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Exposure to ambient air pollution during pregnancy and inflammatory biomarkers in maternal and umbilical cord blood: The Healthy Start study

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Abstract

Background: Air pollution exposure during pregnancy has been associated with adverse pregnancy and birth outcomes. Inflammation has been proposed as a potential link. We estimated associations between air pollution exposure during pregnancy and inflammatory biomarkers in

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

maternal and cord blood. We evaluated whether maternal inflammation was associated with infant outcomes.

Methods: Among 515 mother-infant dyads in the Healthy Start study (2009-2014), trimester-long, 7- and 30-day average concentrations of particulate matter ≤ 2.5 micrometers ($PM_{2.5}$) and ozone (O_3) during pregnancy were estimated, using inverse-distance-weighted interpolation. Inflammatory biomarkers were measured in maternal blood in mid-pregnancy (C-reactive protein [CRP], Interleukin [IL]-6, and tumor necrosis factor- α [TNF α]) and in cord blood at delivery (CRP, IL-6, IL-8, IL-10, monocyte chemoattractant protein-1 [MCP-1], and TNF α). We used linear regression to estimate associations between pollutants and inflammatory biomarkers and maternal inflammatory biomarkers and infant weight and body composition.

Results: There were positive associations between $PM_{2.5}$ during certain exposure periods and maternal IL-6 and TNF α . There were negative associations between recent O_3 and maternal CRP, IL-6, and TNF α and positive associations between trimester-long O_3 exposure and maternal inflammatory biomarkers, though some 95% confidence intervals included the null. Patterns were inconsistent for associations between $PM_{2.5}$ and O_3 and cord blood inflammatory biomarkers. No consistent associations between maternal inflammatory biomarkers and infant outcomes were identified.

Conclusions: Air pollution exposure during pregnancy may impact maternal inflammation. Further investigations should examine the health consequences for women and infants of elevated inflammatory biomarkers associated with air pollution exposure during pregnancy.

Keywords

Air pollution; Inflammation; Pregnancy; Infant

1. Introduction

There is a growing body of evidence linking prenatal ambient air pollution exposure with fetal, infant, and child health outcomes.¹⁻³ Several previous epidemiologic studies have reported associations between pregnancy exposure to air pollutants, such as particulate matter ≤ 2.5 micrometers ($PM_{2.5}$) and ozone (O_3), and adverse birth outcomes, including low birth weight, small for gestational age (SGA), and preterm birth.⁴⁻⁹ Evidence from a limited number of animal^{10, 11} and human studies^{12, 13} shows that prenatal exposure to components of air pollution may also adversely affect offspring cardio-metabolic health during childhood. These findings suggest that the prenatal time may represent a critical period of fetal susceptibility to environmental stressors, such as air pollution.^{14, 15}

Prenatal air pollution exposure has been linked to inflammatory airway-related conditions in infancy and childhood, including decreased lung function, wheezing, and asthma.¹⁶⁻¹⁹ Further, differences in the levels of inflammatory biomarkers in cord blood at delivery have been identified between SGA and appropriate for gestational age infants,²⁰⁻²² suggesting a role for inflammation in this outcome. Since low birth weight and subsequent rapid growth during infancy are associated with greater risk of cardio-metabolic disease during childhood,²³⁻²⁶ and inflammatory processes are key components of cardio-metabolic disease

pathophysiology,^{27–29} systemic inflammation may be a plausible biological pathway through which air pollution may lead to cardio-metabolic disease later in life.

Several previous studies from the Netherlands,³⁰ Germany,³¹ France,³² the Czech Republic,^{33–35} Canada,³⁶ and the United States (US)³⁷ have reported that exposure to air pollution during pregnancy was associated with levels of immune biomarkers in the mother and the neonate. However, the Dutch study was the only to investigate the relationship between the air pollutants and an inflammatory marker, C-reactive protein (CRP), in both maternal and cord blood.³⁰ Greater PM₁₀ exposure in the prior 1-2 weeks was associated with elevations in maternal CRP levels during trimester 1, and higher full pregnancy maternal PM₁₀ and nitrogen dioxide (NO₂) exposure were associated with higher levels of CRP in cord blood at delivery.³⁰ Given that the inflammatory response is regulated by several different cytokines with diverse and context-dependent roles, further characterization of the relationship between air pollution and pro- and anti- inflammatory biomarkers is warranted.

We investigated the associations between ambient air pollution exposure (PM_{2.5} and O₃) during pregnancy and maternal and cord blood inflammatory biomarkers. In both maternal and cord blood, we assessed these relationships in a core group of three pro-inflammatory biomarkers, CRP, Interleukin (IL)-6, and tumor necrosis factor- α (TNF α) that have been associated with adverse pregnancy and birth outcomes.^{38–41} We investigated these relationships with an expanded panel in cord blood, with the addition of IL-8, IL-10, and monocyte chemoattractant protein 1 (MCP-1). We hypothesized that greater exposure to ambient air pollution would be associated with higher concentrations of pro-inflammatory biomarkers in maternal mid-pregnancy serum, and these relationships would differ according to timing of exposure to air pollution during pregnancy. We also hypothesized that greater exposure to air pollution during pregnancy would be associated with alterations in concentrations of inflammatory biomarkers in cord blood at delivery. Additionally, we examined whether maternal inflammatory biomarkers during pregnancy were associated with offspring birth weight and body composition.

2. Methods

2.1 Study sample

We included participants from the Healthy Start study, which is an ongoing longitudinal cohort study of 1,410 ethnically diverse mother-infant dyads. Briefly, pregnant women who had not yet reached 24 completed weeks of gestation were recruited at obstetrics clinics at the University of Colorado Hospital between 2009 and 2014. Participants attended follow-up research visits through pregnancy, delivery, and now into childhood for the offspring. Eligibility criteria included singleton pregnancy, age ≥ 16 years, and no history of chronic disease (diabetes, cancer, asthma treated with steroids, or medication-dependent psychiatric illness) or a birth <25 weeks gestation. The participation rate for the Healthy Start study was approximately 50%. The median gestational age at time of enrollment was 17 weeks (range, 11–20). Study participants provided written informed consent. The Colorado Multiple Institutional Review Board approved all study protocols.

