46 ± 0.26	1.29 ± 0.28	94.9* ± 18.6

AHR: aryl hydrocarbon receptor, *AhRR*: Aryl-Hydrocarbon Receptor Repressor, *ARNT*: aryl hydrocarbon receptor nuclear translocator, *CYP1A1*: cytochrome P4501A1, *CYP1B1*: cytochrome P4501B1, *EPHX-1*: Epoxide hydrolase 1, *GSTA-4*: Glutathione S-Transferase Alpha 4, PM2.5-0.3: fine particulate matter, NEM2.5-0.3: Non-extractable matter fraction, OEM2.5-0.3: fine organic extractable fraction, OEM0.3: quasi-ultrafine organic extractable matter, BaP : Benzo[a]pyrene, 1-NPyr: 1-nitropyrene, 9-FluO : 9-FluOrenone.

			PM	2.5-0.3	NEM	I _{2.5-0.3}	OEM	а _{2.5-0.3}	OE	M _{0.3}	BaP	1-NPyr	9-FluO
		Control	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	3 μg PM- éq/cm ²	12 μg PM- éq/cm ²	25 μΜ	25 μΜ	25 μΜ
	ATR	1 ± 0.07	1.07 ± 0.06	$\begin{array}{c} 1.08 \\ \pm \ 0.18 \end{array}$	1.02 ± 0.13	$\begin{array}{c} 1.05 \\ \pm \ 0.1 \end{array}$	$\begin{array}{c} 0.92 \\ \pm \ 0.07 \end{array}$	1.22 ± 0.29	1.014 ± 0.137	1.03 ± 0.13	1 ± 0.224	$\begin{array}{c} 1.06 \\ \pm 0.01 \end{array}$	1 ± 0.209
	p-CHK1	1 ± 0.09	0.97 ± 0.16	0.94 ± 0.13	0.94 ± 0.14	$\begin{array}{c} 0.92 \\ \pm \ 0.07 \end{array}$	$\begin{array}{c} 0.93 \\ \pm \ 0.06 \end{array}$	1.369 ± 0.517	1.038 ± 0.179	1.24* ± 0.12	1.23 ± 0.22	$\begin{array}{c} 1.16 \\ \pm \ 0.08 \end{array}$	1.03 ± 0.21
	p-CHK2	1 ± 0.09	1 ± 0.14	1 ± 0.18	$\begin{array}{c} 0.93 \\ \pm \ 0.13 \end{array}$	0.94 ± 0.09	$\begin{array}{c} 0.95 \\ \pm \ 0.09 \end{array}$	1.464 ± 0.622	0.984 ± 0.152	1.08 ± 0.13	1.34* ± 0.19	3.63* ± 0.29	1.04 ± 0.22
24 h	p-H2AX	1 ± 0.13	0.88 ± 0.25	0.756 ± 0.164	$\begin{array}{c} 0.92 \\ \pm \ 0.12 \end{array}$	0.83 ± 0.1	$\begin{array}{c} 0.95 \\ \pm \ 0.26 \end{array}$	1.110 ± 0.383	0.806 ± 0.137	$\begin{array}{c} 0.85 \\ \pm \ 0.18 \end{array}$	1.42* ± 0.08	1.19 ± 0.24	0.83 ± 0.27
	MDM2	1 ± 0.12	1.27* ± 0.11	1.34* ± 0.24	1.02 ± 0.13	1.21* ± 0.1	$\begin{array}{c} 0.92 \\ \pm \ 0.107 \end{array}$	1.251 ± 0.408	0.918 ± 0.131	1.22*	1.24 ± 0.20	$\begin{array}{c} 1.15 \\ \pm \ 0.03 \end{array}$	1.06 ± 0.25
	P21	1 ± 0.10	0.89 ± 0.2	0.55 ± 0.1	$\begin{array}{c} 0.8 \\ \pm \ 0.07 \end{array}$	$\begin{array}{c} 0.61 \\ \pm \ 0.08 \end{array}$	$\begin{array}{c} 0.88 \\ \pm \ 0.05 \end{array}$	1.065 ± 0.215	0.881 ± 0.128	1.60* ± 0.10	0.88 ± 0.15	$\begin{array}{c} 0.95 \\ \pm \ 0.07 \end{array}$	1.17 ± 0.26
