Urban Airborne Particle Exposure Impairs Human Lung and Blood *Mycobacterium tuberculosis* Immunity

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10 Online Data Supplement

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12 METHODS

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14 Human Participants

Participants were residents of Iztapalapa (the most populous municipality in Mexico City) or the neighboring Iztacalco for a minimum of one year prior to their study inclusion. Of the 35 participants, six were TST positive (>10 mm, Mantoux method, 5U Turbesol, Canada), 29 TST-negative, two interferon gamma release assay (IGRA)positive and 33 IGRA-negative. Participant numbers for each experiment are shown in the corresponding figures and their legends

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22 Primary Human Immune Cells

PBMC were obtained by gradient centrifugation from whole heparinized peripheral
 venous blood using Lymphocyte Separation Medium (Bio Whittaker, LONZA) as

described previously (<u>Supplementary Fig. 1</u>). BAC from participants were obtained by
bronchoalveolar lavage at the bronchoscopy suite of the INER using 150mL of sterile
0.9% sodium chloride solution in a segment of each right and left lung as described
previously (<u>Supplementary Fig.2</u>).

PBMC and BAC (viability ≥93%, trypan blue exclusion) were suspended in 29 complete culture medium [RPMI (BioWhittaker, Walkersville, MD, USA) + 10% pooled 30 human AB serum (Valley Biomedical, Inc.) + 2 mM L-glutamine. By cytospin analysis 31 (n= 32) BAC consisted of 91.5% AM (mean; max. 99.3%, min. 76.3%), 8.5% alveolar 32 lymphocytes (AL, mean; max. 23.7%, min. 0.7%) and scarce neutrophils. In addition, 33 the macrophage population in BAC (n=8) was also determined based on forward angle 34 and side scatter in flow cytometry (Supplementary Fig.1) and consisted of 85.64% AM 35 (mean; max. 89%, min. 81.9%) and 10.8% AL (mean; max. 16.4%, min. 5.7%). 36

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38 Collection of Outdoor PM_{2.5} in Mexico City and Preparation for *in vitro* Studies

PM_{2.5} was sampled for 24 h on Mondays, Wednesdays, and Fridays of every week 39 between May 2012 and April 2013 comprising of the three weather seasons cold-dry 40 (CD, November - February), warm-dry (WD, March - May) and rainy (R, June -41 using a high volume sampler (TE6070V-2.5, Tisch Environmental, Inc.; October) 42 Village of Cleves, OH, USA, airflow rate 1.13 m³min⁻¹) to obtain adequate amounts of 43 PM_{2.5} on modified nitrocellulose membranes (Supplementary Figs. 3, 4). PM_{2.5} was 44 mechanically recovered from the membranes, pooled by month and stored in endotoxin-45 free glass vials at 4°C, in the dark, in a desiccator after assessing the total PM_{2.5} 46 47 monthly mass. A yearly pool and pools from each weather season were created using

monthly PM_{2.5} aliquots according to the proportional contribution to the total PM_{2.5} mass
by the monthly PM samples. Immediately prior to *in vitro* exposure experiments, PM_{2.5}
was weighed on a precision balance Sartorius CPA225D (Göttingen, Germany),
autoclaved (121°C, 15 min), and suspended in complete culture medium. Following
sonication for 5 minutes (351OR-DTH; Branson, Danbury, CT, USA), PM_{2.5} suspensions
were diluted to final concentrations of 0.1, 1, 5 and 10 µg/ml.

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55 Preparation of *M.tb* for *In vitro* Infections

M.tb H37Ra (ATCC 25177, Manassas, VA) was grown in Middlebrook 7H9 broth (BD 56 Difco, Sparks, Maryland, USA) supplemented with BBL Middlebrook ADC Enrichment 57 (10% albumin dextrose catalase, BD Difco) and 0.2% glycerol (Fisher Scientific, Fair 58 Lawn, NJ, USA). After 21 days of incubation at 37°C in a shaking incubator, M.tb 59 cultures were harvested, centrifuged at 2095 x g (3000 rpm) for 30 min, room 60 temperature (RT) and suspended in Middlebrook 7H9 broth medium with 6% glycerol 61 (Fisher Scientific, Fair Lawn, NJ). Mycobacteria were then aliquoted in cryotubes and 62 stored at -70°C until use. Colony-forming unit (CFU) counts from serial dilutions on 63 7H10 agar plates (Sigma-Aldrich, Fluka. MO, USA) were performed to determine 64 concentrations of the resulting *M.tb* stock. For *in vitro* infection experiments, frozen 65 aliquots of *M.tb* H37Ra (ATCC 25177, Manassas, VA) were thawed, centrifuged for 8 66 67 minutes at 5220xg (8000 rpm), resuspended in complete medium, disaggregated by vortexing five minutes in the presence of five sterile 3-mm glass beads and centrifuged 68 at 350 x g (1000 rpm) for 2 minutes to generate single-cell suspensions of *M.tb*. 69

