Supplemental Material

Short-term Exposure to Ambient Air Pollution and Biomarkers of Systemic Inflammation: The Framingham Heart Study

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Materials and Methods

Study Sample

We included participants from the Framingham Offspring cohort examination cycle 7 (1998-2001), cycle 8 (2005-2008), Third Generation cohort examination cycle 1 (2002-2005), and cycle 2 (2008-2011). The study design and selection criteria of the two cohorts have been described elsewhere.^{1, 2} First, we restricted our analyses to 4,110 participants who lived within 50 km from the Harvard Supersite air pollution monitor in Boston, Massachusetts and who were not current smokers at the time of their examination visit. We excluded current smokers (1,139 observations) because levels of inflammatory biomarkers were likely already elevated among current smokers and may affect our ability to assess the relatively small variation in levels of inflammatory biomarkers that could be attributable to ambient air pollution,³ similar to our previous work.⁴ Then we assigned "missing status" to biomarker levels that were under the minimum detection limit (27 observations for CRP, 10 observations for interleukin-6, and 1 observation for TNFR2) and 1 TNFR2 measurement that had an extremely high value (948,114 pg/ml). Last, we excluded 65 participants who had no measurement of any biomarkers and 49 participants who had missing information on covariates including body mass index, alcohol intake, or pack years of smoking, leaving 3,996 participants in the final analytic dataset. At each examination visit, physical examinations were performed following standardized protocols, and data on demographics, medication history, smoking history, and alcohol intake were collected using standardized questionnaires. All participants provided written informed consent at each examination, and the Institutional Review Boards at Beth Israel Deaconess Medical Center and Boston University Medical Center approved the study.

Air Pollution Assessment

Previous studies have shown that the blood levels of the inflammatory biomarkers could increase within a few days after inflammation.^{5, 6} thus we calculated 1-, 2-, 3-, 5-, and 7-day moving averages of the air pollutants prior to the date of examination visit, based on hourly measures of PM_{2.5}, black carbon (BC), and sulfate (SO₄²⁻) from the Boston Harvard Supersite air pollution monitoring station, and measures of nitrogen oxides (NO_x) and ozone (O_3) from local state monitors within the Greater Boston area.⁷ The central monitor is located on the rooftop of the Francis A. Countway Library of Medicine (5 stories above ground level) and 50 m from the nearest street. PM_{2.5} was measured using a tapered element oscillating microbalance (Model 1400A, Rupprecht & Patashnick Co., Inc.); BC was measured using an aethalometer (Model AE16, Magee Scientific Corp.). We calculated daily average SO₄²⁻ from elemental sulfur that was measured by X-Ray Fluorescence analysis of the PM_{2.5} filter samples, and used an SO₄²⁻ analyzer (Model 5020, Thermo Electron Corp.) on days when SO₄²⁻ X-Ray Fluorescence data were not available. The detailed measurement methods have been described previously.⁸ We also obtained hourly temperature and relative humidity data from the Boston Logan International Airport Weather Station, located 12 km from the central site.⁷

Biomarker Assessment

Blood samples were collected after an overnight fast and stored at -80°C until assayed. Plasma samples were analyzed to measure TNFα, TNFR2, and fibrinogen, and serum samples were used for CRP and IL-6. CRP was measured twice in each cohort (Offspring examination 7 and Third Generation examination 1: immunonephelometry, Dade Behring BN 100 nephelometer; Offspring examination 8 and Third Generation examination 2: immuno-turbidometry, R&D Systems). Fibrinogen was measured using the Clauss method (Diagnostica Stago Reagents) in Offspring examination 7 and Third Generation examination 1. TNFR2 was measured twice in the Offspring cohort and once in the Third generation cohort, Interleukin-6 was measured in Offspring examination 7, all by commercially available enzyme-linked immunosorbent assay (ELISA) kits (R&D Systems). Details regarding biomarker measurements have been reported elsewhere ⁹⁻¹¹ and can be found at the Framingham Heart Study website: https://www.framinghamheartstudy.org/

researchers/description-data/noninvasive-biomarker.php.

Statistical Methods

We fit multivariable linear regression models for fibrinogen, interleukin-6, and TNFα, and multivariable linear mixed effects models with participant-specific random intercepts for CRP and TNFR2 for each of the 1-, 2-, 3-, 5-, and 7-day moving averages before the date of examination visit. Levels of the biomarkers were log_e transformed to approximate a normal distribution. In each model, we adjusted for demographic and socio-economic position variables including centered age at the time of examination visit, (centered age)², sex, educational attainment (high school or less, some college, and college graduate), census tract-level median household income (continuous); lifestyle variables including body mass index, alcohol intake (drinks/week; continuous, standardized to 0.5 oz [15 ml] alcohol/drink),¹² pack years of smoking (continuous), and smoking status (never or former smoker); and an examination identifier (categorical). All census data were from U.S. Census 2000. We adjusted for seasonality and time by adding sine and cosine terms of the examination date and a linear term of the examination date, and adjusted for meteorology by adding corresponding moving averages of temperature and relative humidity.

For sensitivity analyses, we examined whether the associations differed if we: 1) excluded observations that had a daily average PM_{2.5} concentration>35 µg/m³ (U.S. Environmental Protection Agency 24-hour PM_{2.5} standard) in any one of the 7 days prior to the examination date; 2) included current smokers in the analyses; 3) restricted study participants to those who lived within 40 km from the central monitoring site; and 4) additionally adjusted for usual occupation (laborer; sales/homemaker/clerical; professional/executive/supervisory/technical; and unspecified),¹³ census tract median value of owner occupied housing units (continuous), census tract population density (people/km², continuous), physical activity index (in tertiles),¹⁴ cardiovascular disease status, antihypertensives use, and statins use. We also explored whether associations differed by median age of the overall available study sample (53 years old), sex, education level (high school or less *versus* higher than high school), diabetes status, cardiovascular disease status, anti-hypertensives use, statins use, and season (warm [April to September] *versus* cold [October to March]) by adding an interaction term to

the models. Because we pooled participants from both cohorts together in the primary analyses, we examined the associations separately in each cohort. Further, because not all participants have data on all moving averages, we conducted a sensitivity analysis where we restricted analyses to the same participants across all moving averages for each pollutant. For all analyses, we created missing indicators for participants with missing data on educational attainment (35 observations) or physical activity (96 observations).

Parameter estimates from the analyses were scaled by a factor close to the interquartile range for the 1-day moving average of each pollutant: $5 \mu g/m^3$ for PM_{2.5}, 0.5 $\mu g/m^3$ for BC, 2 $\mu g/m^3$ for SO₄²⁻, 20 ppb for NO_x, and 10 ppb O₃. We reported estimated percent differences with 95% confidence intervals (CIs). As to interpreting the results from our primary analyses, we focused on describing the observed association patterns between pollutants and the biomarkers across moving averages and pollutants. For sensitivity analyses in which effect modification was examined, we used the two-tailed *p*-value from the Wald test of the interaction term to judge whether the associations differed between subgroups (considered statistically significant if p<0.05). We reported and highlighted association patterns that we considered consistent across moving averages or pollutants. Analyses were performed using PROC GENMOD and PROC MIXED in SAS 9.4 (SAS Institute, Inc., Cary, NC). Figures were plotted using Stata 13 (StataCorp LP, College Station, TX).

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