Of the 1,410 participants enrolled in the Healthy Start study, 11 withdrew from the study prior to delivery and 17 experienced fetal demise. Of the 1,382 eligible participants, we measured inflammatory biomarkers in 536 dyads with sufficient volume of both maternal and cord blood stored serum samples, and we excluded participants who resided outside of the Denver, Colorado metropolitan area or with no O₃ and/or PM_{2.5} monitors within 50 km of the residence^{8, 42} (n=21), ultimately resulting in a sample size of 515 for the present analysis (Supplementary Figures 1 and 2). For the analysis of the offspring outcomes, which consisted of birth weight, gestational age, and body composition (% fat mass [adiposity], fat mass, and fat free mass) at birth and at 5 months of age, some participants had missing data. There were 497 and 334 mother infant-pairs included in the analysis of body composition at birth and at five months, respectively.

2.2 Assessment of exposure

We estimated average ambient air pollution exposure levels (PM_{2.5} and O₃) for trimester-long and 7- and 30- day exposure windows during pregnancy using measurements obtained from stationary monitors in the United States Environmental Protection Agency Air Quality System. Participants reported their place of residence via questionnaire completed at the time of enrollment in the Healthy Start study. Details have been described previously.⁴²

Briefly, there were 10 stationary monitors that measured average 24-hour PM_{2.5} and 19 stationary monitors that measured hourly O₃, which were located within 50 km of at least one study participant. The 50 km distance was selected based on previous literature.^{43, 44} Most of the monitors for PM_{2.5} recorded measurements every three days, with some measuring daily or every six days. All of the monitors for O₃ recorded measurements hourly each day. We calculated trimester-specific average and full-pregnancy average PM_{2.5} levels by averaging the daily values from each monitor within 50 km over the relevant exposure period for each participant. Additionally, we calculated average PM_{2.5} levels for different time windows (30 days and 7 days) preceding the maternal blood sample collection and cord blood collection at delivery, to capture both recent and longer term exposure and to enhance comparability with prior literature that has included similar exposure windows.^{30, 31, 37} For O₃, the maximum 8-hour average for each 24-hour period was determined, and daily 8-hour maximum values were averaged over the same time-periods that were estimated for PM_{2.5}. For both air pollutants, we used inverse-distance-weighting ($1/(\text{distance}^2)$) to calculate the average exposure at each participant's residence using the period-specific averages from all monitors within 50 km.⁴⁵ We included averages from each monitor if the number of non-missing values was at least 75% of the expected values.^{44, 45}

2.3 Assessment of outcomes

Fasting maternal blood samples were collected at a median gestational age of 27 weeks, and immediately separated and stored at -80°C. Inflammatory biomarkers in maternal serum were analyzed by the University of Colorado Clinical and Translational Sciences Institute Core Laboratories. Maternal CRP was measured using immunoturbidimetric methodology (Beckman Coulter, Inc, Indianapolis, IN). Maternal IL-6 was measured using Luminex MAP technology (R&D Systems, Inc, Minneapolis, MN, USA). Maternal TNF α was measured

using enzyme-linked immunosorbent assay (ELISA) (R&D Systems, Inc, Minneapolis, MN, USA).

Infant inflammatory biomarkers were measured in cord blood drawn at delivery. Samples were stored on ice for up to 20 minutes and then centrifuged for processing. They were next stored at 4 °C for up to 24 hours, before being transferred for long-term storage in a freezer kept at -80 °C. Cord blood CRP levels were determined by an ELISA (Alpco, Salem, New Hampshire, USA). Cord blood inflammatory biomarkers, IL-10, IL-1/ β , IL-4, IL-6, IL-8, MCP-1, and TNF α were processed by the University of Colorado Cancer Center Flow Cytometry Shared Resource. Cord blood cytokine levels were determined by multiplex panel immunoassay, according to manufacturer's instructions (EMD Millipore Corporation, Billerica, MA, USA). We restricted analyses to those cytokines present in at least 70% of the study sample (IL-6, IL-8, IL-10, MCP-1, TNF α). The limit of detection of the assays varied by panel. There were no values below the minimum limit of detection for the five cord blood cytokines included. For cord blood cytokine values that were above the maximum detectable concentration (IL-6, n=1; MCP-1, n=3), we used the value of the maximum detectable concentration for that panel.

Additional infant outcomes included: birth weight, gestational age at birth, and body composition (fat mass, fat free mass, and fat mass as percent of body mass, hereafter referred to as adiposity) at birth and at approximately five months of age (mean \pm standard deviation: 5.1 \pm 1.2 months). Birth weight and gestational age at birth were obtained from medical records. Body composition at birth and at approximately five months was measured using whole body air displacement plethysmography (PEA POD, COSMED, Rome, Italy)⁴⁶ by trained research staff. Body composition was measured up to three times for each infant, and the two closest measurements were averaged.

2.4 Other variables

Participants reported their age, race/ethnicity, education completed, and the number of previous pregnancies at the time of their study enrollment. We calculated maternal pre-pregnancy body mass index (BMI) using height measured at the first study visit and weight obtained from the medical record (87%) or self-report at study enrollment (12%) if unavailable in the medical record. Participants self-reported their smoking status at two study visits during pregnancy and at delivery, and we created a binary variable to indicate any smoking during pregnancy. Census tract-level median income was obtained from the 2012-2016 American Community Survey and linked to the participant's geocoded enrollment address. Infant sex, season of birth, and mode of delivery were obtained from the medical record.

2.5 Statistical analysis

Because ambient air pollution exposure was estimated using inverse-distance-weighted interpolation of data obtained from stationary monitors, and not directly measured, there is the possibility of measurement error due to spatial heterogeneity. We calculated period-specific intra-class correlation coefficients (ICC) for the three monitors closest to the participant's address that contributed to the average exposure estimate. The ICC was

interpreted as the degree of spatial homogeneity in the study area. We calculated Spearman correlations between period-specific air pollution exposures and between maternal and cord blood inflammatory biomarkers.

We used separate multivariable linear regression models to estimate the associations between the period-specific air pollution exposures and each of the maternal and cord blood inflammatory marker outcomes. For maternal inflammatory marker outcomes, which were measured at a median of 27 weeks of gestation, we only examined the 30- and 7-day periods preceding the sample collection and the first and second trimester exposures. For cord blood inflammatory biomarkers, which were measured at delivery, we considered all period-specific average exposures. We natural log-transformed all maternal and cord blood inflammatory biomarkers to minimize the effect of potential outliers. Results are presented as the percent change and 95% confidence intervals (CI) for an IQR increase in the pollutant. We also used multiple linear regression to investigate the association between maternal inflammatory biomarkers and infant birth weight and body composition outcomes.