	p-P53	1 ± 0.18	1.17 ± 0.16	1.11 ± 0.3	1.11 ± 0.26	1.06 ± 0.12	1.16 ± 0.15	2.53* ± 1.639	1.574* ± 0.384	1.76* ± 0.24	4.29* ± 0.47	1.71* ± 0.09	1.12 ± 0.29
	ATR	1 ± 0.14	0.98 ± 0.19	0.916 ± 0.182	1.14 ± 0.05	0.73 ± 0.13	$\begin{array}{c} 0.97 \\ \pm \ 0.08 \end{array}$	0.74 ± 0.14	0.886 ± 0.077	1.29* ± 0.75	1.1 ± 0.15	$\begin{array}{c} 0.93 \\ \pm \ 0.33 \end{array}$	0.88 ± 0.14
	p-CHK1	1 ± 0.15	1.07 ± 0.19	0.916 ± 0.195	1.26 ± 0.13	1 ± 0.27	1.21 ± 0.11	0.98 ±0.09	1.37* ± 0.18	5.21* ± 2.85	1.56* ± 0.16	1.24 ± 0.21	1.37 ± 0.32
	p-CHK2	1 ± 0.13	$\begin{array}{c} 1.05 \\ \pm 0.2 \end{array}$	0.889 ± 0.161	1.24 ± 0.18	$\begin{array}{c} 1.01 \\ \pm \ 0.32 \end{array}$	1.25 ± 0.10	$\begin{array}{c} 1.08 \\ \pm \ 0.08 \end{array}$	1.83* ± 0.32	4.21* ± 2.31	1.65* ± 0.17	1.31 ± 0.27	1.46 ± 0.37
72 h	p-H2AX	1 ± 0.13	1.36 ± 0.38	1.005 ± 0.224	0.8* ± 0.09	0.49* ± 0.06	0.42* ± 0.03	0.43* ± 0.04	0.74* ± 0.11	1.36* ± 0.73	0.94 ± 0.38	0.65 ± 0.15	0.46 ± 0.11
	MDM2	1 ± 0.12	1.16 ± 0.15	1.054 ± 0.259	1.37* ± 0.20	$\begin{array}{c} 1.01 \\ \pm 0.22 \end{array}$	1 ± 0.07	0.79 ± 0.07	0.94 ± 0.16	3.13* ± 1.98	1.03 ± 0.11	0.80 ± 0.193	0.79 ± 0.15
	P21	1 ± 0.24	1.11* ± 0.25	0.912 ± 0.199	1.42 ± 0.29	$\begin{array}{c} 0.79 \\ \pm \ 0.06 \end{array}$	$\begin{array}{c} 0.86 \\ \pm \ 0.07 \end{array}$	0.7 ± 0.15	1.03 ± 0.23	18.49* ± 7.89	1.40 ± 0.20	$\begin{array}{c} 0.88 \\ \pm \ 0.26 \end{array}$	0.94 ± 0.25
	p-P53	1 ± 0.20	0.96 ± 0.39	0.860 ± 0.132	1.67* ± 0.41	0.94 ± 0.17	$\begin{array}{c} 0.77 \\ \pm \ 0.08 \end{array}$	0.89 ± 0.18	2.24* ± 0.33	14.91* ± 9.31	3.51* ± 0.56	1.42* ± 0.17	1.51 ± 0.63

CHK-1: checkpoint kinase 1, CHK-2: checkpoint kinase 2, H2AX: H2A histone family member X, P53: protein 53, ATR: ataxia telangiectasia and Rad3-related protein, MDM2: mouse double minute 2 homolog P21: protein P21, PM_{2.5-0.3}: fine particulate matter, NEM_{2.5-0.3}: Non-extractable matter fraction, OEM2.5-0.3: fine organic extractable fraction, OEM_{0.3}: quasi-ultrafine organic extractable matter, BaP : Benzo[a]pyrene, 1-NPyr: 1-nitropyrene, 9-FluO : 9-FluOrenone.