Multi-pollutant exposure profiles associated with term low birth weight in Los Angeles County

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ABSTRACT

Research indicates that multiple outdoor air pollutants and adverse neighborhood conditions are spatially correlated. Yet health risks associated with concurrent exposure to air pollution mixtures and clustered neighborhood factors remain underexplored. Statistical models to assess the health effects from pollutant mixtures remain limited, due to problems of collinearity between pollutants and area-level covariates, and increases in covariate dimensionality. Here we identify pollutant exposure profiles and neighborhood contextual profiles within Los Angeles (LA) County. We then relate these profiles with term low birth weight (TLBW). We used land use regression to estimate NO2, NO, and PM2.5 concentrations averaged over census block groups to generate pollutant exposure profile clusters and census block group-level contextual profile clusters, using a Bayesian profile regression method. Pollutant profile cluster risk estimation was implemented using a multilevel hierarchical model, adjusting for individual-level covariates, contextual profile cluster random effects, and modeling of spatially structured and unstructured residual error. Our analysis found 13 clusters of pollutant exposure profiles. Correlations between study pollutants varied widely across the 13 pollutant clusters. Pollutant clusters with elevated NO2, NO, and PM2.5 concentrations exhibited increased log odds of TLBW, and those with low PM2.5, NO2, and NO concentrations showed lower log odds of TLBW. The spatial patterning of pollutant cluster effects on TLBW, combined with between-pollutant correlations within pollutant clusters, imply that traffic-related primary pollutants influence pollutant cluster TLBW risks. Furthermore, contextual clusters with the greatest log odds of TLBW had more adverse neighborhood socioeconomic, demographic, and housing conditions. Our data indicate that, while the spatial patterning of high-risk pollutant clusters largely overlaps with adverse contextual neighborhood cluster, both contribute to TLBW while controlling for the other.

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1. Introduction

Evidence has been accumulating that birth outcomes may be particularly sensitive to air pollution mixtures, specifically components related to traffic sources of airborne particulate matter (PM) (Bell et al., 2010; Laurent et al., 2014; Wilhelm et al., 2011a). While earlier research has linked increased prevalence of term low birth weight (TLBW) with various outdoor air pollutants including NO2, NO, and PM2.5 (Geer, 2014; Ritz and Wilhelm, 2008), most evidence relied on single pollutant modeling of exposures (Ritz and Wilhelm, 2008). A number of studies (Brauer and Tamburic, 2009; Ghosh et al., 2012; Gouveia et al., 2004; Laurent et al., 2014; Le et al., 2012; Morello-Frosch et al., 2010; Wilhelm et al., 2011a, 2011b) investigated exposures to multiple pollutants in relation to birth outcomes; however, these studies are limited in assessing which combination of pollutants are most hazardous or how multipollutant health effects vary spatially. Despite there being no single exposure-measure-framework to holistically address the health effects of multipollutant exposures (Oakes et al., 2014), investigating health effects of profiles of multiple pollutants using clustering techniques has recently shown promise (Gu et al., 2012; Molitor et al., 2014a; Papathomas et al., 2010; Pirani et al., 2015; Qian et al., 2004; Zanobetti et al., 2014).

1.1. Multipollutant exposures

Considerable intra-urban spatial variations in outdoor air pollution concentrations exist, and recent research indicates that between-pollutant correlations and PM2.5 composition exhibit highly localized spatial patterns to create complex mixtures (Austin et al., 2013; Austin...
Spatially correlated air pollution mixtures correspond to localized sources, such as transportation-related emissions (Laurent et al., 2014), local industrial activities (Morello-Frosch et al., 2002; Zhu et al., 2011), or small-area commercial land uses (Morello-Frosch et al., 2002; Morello-Frosch et al., 2001). Factors that combine to determine exposure to spatially correlated pollutants from a particular source are complex and diverse, e.g., traffic-source driven exposures are influenced by traffic volumes and congestion, proximity to traffic, the types of fuel and engines, operating conditions of emitting sources, types of emitting sources, background air pollution levels, local meteorology, chemical reactions between pollutants, and local topographies (Austin et al., 2012; Boehmer et al., 2013; Cho et al., 2009; Greco et al., 2007; Hu et al., 2012; Janhäll et al., 2012; U.S. EPA, 2008; Zhang and Batterner, 2013).