We included potential confounders identified via a directed acyclic graph that represented hypothesized causal relationships and associations reported in previous literature (Supplementary Figure 3). All models were adjusted for the corresponding co-pollutant (PM_{2.5} or O₃) during the specified pregnancy period, maternal age at delivery (years), pre-pregnancy BMI (kg/m²), race/ethnicity (Hispanic, non-Hispanic white, non-Hispanic black, all others), smoking during pregnancy (any, none), median income in the Census tract (dollars), maternal education (less than 12th grade, high school degree or GED, some college or associate's degree, four years of college, graduate degree), any previous pregnancies (yes, no), and infant sex. Models with maternal inflammatory biomarkers were additionally adjusted for gestational age at maternal blood sample collection and season of conception (winter, spring, summer, fall), and models with cord blood inflammatory biomarkers were additionally adjusted for mode of delivery (vaginal, Caesarean) and season of delivery (winter, spring, summer, fall). Models for associations between maternal inflammatory biomarkers and birth and body composition outcomes were also adjusted for gestational age at maternal blood sample collection, and the models for body composition at five months were additionally adjusted for infant age at the time of the body composition measurement. We performed a sensitivity analysis excluding infants who were born preterm (<37 weeks gestation) (n=21).⁴⁷ We hypothesized that gestational age was a causal intermediate between maternal inflammatory biomarkers and birth weight and body composition, therefore, the models for the primary analyses did not include gestational age. However, we performed a sensitivity analysis adjusting for gestational age in models for these outcomes. We evaluated model assumptions by visual inspection of scatter plots between exposures and outcomes to detect departures from linearity and the presence of outliers and residual plots to identify violations of homoscedasticity.

Because there is some evidence that the maternal immune response may differ according to fetal sex, we investigated whether infant sex was a potential effect modifier by including product interaction terms between each air pollutant and infant sex, in adjusted models. We considered the corresponding co-pollutant as a potential confounder, due to the inverse relationship between PM_{2.5} and O₃. We conducted a sensitivity analysis with models that

only included a single pollutant. We performed two additional sensitivity analyses: the first in which we excluded participants who smoked during pregnancy, and the second, for cord blood only, in which we excluded values above the maximum detectable concentration for that panel.

We conducted all analyses in SAS (Version 9.4, The SAS Institute, Cary, NC, USA) and in R version 4.0.2 (R Foundation for Statistical Computing, Vienna, Austria).

3. Results

The characteristics of the participants in this study sample were comparable to those not included in this study sample and to those of the entire Healthy Start cohort (Supplementary Table 1). The average maternal age at delivery was 28 ± 6 years, and the average pre-pregnancy maternal BMI was 26 ± 7 kg/m². The study sample was racially and ethnically diverse (Table 1).

The median percent of stationary monitors within 50 km of the participants' residences with less than 25% missing observations was 75% for PM_{2.5} and 78.6% for O₃ (Supplementary Table 2). The median full pregnancy average PM_{2.5} and average 8-hour O₃ maximum levels were 7.51 µg/m³ (range 5.41-9.39 µg/m³) and 44.0 parts per billion (ppb) (range 30.6-54.0 ppb), respectively (Table 1). Air pollution exposures varied seasonally according to the timing of the pregnancy and were generally weakly or inversely correlated between trimesters, and more highly correlated between shorter periods (Supplementary Figure 4).

The ICCs for period-specific average levels at the three nearest monitors were moderate to high, ranging from 0.56-0.80 for PM_{2.5} and 0.73-0.86 for O₃, suggesting reasonable levels of spatial homogeneity in the study area.

There was a moderately high Spearman correlation (r_s) between maternal CRP (CRP_m) and IL-6_m ($r_s=0.50$), with weak correlations observed between CRP_m and TNFα_m ($r_s=0.02$) and IL-6_m and TNFα_m ($r_s=0.10$) (Table 2).

Median concentrations of cord blood biomarkers varied by several orders of magnitude, with the lowest median concentration observed for cord blood IL-6 (IL-6_{cb}) (9.85 pg/mL, IQR 4.80-24.46 pg/mL) and the highest median concentration observed for CRP_{cb} (0.09 mg/L, IQR 0.04-0.18 mg/L). The pairwise Spearman correlations between the cord blood inflammatory biomarkers ranged from 0.02 to 0.64, with the highest correlations observed between IL-6_{cb} and IL-8_{cb} ($r_s=0.64$) and between MCP-1_{cb} and TNFα_{cb} ($r_s=0.64$) (Table 2).

For the three inflammatory biomarkers that were measured in both cord blood and maternal blood, the observed median concentrations of CRP were lower in cord blood compared to maternal blood (0.09 mg/L cord versus 4.48 mg/L maternal), whereas the median levels of both IL-6 and TNFα were higher in cord blood compared to maternal blood (IL-6: 9.85 pg/mL cord versus 1.36 pg/mL maternal; TNFα: 32.33 pg/mL cord versus 1.11 pg/mL maternal). Cord blood and maternal inflammatory biomarkers were largely uncorrelated with each other (Spearman correlation range: -0.02, 0.15), except CRP which was moderately correlated between maternal and cord blood samples ($r_s=0.26$) (Table 2).

In adjusted models including the other co-pollutant, we observed higher TNF α _m with higher trimester-long concentrations of PM_{2.5} and O₃. For example, an IQR increase in trimester 1 O₃ was associated with 30.39% (95% CI; 14.46, 48.54) higher TNF α _m (Figure 1 and Supplementary Table 3). Higher trimester 1 and 2 PM_{2.5} was associated with higher TNF α _m, as well as prior 30-day PM_{2.5} though the 95% CI included the null for this time period. PM_{2.5} was generally positively associated with IL-6_m, notably for the prior 30- days to sample collection and trimester 2. For example, in the prior 30- days to sample collection, an IQR increase in PM_{2.5} was associated with 6.54% higher IL-6_m (95% CI; 0.86, 12.53). PM_{2.5} in the 7-days prior had a non-significant negative association with CRP_m. Higher trimester 1 and 2 O₃ exposure was associated with higher IL-6_m, though the 95% CI for trimester 2 included the null. Higher prior 30- and 7- day O₃ exposure was non-significantly associated with lower TNF α _m. For example, in the prior 7-days to sample collection, an IQR increase in O₃ was associated with 11.43% (95% CI; -21.83, 0.34) lower TNF α _m. There were non-significant positive associations between trimester 1 and 2 O₃ exposure and CRP_m. Results from unadjusted models were generally similar in direction and magnitude to adjusted results, except that associations between trimester 1 and 2 PM_{2.5} and TNF α _m were attenuated and no longer significant (Supplementary Table 4). Results from single pollutant models were comparable (Supplementary Table 6).

