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Method Article

An in vitro method to survey DNA methylation in peripheral blood mononuclear cells (PBMCs) treated by airborne particulate matter (PM₁₀)



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ABSTRACT

The International Agency for Research on Cancer (IARC) has defined outdoor air pollution and PM as the human carcinogen (Group 1), which mainly cause an increased risk of lung cancer. Scientists have considered epigenetic modifications as a possible mechanism to deal with adverse effects of air pollution. This study aimed to compare the effect of airborne PM_{10} (PM with aerodynamic diameter $\leq 10 \,\mu$ m) on in vitro global methylation in human peripheral blood mononuclear cells (PBMCs). PM_{10} was sampled in metropolitan Tehran, the capital of Iran. PBMCs were extracted from whole blood of healthy males and treated with PM_{10} suspension at concentrations of 50–300 μ g/mL for 4 h. Untreated cells were used as the negative control. Genomic DNA was extracted from each sample using the DNA blood mini kit according to the manufacturer's instruction. Moreover, 5-methylsytosine (%5-mC) and 5-hydroxymethylcytosine (%5-hmC) were measured by the enzyme-linked immunosorbent assay (ELISA) method. %5-mC and %5-hmC in each sample was compared with negative control and reported as difference %5-mC and %5-hmC.

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Specifications Table

Subject Area	Select one of the following subject areas: • Environmental Science
More specific subject area:	Epigenetic in environmental science
Method name:	Effect of airborne particulate matter on DNA methylation
Name and reference of	For DNA extraction see: https://www.qiagen.com/us/shop//sample-technologies/dna/
original method	qiaamp-dna-mini-kit/#resources
	For analysis of 5-mC see:
	https://www.zymoresearch.eu/5-mc-dna-elisa-kits
	For analysis of 5-hmC see:
	https://www.zymoresearch.eu/quest-5-hmc-dna-elisa-kits
Resource availability	NA

Method

The study stages were as follows: PM₁₀ sampling; preparation of PM₁₀ suspension; blood sampling and isolation of human peripheral blood mononuclear cells (PBMCs); cell culture and treatment; DNA extraction and analysis of DNA methylation and statistical analysis. All the stages are exhibited in Fig. 1.

PM₁₀ sampling

Sampling was described in detail in our previous work [1]. Briefly, PM₁₀ was sampled in Tehran, Iran (35°70′66.00″N, 51°39′38.55″E) (Fig. 2). Sampling was done as 24 h using a high-volume sampler $(1.3-1.7 \text{ m}^3/\text{min})$ (Grasebey, USA) included fiberglass filter (8 \times 10 inch, grade G 653 Whatman, USA). Firstly, filters were heated for 2 h at 180 °C and maintained in a dark and dry place. Then, they were weighed with an analytical balance $(\pm 10 \text{ mg})$ before and after sampling to calculate the sampled PM mass. Weather information was obtained from a local meteorological monitoring station (Mehrabad).

Preparation of PM₁₀ suspension

Particle extraction for biological assay was done as the dry ultrasonic method [2]. Firstly, filters were cut into the small pieces and placed in 50 ml centrifuge tubes. Tubes were placed in an ultrasonic bath (Elma-ultrasonic, Germany) for 45 min followed by smooth sweeping with a brush. PM₁₀ was collected into endotoxin-free vials, weighted and then stored at -18 °C until their use [3-5]. In the dry extraction method, presence of fiberglass fibers in PM samples is negligible and cannot cause noticeable biologic impacts on cells [5]. Morphological structure of blank filter and extracted samples were investigated by using a scanning electron microscope (SEM) (HITACHI, SU3500, Japan) (Fig. 3). Fiberglass fibers in the blank filter (Fig. 3a) were not observed in the extracted particles (Fig. 3b). Therefore, it can be concluded that dry extraction method used in this study prevented contamination of the PM₁₀ samples with fiberglass fibers. The presence of fibers in the extracted samples is serious for biological assay because these minerals have their own inherent toxicity [6]. Immediately before use,



Fig. 1. Flow diagram of the study stages.

extracted particles were autoclaved and suspended in RPMI-1640 culture medium to prepare PM₁₀ suspension at different concentrations.