Correlations across different pollutants hinder our ability to assess their individual or combined health effects, since estimates of effects may become unstable when adjusting for co-pollutants using regression techniques (Mauderly et al., 2010). Correlations between PM2.5 concentrations and nitrogen oxides (NO2 and NOx) are typically weak to moderate (Greco et al., 2008) or secondary PM2.5 (particles formed in the atmosphere (Fine et al., 2008)). Therefore, since some PM2.5 components represent “fresh” traffic emissions (i.e. ultrafine PM and black carbon), they can exhibit high correlations with outdoor concentrations of NOx as studies of urban air pollution from Asia, Europe, and North America demonstrate (Brauer et al., 2011; Dionisio et al., 2014; Janhäll et al., 2012; Levy et al., 2013; Tsai et al., 2015; Wang et al., 2014). Furthermore, the spatial variation in between-pollutant correlations also suggests a strong potential for a unique spatial pattern of multipollutant-related health risks, yet research on this question is lacking.

1.2. Contextual factors

Neighborhood-level “contextual” factors may also affect risk of birth outcomes (English et al., 2003; Morello-Frosch and Shenassa, 2006). Contextual effects are non-chemical stressors (Lewis et al., 2011) that arise when grouped neighborhood-level factors – such as socio-demographic or built environment (e.g. housing as one component of the built environment) factors – influence health outcomes across populations (Sheppard et al., 2012). Data from Southern California indicates that spatially clustered socioeconomic deprivation and racial segregation correlate with air pollution exposures, including pollutant mixtures (Molitor et al., 2011; Morello-Frosch et al., 2011; Morello-Frosch et al., 2002; Morello-Frosch and Shenassa, 2006; Su et al., 2012). In addition, compared to newer homes, older homes are shown to have higher indoor air pollution levels within LA County (Spengler et al., 1994). Older housing stock may further correlate with higher poverty, residential racial segregation patterns, substandard housing conditions, and a lack of compliance with building or sanitary codes (Shennassa et al., 2004). Moreover, research from LA County indicates that housing ventilation conditions may be associated with TLBW (Greco et al., 2013).

Air pollution and birth outcomes research studies, however, generally do not account for spatial clustering of multiple neighborhood-level vulnerabilities (i.e. race-ethnicity, poverty, and adverse housing conditions) related to exposure. This may confound multipollutant exposure-response relationships (Geer, 2014; Morello-Frosch et al., 2011; Morello-Frosch and Shenassa, 2006; Ponce, 2005; Ritz and Wilhelm, 2008). Given this gap in air pollution and health effects literature, our study includes contextual factors, as clustering covariates, to better control for highly correlated contextual factors known to influence differential exposures across socio-demographic groups and risk of TLBW.

1.3. Study objectives

The primary objectives of our study are to first identify profiles of exposure to multiple different air pollutants (pollutant profiles) for pregnant women within LA County, and secondly to assess whether and which pollutant profiles relate with elevated prevalence of TLBW. Additionally, our approach identifies pollutant profiles most likely related to primary traffic emissions, based on examination of the spatial patterning of pollutant exposure profiles and well established pollutant source emissions relationships.