PM_{2.5} exposure was not consistently significantly associated with any cord blood inflammatory biomarker, however, there were several notable trends. Specifically, there were non-significant positive relationships between trimester 1, 2, and full-pregnancy PM_{2.5} and CRP_{cb}. Additionally, PM_{2.5} exposure in trimester 1 and trimester 3 was associated with higher and lower IL-8_{cb} levels, respectively. Finally, trimester 2, 3, and full-pregnancy PM_{2.5} was generally associated with lower MCP-1_{cb} and TNF α _{cb}, but higher levels of these inflammatory biomarkers when considering prior 30-day PM_{2.5} exposure (Table 3 and Supplementary Figure 5).

Similarly, O₃ was not consistently significantly associated with any cord blood inflammatory marker, but there were some trends. There were non-significant positive associations between O₃ during all of the examined periods of pregnancy and CRP_{cb}. Trimester 2 O₃ was positively associated with IL-6_{cb}, IL-8_{cb}, and MCP-1_{cb}, and trimester 1, and prior 7- and 30-day O₃ was negatively associated with IL-6_{cb} and IL-8_{cb} (Table 3 and Supplementary Figure 6).

We did not find evidence for exposure-by-sex interaction in cord blood at alpha of 0.05. Results from unadjusted models are presented in Supplementary Table 5. In single pollutant models, results were similar (Supplementary Table 7).

In the sensitivity analysis excluding participants who smoked during pregnancy, the results for the associations between the air pollutants and cord blood and maternal inflammatory biomarkers did not differ appreciably (Supplementary Tables 8 and 9). Similarly, the associations between the air pollutants and cord blood cytokines, excluding values that were above the maximum detectable concentration (n=4 total), were comparable to the results of models in which we substituted the maximum detected value (Supplementary Table 10).

CRP_m was positively associated with infant gestational age at birth (β ; 1.30, 95% CI; 0.40-2.21). There were no associations between CRP_m, IL-6_m, and TNF α _m and birthweight, adiposity at birth, and adiposity at five-month follow-up (Table 4). We did not observe associations between the maternal inflammatory biomarkers and fat mass and fat free mass at birth and at five-month follow-up (Supplementary Table 11). Results remained null, after excluding preterm births and adjusting for gestational age (Supplementary Tables 12, 13, 14, and 15).

4. Discussion

In this racially and ethnically diverse sample of mother-infant dyads, we found associations between exposure to PM_{2.5} and O₃ during several time windows and some maternal inflammatory biomarkers during mid-pregnancy. We also observed associations between air pollution during some time windows in pregnancy and inflammatory biomarkers in cord blood at delivery, though patterns were inconsistent.

Dysregulation of the maternal immune response has been identified in pregnancy-related diseases, including pre-eclampsia, gestational diabetes (GDM),^{38, 39} and adverse birth outcomes, including fetal growth restriction and preterm birth.^{40, 41} Given the impact of maternal inflammation on both maternal and fetal health, a greater understanding of potentially preventable sources of alterations in the immune profile, such as air pollution, is pertinent. A prior study investigated the relationship between air pollution and the maternal lymphocyte immune profile.³³ In a cross-sectional study in the Czech Republic, Hertz-Picciotto and colleagues found mothers who gave birth in a city with high levels of air pollution had lower percentages of T cells and a lower CD4+/CD8+ ratio compared to those who lived in a less polluted city.³³ However, this study did not examine associations between ambient air pollution with cytokines. Cytokines are pleiotropic factors responsible for mediating much of T-lymphocyte function,⁴⁸ as well as several other functions, and they may be mediators in the relationship between pregnancy environmental exposures and adverse offspring outcomes.

Dysregulation of IL-6,⁴⁹ TNF α ,⁵⁰ and CRP^{40, 51} has been linked to pregnancy complications associated with the cardio-metabolic health of the mother and the neonate, including preeclampsia and preterm birth. During healthy pregnancy, upregulation of cytokines may promote the typical increase in maternal insulin resistance necessary for normal fetal growth; however, an excessive inflammatory response may induce abnormal insulin resistance and in some cases, the development of GDM.⁵²⁻⁵⁴ Higher CRP during pregnancy is found in women with obesity, which is associated with decreased insulin sensitivity during pregnancy.⁵⁵ Higher levels of IL-6 and in some studies TNF α have also been associated with insulin resistance⁵⁶ and GDM, respectively.⁴⁹

We found positive associations between PM_{2.5} and IL-6_m levels during mid-pregnancy. We also observed positive associations for trimester-long O₃ exposure and IL-6_m. While there is a lack of literature exploring this relationship in pregnant women, this finding is consistent with epidemiologic evidence suggesting an association between acute exposure to particulate matter and IL-6 in adult cohorts.⁵⁷⁻⁵⁹ However, other studies have reported null

results.^{60, 61} IL-6 is a pro-inflammatory cytokine that has roles in both acute and chronic inflammatory responses, and in rodents, elevated levels of IL-6 and lower levels of its inhibitor (soluble gp130) have been associated with fetal loss, preterm birth, and infertility, and they have also been observed in pregnant women with preeclampsia and gestational diabetes.⁴⁹ In early pregnancy, IL-6 is involved in placental development, in mid-pregnancy IL-6 levels are highly regulated and its production and activity are suppressed, and finally during labor, IL-6 levels raise to facilitate parturition.⁴⁹ Disruptions in the tightly regulated and temporally-controlled levels of maternal IL-6 during pregnancy, may represent a pathophysiological rationale for the adverse pregnancy and birth outcomes observed in association with environmental exposures, such as air pollution.

We also observed positive associations between PM_{2.5} and O₃ exposure and maternal mid-pregnancy TNF α . Elevated TNF α levels have been observed following acute exposure to particulate matter⁵⁷ and diesel exhaust⁶² in healthy adults, however, we only observed associations with trimester-average, and not more short-term, exposure. The placenta and adipose cells are two of the main sources of TNF α production during pregnancy, and higher TNF α levels are observed during the pregravid and late pregnancy periods, with a decline in serum levels in early pregnancy.^{56, 63} TNF α is a known inhibitor of the tyrosine phosphorylation cascade of the insulin receptor, and as a result, it is associated with the normal insulin resistance that is characteristic of pregnancy.⁶³ However, further elevated levels of TNF α have been identified in GDM, and consequently, it has been hypothesized to be a mediator of the growth-related birth outcomes associated with GDM and environmental exposures, such as tobacco smoke.^{63, 64} The present study suggests air pollution exposure during pregnancy may also disturb regulation of TNF α and given the between link between TNF α and metabolic homeostasis, this may have health implications for both the mother and the offspring.