Blood sampling and isolation of peripheral blood mononuclear cells (PBMCs)

This study was approved by the ethics committee of the Tehran University of Medical Sciences (IR. TUMS.SPH.REC.1395.841). Written informed consent was obtained from volunteers and their parents before initiating the study.

PBMCs were isolated from 20 mL whole blood sample from five males (16–18 years old), that enrolled in this study according to defined inclusion criteria, by density centrifugation on Ficoll-Hypaque gradient. Briefly, whole blood was diluted using 40 mL Ca²⁺/Mg²⁺ -free PBS (Biosera, France) in a laminar flux hood and gently mixed. Then, 30 mL Ficoll-Hypaque solution (Biosera, France) was added into the diluted blood and isolated PBMCs by density centrifugation (22 min, 2000 rpm, no acceleration, no brake). Layer of PBMCs was collected and washed by lysis buffer and isolation buffer and again centrifuged (400 g, 14 min, acceleration 6, brake 4). Next, PBMCs were suspended in 1 mL complete RPMI-1640 culture medium (Gibco BRL, San Diego, CA) and counted by trypan blue exclusion [7]. The viability of the cells was over 95%. Subsequently, the cells were maintained in a humidified incubator at 37 °C with 5% (v/v) CO₂ in air.





Fig. 3. SEM images of the blank filter (a) and extracted PM_{10} (b).

Cell cultures and treatment

PBMCs were cultured at a density of 500,000 cells/mL in RPMI-1640 culture medium for 24 h before treatment with PM suspension in an incubator at 37 °C with 5% (v/v) CO₂. Further, cells from each donor were separately treated during 4 h with PM₁₀ suspension at six concentrations of 50, 100, 150, 200, 250, and 300 μ g/mL. The untreated cells were used as negative control in all the experiments. All the experiments at six concentrations were conducted in triplicate to confirm repeatability of results. After 4 h, the culture medium was collected and centrifuged at 5000 rpm for 8 min to separate cells from the supernatant. Finally, the cells were stored at -20 °C until DNA extraction [1].

DNA extraction and analysis of DNA methylation

Genomic DNA was extracted from each sample using the QIAamp DNA blood mini kit (Qiagen, Hilden, Germany) according to the manufacturer's instruction. DNA concentration was measured based on absorbance at 260 nm using Nanodrop spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA).

Global %5-mC and %5-hmC in the extracted DNA samples were measured by the enzyme-linked immunosorbent assay (ELISA) method with 5-mC and Quest 5-hmCTM DNA ELISA kits (both from Zymo Research, Orange, CA, USA) using 100 ng extracted genomic DNA according to the manufacturers' protocols. All samples were assayed in duplicate. The detection limit per 100 ng single-stranded DNA was 0.5% for 5-mC and 0.02% for 5-hmC [1].

Statistical analysis

 PM_{10} concentrations in the methylation test at six levels (50, 100, 150, 200, 250 and 300 µg/mL) were considered as independent variable. The effect of PM_{10} concentration on the difference of %5-mC or %5-hmC (change of %5-mC and %5-hmC in the treated cells in respect to the negative control) as the response variables was investigated using R software version 3.4.3 [8,9]. Significant level was defined as p-value lower than 0.05. According to the Shapiro–Wilk test for the 5-mC difference (p = 0.01), the effect of PM_{10} concentration on the value of 5-mC was investigated using the non-parametric test of Scheirer Ray Hare in Fisheries Stock Analysis (FSA) package [10] and the results were reported as median \pm interquartile range (IQR). Normality and homogeneity of the 5-hmC difference were approved considering the p-value of concentration 0.11 in the Shapiro–Wilk test and 0.89 in the Bartlett test. Thus, analysis of variance (ANOVA) test was applied to compare the 5-hmC average. Moreover, Dunn test and TukeyHSD were respectively selected as post-hoc tests to assess the 5-mC and 5-hmC average between different PM_{10} concentrations.

Conflict of interest

The authors declare that they have no competing financial interests.

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