2. Methods

2.1. Study population and birth outcomes

Electronic birth certificates from the California Department of Public Health provided the data on baby’s birth weight and individual-level covariates for LA County births during the years 2000–2006. Individual data from the birth records included maternal characteristics (age, race and ethnicity, education, total number of previous maternal births, and residential address) and information on the infant and birth (date of birth, abnormalities, birth season, gestational age at birth [as determined by self-report of last menstruation], birth weight and baby’s sex). The dataset was restricted to singleton births without apparent abnormalities, while births with extreme gestational days (<140 days or >320 days) and births with weight <500 g or >5000 g were excluded from the analysis. Such extreme values are likely attributable to recording errors. These data restrictions provided a sample size of 899,554. Finally, we defined TLBW as full-term (>259 gestation days) infants with a birth weight <2500 g, which further restricted the study population to term births, to provide a final sample size of 804,726 to assess the relationship between TLBW with neighborhood-level pollutant profile exposures. Human subjects research was approved through the University of California, Los Angeles’ Office of the Human Subjects Protection Program, the California Committee for the Protection of Human Subjects, and the University of Southern California’s Office for the Protection of Research Subjects. Geocoding of residential addresses are explained elsewhere (see Goldberg et al., 2008).

2.2. Exposure estimation

Two separate land use regression (LUR) models estimated individual-level exposures for PM2.5, NOx, and NO2 (Beckerman et al., 2013a, 2013b; Su et al., 2009). LUR estimates were temporally adjusted to derive the entire pregnancy average exposures. LUR exposure predictions for NO and NO2 were based on traffic volumes, truck routes, road networks, land use data, satellite-derived vegetation greenness and soil brightness, truck route slope gradients, and air monitoring data. NO2 and NO data were collected during 2-week time periods in Summer of 2006 and Winter of 2007, from over 200 monitoring locations (Su et al., 2009). The PM2.5 exposure estimates came from a LUR model that utilized long-term governmental monitoring data of PM2.5 measurements collected between 1998 through 2002 (Beckerman et al., 2013b). A machine learning deletion/substitution technique (Beckerman et al., 2013a) assessed as many as 70 covariates to develop the final PM2.5 LUR model, such as land use data (i.e. agricultural, barren, all developed land, high-density development, green space, water, and wetland), long-term traffic counts (1990–2001), and road networks from the year 2000 (Beckerman et al., 2013a; Jerrett et al., 2013).

We adjusted LUR exposure estimates temporally to derive “seasonalized” values that correspond to each pregnancy time span. For temporal adjustments, we first used daily air monitoring data from LA County between the years 1999–2006, for all monitors to
calculate an overall daily average for PM$_{2.5}$, NO$_2$, and NO$_x$. Pregnancy time period averages were then calculated for each pregnancy from these daily averages. As NO was not directly measured by these monitors, we subtracted the NO$_2$ pregnancy average from the NO$_x$ pregnancy average to derive NO pregnancy time period estimates. Temporal adjustments of the LUR estimates for NO$_2$ and NO were achieved using the following equation, which is similar in approach to our earlier work (see (Ghosh et al., 2012)):

\[
\text{NO}_2 \text{ LUR Pregnancy Average} = \text{NO}_2 \text{ LUR} \times \frac{\text{monitoring station pregnancy average NO}_2}{\text{2006 NO}_2 \text{ average for all monitoring stations}}.
\]

\[
\text{NO LUR Pregnancy Average} = \text{NO LUR} \times \frac{\text{monitoring station pregnancy average NO}}{\text{2006 NO average for all monitoring stations}}.
\]

Since the PM$_{2.5}$ LUR estimates represent long-term estimates spanning the study time period and some pregnancies began in 1999, we performed seasonal adjustments with the following equation:

\[
\text{PM}_{2.5} \text{ LUR Pregnancy Average} = \text{PM}_{2.5} \times \frac{\text{monitoring station pregnancy average PM}_{2.5}}{\text{PM}_{2.5} \text{ average for all monitoring stations (1999–2006)}}.
\]

Such temporal adjustment via region-wide monitoring station ratios has been validated for the purposes of estimating pregnancy exposures in birth outcomes studies (Ross et al., 2013).

All of the available data from the temporally adjusted LUR model estimates (N = 899,554) were then averaged over census block groups to develop air pollution exposure profiles at the census block group level. Data aggregation at the census block group-level for individual estimates was performed since we were interested in assessing between neighborhood multipollutant exposure-related TLBW risks. Moreover, implementation of the Bayesian profile regression using individual-level estimates with a dataset as large as ours is not feasible given the current computational limitations of the PReMiuM package (described below).