We noted non-significant positive associations for trimester-long O₃ exposure and maternal CRP levels. Previous findings from 5,067 mothers in a population-based prospective cohort in the Netherlands identified higher PM₁₀ exposure during the prior 1-2 weeks to be associated with elevated concentrations of maternal CRP (>8 ng/mL) in trimester 1.³⁰ Further, a study of 1,696 mothers in Pennsylvania found PM_{2.5} in the 28 days prior to the blood sample to be associated with higher odds of elevated maternal CRP (> 8 ng/mL) in early pregnancy for non-smokers, along with positive, but non-significant, associations with PM₁₀ and O₃ exposure.³⁷ One potential explanation for this discrepancy is the variation in air pollution exposure levels by geographic location, as average particulate matter concentrations in Colorado⁶⁵ are much lower than those found in major cities in Europe⁶⁶ and the northeastern US.⁶⁷

Previous literature examining associations between maternal air pollution exposure and changes in the cord blood immune profile has noted alterations in absolute numbers and percentages of lymphocyte populations.^{32–35} Among those that included cord blood cytokines that overlap with the set included in our study, none included PM_{2.5} and O₃ as exposures. It is therefore difficult to directly compare findings. Latzin and colleagues found PM₁₀ exposure late in pregnancy to be associated with lower IL-10 and higher IL-1 β levels, but no associations with MCP-1 and IL-6, in cord blood.³¹ Van den Hooven and colleagues

noted positive associations between full pregnancy PM₁₀ and NO₂ exposure and cord blood CRP levels.³⁰ Future investigations are necessary to confirm our findings.

Interestingly, levels of cytokines in maternal blood were not highly correlated with levels of the same cytokines in cord blood. Samples of maternal blood that were used for cytokine measurement were collected in mid-pregnancy, whereas the cord blood samples were collected at delivery, and thus the difference in the timing of sample collection may be a potential explanation for the discordance. Additionally, maternal and cord blood inflammatory biomarker levels were measured using different assays and labs and were sampled from different tissues, which may explain the large differences in concentrations we observed. It is also possible that this contributed to the low correlations, however, by using the Spearman correlation coefficient, which is calculated based on rank order, we expect the correlations to be somewhat robust to measurement error. There is scarce evidence regarding concordance between maternal and cord blood immune profiles,^{68, 69} and a handful of *ex vivo* studies reported conflicting results regarding the degree of transplacental transfer of different cytokines.^{70–72} Studies comparing inflammatory biomarkers in maternal blood with those in amniotic fluid during mid-pregnancy have found some, but not strong, concordance.^{73, 74}

Changes in the maternal immune response during pregnancy have been linked to preterm birth^{41, 75, 76} and changes in body composition at birth,^{77–80} though not consistently.⁸¹ Our finding that elevated levels of maternal CRP were associated with modestly older gestational age at birth was unexpected, in light of some previous studies that have found positive associations between levels of CRP in amniotic fluid and plasma and preterm birth, albeit a weaker association in plasma compared to amniotic fluid.⁷⁶ Epidemiologic studies have found inverse associations between maternal CRP and birthweight,^{40, 80} length,⁷⁸ percent body fat,⁸⁰ and sum of skin folds.^{78, 79} While associations between maternal inflammation during pregnancy and adiposity have been noted in mid-childhood,⁸² we did not observe associations between maternal inflammatory biomarkers and body composition outcomes at five months postnatally. Since growth patterns in the first year differ from those later in life, it is possible that five months may be too early to observe alterations in body composition associated with maternal inflammation during pregnancy.

A recent study from the Healthy Start cohort found limited evidence for associations between the air pollution measures used in the present study and birth outcomes.⁴² We did not find associations between maternal inflammatory biomarkers and birth weight and infant body composition. Nevertheless, inflammation has been posited as a potential mechanistic link between *in utero* air pollution exposure and birth and outcomes in offspring.⁸³ Findings from an animal study⁸⁴ and a case-control study⁶⁴ have begun to elucidate this relationship; future prospective observational studies are necessary.

Our study has several strengths and limitations. We utilized data from a prospective, pre-birth cohort, which allowed for adjustment of several potential confounders. The air pollution profile of the Denver metropolitan area is unique, as it is characterized by relatively low levels of PM_{2.5} compared to other major urban areas, but comparatively high levels of O₃, which differs from other more highly studied geographic locations in the US.⁶⁵

Even with the relatively low levels of particulate matter and limited variability, we observed associations with maternal pro-inflammatory cytokines. Few studies have investigated this relationship with a comprehensive panel of cytokines in maternal and cord blood concomitantly. The findings from this study contribute to existing literature characterizing the poorly understood relationship between maternal and fetal immune profiles.

We estimated exposure to air pollution using inverse-distance-weighted interpolation from stationary monitoring stations and were therefore unable to capture smaller-scale spatial variability of living close to roads and other sources of air pollution. Participants self-reported their residential address at enrollment, thus, there is potential for misclassification due to participants moving during pregnancy, as well as indoor air pollution exposures⁸⁵ and exposures occurring outside of the area of residence, such as exposure during commuting and work. A prior study found pregnant women in New York did not move often, and when they did, it was over short distances.⁸⁶ Thus, large changes in exposure estimates due to residential mobility, were not observed.⁸⁶ While this information is unavailable for the Colorado-based Healthy Start study, it provides evidence that residential mobility may not result in substantial measurement error. Further, the air pollution measurement error is not likely to be related to the inflammatory biomarkers, and thus, may result in an underestimation of associations or a reduction in power.⁸⁷ Our analysis was limited to PM_{2.5} and O₃, however, other components of air pollution and mixtures of air pollutants may be of additional interest.