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71 Phagocytosis of *M.tb* by human AM

To assess phagocytosis of *M.tb*, BAC (2 x 10^4 /well) were placed in 8-well chamber slides and exposed to PM_{2.5} (1µg and 5µg/ml) for 18 h, subsequently infected with *M.tb* for 2h (37°C, 5% CO₂) at multiplicities of infection (MOI) of 0.1, 0.5, 1 and 5, washed (RPMI 3x) to remove extracellular *M.tb* and stained by Kinyoun (AlphaTec. WA. USA). Proportions of cells containing one or more intracellular *M.tb* were determined by bright field microscopy (1000x, oil immersion) on cytospin preparations in a total of 300 randomly selected AM from each study participant at each experimental condition.

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80 Cell viability, necrosis and apoptosis

Briefly, 10⁴ to 10⁶ cells were resuspended in 1 ml of protein-free buffer, mixed with 1 µl of diluted staining reagent and incubated 30 minutes on ice. After washing once with 1 ml of PBS, cells were resuspended in 200 µl of PBS and viability analyzed by flow cytometry with fluorescence detection at 660 nm (FACSfusion; BD Biosciences, San Jose, CA). Differences in intensity between live and dead cell populations were typically greater than 50-fold. Cell viability was expressed in percentages of untreated cells (controls).

For MTS assay, BAC and PBMC were seeded in 96-well plates at densities of 200,000 and 50,000 cells/well, respectively, and incubated in a total volume of 200 μ L in triplicate. Cells were incubated with PM_{2.5} suspensions (0.1, 1, 10 and 25 μ g/ml) and culture medium (viability control) for 24, 48 and 72 h (humidified, 37°C, 5% CO₂). Absorbance was quantified at 490 nm using a microplate reader (Multiskan Ascent, Thermo Electron Co., Shanghai, China). Cell viability was expressed as a percentage of

94 unexposed cells (control). For the detection of Apoptosis/Necrosis, 10^{6} BAC or PBMC 95 were placed in ultra-low attachment 24-well plates, exposed to PM_{2.5} at final 96 concentrations of 0.1, 1, 10 and 25 µg/ml and incubated for 24, 48 and 72h. Cells 97 incubated in culture medium alone or in culture medium with 2 µM of staurosporine 98 (apoptosis inducer) were included as negative and positive controls, respectively.

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100 Assessment of *in vivo* PM burden in AM

101 The ranges used for the size and shape of the particles were: size 0 - ∞ µm and 102 circularity 0.00 - 1.00, where 1 equals the perfect circle. Image adjustment was used 103 with color threshold images analyzed with a range of (HSB): Hue 50-229, saturation 60 -104 255, brightness 25 - 140. Thresholding method: default. Threshold color: network. Color 105 Space: HSB. The Overlay Mask function was used to show particle contours in the 106 superposition of images.

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108 Cytokine ELISPOT Assays

BAC and PBMC were added into Multi Screen HTS 96-flat bottom well plates coated 109 with appropriate antibodies in duplicate (200 µl/well) at optimal densities determined in 110 earlier studies and based on frequency of spot-forming cells (IFN- γ : 2x10⁴ BAC, 2x10⁵ 111 PBMC; TNF-α: 2x10² BAC, 2x10³ PBMC; IL-1β: 2x10⁴ BAC, 2x10⁵ PBMC; IL-6: 10³ 112 BAC, $2x10^3$ PBMC), incubated in the presence of 0, 0.1, 1, 10, 25 µg/ml PM_{2.5} and 113 infected simultaneously with *M.tb* (MOI 1, 5, 10) or stimulated with PPD (10µg/ml), or 6 114 kDa early secretory antigenic target (ESAT-6, 10µg/ml) at 37°C in 5% CO₂ for 24 h 115 (simultaneous exposure). Unexposed/uninfected control cells were incubated in 116

complete culture medium. Cytokine spot-forming cells (SFC) were enumerated with an ImmunoSpot Analyzer and Image Acquisition 4.5 Software (Cellular Technology Ltd, Cleveland, OH, USA) and frequencies of IL-1β, TNF- α , IL-6 and IFN- γ -producing cells calculated by averaging spot numbers from duplicate wells after background subtraction. Results are expressed as SFC/10⁶ BAC or PBMC.