2.3. Bayesian profile regression

We developed the profile clusters using a non-parametric dimension reduction technique known as Bayesian profile regression, based on commonly used Dirichlet process mixture model methods (Neal, 2000). Profile regression is set in a Bayesian framework using Markov chain Monte Carlo (MCMC) methods. Bayesian profile regression uses covariate values to observe joint patterns within the covariate data. This approach was used in recent studies (Hastie et al., 2013; Molitor et al., 2014a; Molitor et al., 2014b; Molitor et al., 2011; Papathomas et al., 2012), including environmental epidemiology studies (Papathomas et al., 2010; Pirani et al., 2015; Vrijheid et al., 2014). This clustering approach is advantageous because it reduces the dimensionality of the covariate data and allows for examining health risks as they relate to joint patterns of exposure, while avoiding the pitfalls of exposure variables that are highly collinear. This approach is also quite flexible because it does not rely on setting a total number of allowable clusters, as seen with k-means clustering procedures (Austin et al., 2013; Austin et al., 2012; Gu et al., 2012). We implemented the profile regression using the PReMiuM package in R (Liverani et al., 2015).

Since our interest is in obtaining clustering that best fits the data for sub-regions within the LA County area, we utilized a feature of the PReMiuM package that excludes the outcome variable from the profile regression model (Liverani et al., 2015). We relied on “hard clustering” (Fang et al., 2011) in the sense that a census block group’s final allocation is to a single cluster. Cluster allocation is based on the “best” clustering derived from the Bayesian averaging process, rather than probabilistic allocation to several different clusters simultaneously (as in fuzzy or soft clustering). Briefly, for each census block group, j, a covariate profile is defined as $x_j = (x_{jk}, x_{j2}, ..., x_{jp})$, where every covariate, $x_{jp}$, $p = 1, ..., P$, within each profile represents a level of exposure for covariate p in region j. The primary model for cluster profiles was defined by a multivariate normal mixture model (Jain and Neal, 2004) that further integrates a Dirichlet process prior into the mixing distribution. For greater details on this Bayesian profile regression approach, the reader is referred to other recent works (Hastie et al., 2013; Liverani et al., 2015; Molitor et al., 2014a; Molitor et al., 2011; Molitor et al., 2010; Papathomas et al., 2012; Papathomas et al., 2010).

We performed two separate profile regressions to develop a set of two unique profile clusters to fit in the TLBW risk model. The first clustering procedure developed pollutant-only profile clusters. The second clustering procedure developed contextual-only profile clusters. The co-pollutants for our pollutant-only profile regression included average census block group-level concentrations for NO$_2$, PM$_{2.5}$, NO. Furthermore, since our LUR estimates were seasonalized and thus provide temporally resolved estimates of exposure for each pregnancy, we also performed pollutant profile regression across different birth seasons and across different birth years. We present the results of these seasonal and yearly pollutant clusters in the Supplemental materials (Figs. S1–S4), however, in this paper we focus on the overall pollutant profiles as described above since the spatial patterning and between-pollutant correlations were very stable across each of these different seasonal and yearly pollutant profile cluster analyses.

The contextual-only exposure profile regression utilized year 2000 U.S. census data and included census block group-level race-ethnicity (percent non-Hispanic White, percent non-Hispanic Black, and percent Hispanic), median household income, and percent of homes built prior to year 1950. Even though our multivariate risk model (described below) adjusts for individual-level maternal race-ethnicity, we included census block group-level racial/ethnic composition as a clustering contextual covariate under the rationale that area-level racial/ethnic composition may act as a contextual risk factor for TLBW separate from an individual’s race-ethnicity (Debbink and Bader, 2011). Similarly, while our multivariate model adjusts for maternal education as a marker of individual-level SES, we included census block group level median household income as a contextual SES variable in the clustering procedure, under the same rationale that area-level SES acts as a contextual risk factor for TLBW independent of individual-level SES (Grady, 2011; Grady, 2006). We included the percentage of homes built before 1950 since disparities in housing quality and other housing characteristics correlated with older housing may act as an important contextual risk factor in TLBW risk (Ghosh et al., 2013; Grady, 2011). Individual mothers were then assigned to both a pollutant cluster and a contextual cluster as determined by which census block group the mother resided in according to their address at time of delivery.