For some women, the blood sample collection occurred prior to the end of the first trimester. While this does not introduce an issue of reverse causality, it is possible that it may lead to a small amount of measurement error. Further, the levels of maternal inflammatory biomarkers fluctuate throughout pregnancy,⁸⁸ and the samples in our study were collected at a single mid-pregnancy timepoint. Therefore, the conclusions based on findings from our study cannot be generalized to levels of inflammatory biomarkers in other time periods during pregnancy. Despite this limitation, there was little variation in the gestational ages at maternal blood sample collection (mean; 191, SD; 17 days), and thus, our data consistently captured the mid-pregnancy time-period; future studies are necessary to characterize these relationships with levels of early- and late- pregnancy inflammatory biomarkers. Additionally, our study was limited to six inflammatory biomarkers, three in maternal blood, and six in cord blood, and high throughput immune profile platforms are now available, which allow for the analysis of several hundred immune biomarkers simultaneously. Finally, we included only pro-inflammatory biomarkers from the mother, which did not allow for the investigation of composite indices with relative measures of pro- to anti-inflammatory measures.⁶⁹

5. Conclusions

We observed estimated ambient concentrations of PM_{2.5} and O₃ in early pregnancy to be associated with maternal levels of pro-inflammatory cytokines, during mid-pregnancy. However, levels of these maternal inflammatory biomarkers were not associated with infant birth weight, nor body composition at birth and at five months. There was some evidence supporting relationships between PM_{2.5} and O₃ exposure and inflammatory biomarkers in

cord blood at delivery, but patterns were inconsistent, and generally 95% confidence intervals included the null. Our findings highlight the potential for environmental pollutants during pregnancy to alter the maternal immune response. Future research may investigate the health implications of a heightened inflammatory response associated with air pollution exposure during pregnancy.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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The Colorado Multiple Institutional Review Board approved all study protocols.

Abbreviations

BMI	body mass index
CI	confidence interval
CRP	C-reactive protein
ELISA	enzyme-linked immunosorbent assay
EPA	Environmental Protection Agency
GDM	gestational diabetes mellitus
ICC	intra-class correlation coefficient
IL	interleukin
IQR	interquartile range
MCP-1	monocyte chemoattractant protein 1
NO₂	nitrogen dioxide
O₃	ozone
PM_{2.5}	particulate matter 2.5 micrometers
PM₁₀	particulate matter 10 micrometers
ppb	part per billion
SGA	small for gestational age

TNF α tumor necrosis factor- α

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Highlights

- Exposure to particulate matter $PM_{2.5}$ and ozone was estimated for 515 pregnant women
- Inflammatory biomarkers were measured in maternal and cord blood
- Early pregnancy air pollution associated with mid-pregnancy maternal inflammation
- Inconsistent associations between prenatal air pollution and cord blood inflammation
- Maternal inflammatory markers did not predict infant outcomes

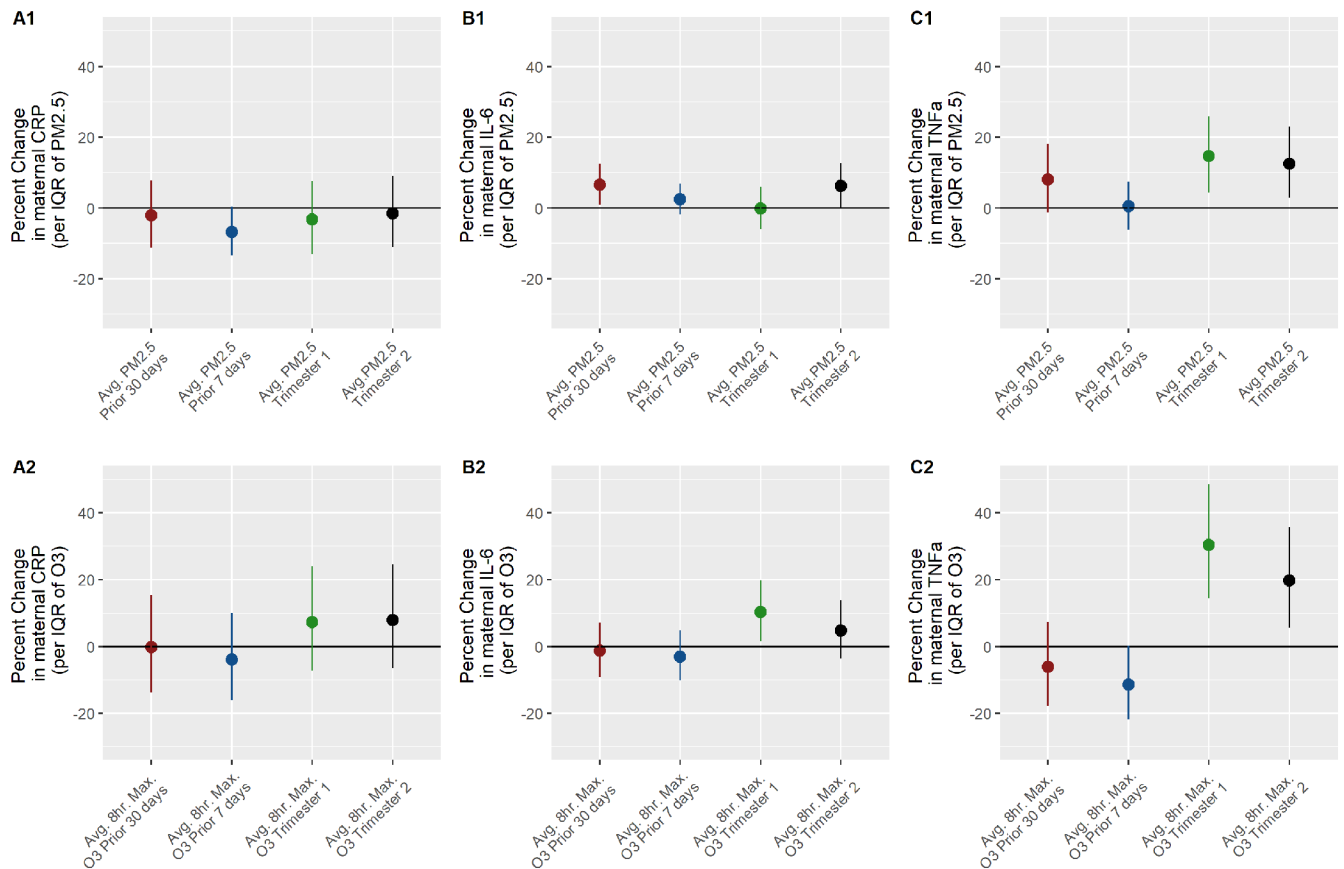


Figure 1:

A1 and A2. Associations between average PM_{2.5} and 8-hour maximum O₃ at the residential address during pregnancy and maternal CRP, respectively; B1 and B2. Maternal IL-6; C1 and C2. Maternal TNFα. Adjusted for maternal age, pre-pregnancy BMI, race, smoking status, median income in the Census tract, education, parity, infant sex, gestational age at sample collection, and the corresponding co-pollutant during the specified pregnancy period. Sample sizes differ by model due to missing PM_{2.5} exposure data (n=number missing): prior 30 days to sample collection, n=31; prior 7 days to sample collection, n=30; trimester 1, n=46; trimester 2, n=44.