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123 Western Blotting

124 To examine the activation of signaling pathways involved in the induction of cytokines, levels of P-IκB-α, P-STAT1, P-ERK1/2 and GAPDH (Cell Signaling Technologies, 125 Danver, MA) were assessed by western blotting. BAC (7.5 x 10^5) and PBMC (2 x 10^6) 126 127 were placed in 24-well plates and lysed by incubating for 30 min on ice in RIPA (radio 128 immunoprecipitation assay) lysis buffer system (Santa Cruz Biotechnology, Dallas, TX) containing 2 mM PMSF, protease inhibitor cocktail, and 1 mM sodium orthovanadate 129 and supernatants were collected following centrifugation at 10,000 g for 10 min at 4°C. 130 Protein lysates were analyzed on 10% SDS/PAGE in Tris-Glycine buffer and transferred 131 onto PVDF membrane. After blocking in 5% milk in TBST (150 mM NaCl, 50 mM Tris-132 HCl, pH 7.5, 0.05% Tween 20) for 1 hour at room temperature, membranes were 133 incubated with appropriate antibody overnight at 4°C. Following incubation, membranes 134 were washed 3-4 times in TBST and incubated with HRP-conjugated secondary 135 antibody at room temperature for 1 hour. After washing four times in TBST, protein 136 specific bands were visualized with enhanced chemiluminescence substrate Western 137 Lightning Plus-ECL (PerkinElmer, Waltham, MA). 138

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140 Statistical analysis

Statistical analyses were performed using GraphPad Prism, version 5.0 (GraphPad
Software Inc., San Diego, CA, USA). Types of statistical analysis used for experiments
and conditions are indicated in the corresponding figure legends in the Results section.

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145 SUPPLEMENTARY FIGURES

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Supplementary Figure 1. Cytotoxic effect of PM_{2.5}. PBMC or BAC were exposed to 147 PM and incubated for 24-72h to evaluate PM2.5 exposure effects on membrane 148 permeability and induction of apoptosis and necrosis. Permeability of PBMC and BAC 149 membranes was evaluated by flow cytometry. Individual cells from an exemplary BAC 150 sample were selected in SS-A and SS-H (A), lymphocyte and macrophage 151 subpopulations identified (B) and single histogram plots generated to identify live 152 lymphocytes and macrophages using LIVE/DEAD Fixable Far Red Dead Cell Stain (C, 153 <u>D</u>). 154

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Supplementary Figure 2. PM_{2.5} does not affect *M.tb* phagocytosis by AM. AM were pre-exposed to PM_{2.5} cold-dry, warm-dry, rainy weather seasons at 0 and 1 μ g/ml (<u>A</u>) and 5 μ g/ml (<u>B</u>) for 24 h, infected with *M.tb* MOI 0.1, 0.5, 1 and 5 for 2 h and proportions of AM with intracellular *M.tb* determined by bright field microscopy (1000x, oil immersion) on Kinyoun-stained cytospins assessing a total of 300 AM in each experimental condition.

Supplementary Figure 3. Effect of PM_{2.5} on *M.tb*-induced Cytokine Production in 163 **BAC and PBMC.** Optimal densities of BAC and PBMC (n=9) were exposed to PM_{2.5} (0, 164 0.1, 1 and 10 µg/ml) and simultaneously infected with *M.tb* or left uninfected for 24h. 165 Frequencies of IL-1 β , TNF- α , IL-6 and IFN- γ spot-forming cells were evaluated by 166 ELISpot assay. Images of representative IL-1 β , TNF- α , IL-6 and IFN- γ -producing BAC 167 showing cytokine spot formation and identification in *in vitro* PM-exposed and *M.tb*-168 infected cells for image analysis and automated counting (A). Frequencies of 169 constitutively (unstimulated) IFN- γ , IL-1 β , IL-6, and TNF- α -producing BAC (open circles) 170 and PBMC (black circles) (B). Frequencies of IL-1 β (C), TNF- α (D) IL-6 (E) and IFN- γ 171 production (F) in response to in vitro PM exposure (left column panels) and M.tb 172 173 infection (right column panels) by BAC (open circles) and PBMC (black circles). Each 174 data point represents an individual participant and horizontal lines indicate medians. Asterisks indicate significant differences ($p \le 0.05$, ANOVA Friedman test followed 175 176 Dunn's multiple comparisons test using the GraphPad prism software for Mac OS X version 6.0) relative to PM_{2.5}-unexposed control cells. 177

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Supplementary Figure 4. Activation of signaling pathways. BAC and PBMC from
three participants (1-3) were infected with *M.tb* at MOI1 and 5 or stimulated with PPD or
left unstimulated (constitutive) for 24h. Protein lysates were analyzed with SDS/PAGE
followed by western blotting for Phospho-IκBα (Ser32/36), Phospho-STAT1 (Tyr701)
and Phospho-Erk1/2 (Thr202/Tyr204), and GAPDH.

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