Abbreviations: IQR, interquartile range; PM_{2.5}, particulate matter with diameter 2.5 micrometers; O₃, average 8-hour maximum ozone concentration in parts per billion; CRP, C-reactive protein; IL-6 interleukin-6; TNFα, tumor necrosis factor-α.

Table 1:

Characteristics of 515 eligible mother-infant pairs in the Healthy Start Study.

Characteristic	Mean \pm SD, Median (IQR), or N (%)
<i>Covariates</i>	
Maternal age at delivery (years)	27.6 \pm 6.2
Pre-pregnancy body mass index (kg/m ²)	25.9 \pm 6.5
Race/ethnicity	
Hispanic	128 (25)
Non-Hispanic white	273 (53)
Non-Hispanic black	79 (15)
All others	35 (7)
Maternal smoking during pregnancy (any)	49 (10)
Median annual income in the Census tract (per \$1,000)	64.1 \pm 26.7
Maternal education completed	
Less than 12 th grade	79 (15)
High school degree or GED	95 (18)
Some college or Associate's degree	106 (21)
Four years of college (BA, BS)	115 (22)
Graduate degree (Master's, Ph.D.)	120 (23)
Season of conception	
Winter	163 (32)
Spring	97 (19)
Summer	90 (18)
Fall	165 (32)
Mode of delivery= Caesarean	110 (21)
Previous pregnancy	323 (63)
Season of birth	
Winter	109 (21)
Spring	136 (26)
Summer	148 (29)
Fall	122 (24)
Infant sex = male	276 (54)
Gestational age at sample collection (days)	191 \pm 17
Gestational age at birth (days)	276 \pm 9
Age at follow-up visit (months) ^a	5.1 \pm 1.2
<i>Exposures</i>	
Average PM _{2.5} ($\mu\text{g}/\text{m}^3$) ^a	
Prior 30 days from sample collection	7.13 (6.08-8.26)
Prior 7 days from sample collection	6.92 (5.53-8.61)
Trimester 1	7.52 (6.89-8.14)
Trimester 2	7.39 (6.78-8.09)

Characteristic	Mean \pm SD, Median (IQR), or N (%)
Trimester 3	7.48 (6.84-8.29)
Full Pregnancy	7.51 (7.13-7.86)
Prior 30 days to delivery	7.24 (6.33-8.42)
Prior 7 days to delivery	6.73 (5.58-8.20)
Average 8-hr max O ₃ (ppb)	
Prior 30 days to sample collection	45.1 (32.7-54.2)
Prior 7 days to sample collection	44.1 (33.4-54.6)
Trimester 1	42.1 (33.8-53.7)
Trimester 2	42.6 (33.9-52.3)
Trimester 3	46.1 (35.0-54.6)
Full Pregnancy	44.0 (40.9-47.1)
Prior 30 days to delivery	48.0 (34.1-55.9)
Prior 7 days to delivery	47.0 (35.4-55.3)
<i>Outcomes</i>	
Birth weight (g)	3275 \pm 427
Fat mass at birth (g) ^a	285.3 \pm 143.1
Fat free mass at birth (g) ^a	2843.4 \pm 333.9
Adiposity at birth (%) ^a	8.9 \pm 3.8
Fat mass at postnatal follow-up visit (g) ^a	1708.6 \pm 526.8
Fat free mass at postnatal follow-up visit (g) ^a	5167 \pm 643.4
Adiposity at postnatal follow-up visit (%) ^a	24.5 \pm 5.6

Abbreviations: SD, standard deviation; IQR, interquartile range; PM_{2.5}, particulate matter with diameter \leq 2.5 micrometers; O₃, average 8-hour maximum ozone concentration in parts per billion.

^aMissing data for the following variables (n=number missing): age at postnatal follow-up, n=181; average PM_{2.5} Prior 30 days to sample collection, n=31; average PM_{2.5} Prior 7 days to sample collection, n=30; average PM_{2.5} Trimester 1, n=46; average PM_{2.5} Trimester 2, n=44; average PM_{2.5} Trimester 3, n=17; average PM_{2.5} Prior 30 days to delivery, n=31; average PM_{2.5} Prior 7 days to delivery, n=32; average PM_{2.5} Full pregnancy, n=4; fat mass at birth, n=18; fat free mass at birth, n=18; adiposity at birth, n=18; fat mass at postnatal follow-up, n=181; fat free mass at postnatal follow-up, n=181; adiposity at postnatal follow-up, n=181.

Table 2:

Concentrations and pairwise spearman correlations of cord blood and maternal inflammatory biomarkers among 515 eligible mother-infant pairs in the Healthy Start Study.

Characteristic	Median (IQR)	Spearman Correlation Coefficient								
		Cord blood				Maternal blood				
		CRP (mg/L) ^a	IL-6 (pg/mL)	IL-8 (pg/mL)	IL-10 (pg/mL)	MCP-1 (pg/mL)	TNF α (pg/mL)	CRP (mg/L)	IL-6 (pg/mL)	TNF α (pg/mL)
Cord blood										
CRP (mg/L) ^a	0.09 (0.04, 0.18)	1.00	0.28	0.28	0.27	0.05	0.02	0.26	0.15	-0.02
IL-6 (pg/mL)	9.85 (4.80, 24.46)		1.00	0.64	0.53	0.32	0.24	0.02	-0.02	0.00
IL-8 (pg/mL)	16.32 (8.20, 33.64)			1.00	0.56	0.42	0.33	0.04	0.02	0.03
IL-10 (pg/mL)	12.85 (8.54, 22.85)				1.00	0.26	0.31	0.05	0.00	0.04
MCP-1 (pg/mL)	665.44 (484.24, 909.20)					1.00	0.64	0.02	-0.03	-0.08
TNF α (pg/mL)	32.33 (24.84, 41.59)						1.00	0.02	-0.06	-0.05
Maternal blood										
CRP (mg/L)	4.48 (2.49, 8.16)							1.00	0.50	0.02
IL-6 (pg/mL)	1.36 (0.97, 1.99)								1.00	0.10
TNF α (pg/mL)	1.11 (0.69, 1.81)									1.00

Abbreviations: IQR, interquartile range; CRP, C-reactive protein; IL-6 interleukin-6; IL-8 interleukin-8; IL-10 interleukin-10; MCP-1 monocyte chemoattractant protein 1; TNF α , tumor necrosis factor- α .

^aMissing data for cord blood CRP (n=1).

Table 3:

Associations between average PM_{2.5} and 8-hour maximum O₃ at the residential address during pregnancy and cord blood inflammatory biomarkers.

	CRP (mg/L)	IL-6 (pg/mL)	IL-8 (pg/mL)	IL-10 (pg/mL)	MCP-1 (pg/mL)	TNF α (pg/mL)
	% change (95% CI) ^d per IQR	% change (95% CI) ^d per IQR	% change (95% CI) ^d per IQR	% change (95% CI) ^d per IQR	% change (95% CI) ^d per IQR	% change (95% CI) ^d per IQR
Trimester 1 (n=469)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	4.98 (-11.26, 24.19)	3.24 (-13.43, 23.11)	12.19 (-2.42, 28.98)	-2.76 (-12.49, 8.04)	-0.16 (-6.27, 6.34)	-0.03 (-5.22, 5.45)
Avg. 8-hr max O ₃ (ppb)	3.46 (-33.48, 60.92)	-16.88 (-47.59, 31.82)	-19.02 (-43.80, 16.70)	-9.70 (-31.48, 19.01)	-2.86 (-17.66, 14.60)	-0.30 (-13.30, 14.64)
Trimester 2 (n=471)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	7.57 (-8.03, 25.81)	4.81 (-11.37, 23.94)	2.67 (-10.10, 17.25)	2.95 (-6.82, 13.76)	-0.70 (-6.50, 5.47)	-4.62 (-9.48, 0.50)
Avg. 8-hr max O ₃ (ppb)	44.94 (-2.43, 115.31)	55.68 (1.68, 138.35)	39.78 (-0.24, 95.86)	6.07 (-17.68, 36.69)	14.89 (-1.42, 33.90)	1.65 (-11.00, 16.10)
Trimester 3 (n=498)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	-1.04 (-16.02, 16.61)	-3.96 (-19.35, 14.38)	-7.11 (-19.31, 6.94)	-3.74 (-13.30, 6.87)	-5.46 (-11.25, 0.71)	-4.82 (-9.91, 0.56)
Avg. 8-hr max O ₃ (ppb)	24.37 (-18.24, 89.19)	-7.09 (-40.59, 45.31)	4.63 (-27.04, 50.05)	12.99 (-13.55, 47.67)	0.79 (-14.26, 18.49)	-2.77 (-15.53, 11.92)
Full pregnancy (n=511)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	6.89 (-8.53, 24.90)	0.07 (-15.38, 18.34)	3.45 (-9.60, 18.40)	-2.17 (-11.50, 8.15)	-3.84 (-9.51, 2.18)	-6.04 (-10.82, -1.01)
Avg. 8-hr max O ₃ (ppb)	20.71 (-5.89, 54.83)	6.52 (-18.51, 39.26)	6.53 (-14.13, 32.16)	-0.67 (-15.38, 16.59)	2.63 (-6.86, 13.09)	-4.60 (-12.23, 3.70)
Prior 30 days to delivery (n=483)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	-2.69 (-16.99, 14.07)	2.61 (-13.22, 21.32)	1.27 (-11.54, 15.93)	-0.29 (-9.85, 10.30)	3.49 (-2.64, 10.00)	1.86 (-3.25, 7.23)
Avg. 8-hr max O ₃ (ppb)	12.32 (-23.03, 63.91)	-32.19 (-54.47, 0.99)	-16.67 (-39.58, 14.93)	12.34 (-11.62, 42.80)	-8.85 (-21.16, 5.37)	-6.01 (-16.82, 6.21)
Prior 7 days to delivery (n=488)						
Avg. PM _{2.5} ($\mu\text{g}/\text{m}^3$)	-0.30 (-9.21, 9.48)	2.37 (-7.23, 12.97)	4.06 (-3.88, 12.66)	0.63 (-5.10, 6.71)	0.31 (-3.23, 3.98)	1.03 (-2.05, 4.21)
Avg. 8-hr max O ₃ (ppb)	5.64 (-24.57, 47.96)	-20.89 (-44.52, 12.79)	-5.51 (-28.99, 25.75)	7.63 (-12.86, 32.94)	4.50 (-8.18, 18.94)	2.75 (-8.10, 14.87)

Abbreviations: 95% CI, 95% Confidence Interval; PM_{2.5}, particulate matter with diameter ≤ 2.5 micrometers; O₃, average 8-hour maximum ozone concentration in parts per billion; IQR, interquartile range; CRP, C-reactive protein; IL-6 interleukin-6; IL-8 interleukin-8; IL-10 interleukin-10; MCP-1, monocyte chemoattractant protein 1; TNF α , tumor necrosis factor- α .

All cord blood inflammatory biomarker outcomes were natural-log transformed.

^aAdjusted for maternal age, pre-pregnancy BMI, race, smoking status, median income in the Census tract, education, mode of delivery, parity, season of birth, infant sex, and the corresponding co-pollutant during the specified pregnancy period.

Table 4:

Associations between maternal inflammatory biomarkers during pregnancy and infant birth and adiposity outcomes.

	Birth weight (g) (n=515)	Gestational age (days) (n=515)	Adiposity at birth (%) (n=497)	Adiposity at postnatal (~5 months) follow-up (%) (n=334)
	β (95% CI) per log-unit ^a	β (95% CI) per log-unit ^a	β (95% CI) per log-unit ^a	β (95% CI) per log-unit ^b
CRP (mg/L)	12.48 (-31.31, 56.27)	1.30 (0.40, 2.21)	0.06 (-0.34, 0.45)	0.43 (-0.26, 1.12)
IL-6 (pg/mL)	10.51 (-66.94, 87.95)	0.24 (-1.37, 1.86)	0.23 (-0.47, 0.93)	-0.04 (-1.30, 1.21)
TNF α (pg/mL)	23.30 (-22.72, 69.31)	0.29 (-0.67, 1.25)	-0.11 (-0.53, 0.31)	-0.14 (-0.91, 0.63)

Abbreviations: 95% CI, 95% Confidence Interval; CRP, C-reactive protein; IL-6 interleukin-6; TNF α , tumor necrosis factor- α

All maternal inflammatory biomarker outcomes were natural log transformed.

^a Adjusted for race, maternal age, median income in the Census tract, pre-pregnancy BMI, education, smoking status, gestational age at sample collection, infant sex, and parity.

^b Additionally adjusted for infant age at the time of adiposity measurement.