2.4. Multilevel risk model

Our multilevel logistic regression model was set in a Bayesian framework with pollutant profile clusters and contextual profile clusters used as separate random effects variables in the regression equation, along with spatially structured and unstructured independent error terms fit as additional random effects. The model specification is detailed in turn:

\[
\logit(p_i) = \alpha + V_i + Y_{i}^{\text{pollutant-cluster}} + \gamma_{i}^{\text{contextual-cluster}} + S_i + \epsilon_{ij}
\]

where $p_i$ denotes the logit of TLBW ($y_i = 1$) for individual $i$, $V_i$ represents the individual-level covariate fixed effects, $Y_{i}^{\text{pollutant-cluster}} \sim N(0, \sigma_{\text{pollutant-cluster}})$, $k = 1, \ldots, 13$ represents the random effects for the pollutant-clusters and $\gamma_{i}^{\text{contextual-cluster}} \sim N(0, \sigma_{\text{contextual-cluster}})$, $c = 1, \ldots, 14$ represents the contextual cluster random effects. Following Gelman and Hill (2006), we use the notation $k[i]$ to denote the pollutant profile group k to which individual i belongs and c[i] to denote the contextual profile group c to which individual i belongs. Thus, each pollutant random error term represents the variation in TLBW prevalence in the pollutant profile clusters and likewise each contextual random error term represents the variation in TLBW in the contextual clusters. In other words, the cluster random effect can be interpreted.
as measuring the change in baseline log odds of TLBW for individual i in cluster k, when all other covariates in the model are set to zero (see Molitor et al., 2010).

Regarding the spatial and independent residual error terms, here $S_j$ and $e_j$ denote spatial and independent residual error terms, respectively, with the restriction $\sum_j S_j = 0$ imposed for indefinability reasons. While the independent error term is defined in the standard way as $e_j \sim N(0, \sigma^2)$, the spatial error term is defined as,

$$S_{jk} \sim N\left(\sum_{j \neq k} w_{jk} S_j, \sum_{j \neq k} w_{jk}\right),$$

where the weights $w_{jk}$ are elements of the zero-one neighborhood adjacency matrix defined to be equal to one when census tracts i, k are adjacent and zero otherwise. This approach implements the Besag-York-Molly (BYM) model (Besag et al., 1991) and has been successfully employed in a variety of exposure/health association studies (Molitor et al., 2007). Given the large number of records in the dataset, we “pre-clustered” exposure profiles as described in our clustering section and then used the R-INLA (integrated nested Laplace approximations) package to implement the Bayesian multilevel random effects model described in equation (1) above. R-INLA estimates Bayesian posterior marginal distributions (Rue et al., 2014; Rue and Martino, n.d.) without relying on computationally intensive Markov chain Monte Carlo techniques (Gilks et al., 1998).

2.5. Assessing uncertainty in pollutant random effects

As our analysis is in a Bayesian framework, with random effects terms for each pollutant cluster and for each contextual cluster, we assessed the uncertainty with respect to the random effect for each cluster on the baseline log odds of TLBW. For instance, we calculated the posterior probability that a specific profile cluster’s posterior distribution of baseline log odds for TLBW $\{\log(P[\text{TLBW}=1]) = \theta_t\}$ is above the overall baseline log odds for TLBW $\{\theta\}$ (Papathomas et al., 2010). Said another way, for each cluster we calculate the probability as $P(h > 0)$, with probability values close to 1 indicative of a high probability for a baseline log odds above zero for each cluster (evidence for adverse effect). Conversely, a posterior probability close to zero is indicative of a low probability for a baseline log odds above zero for each cluster (evidence for no adverse effect). The posterior probabilities for each pollutant cluster and for each contextual cluster were then mapped in ArcGIS V.10.1 (Redlands, CA) to investigate the spatial distribution of these clusters effects on the log odds of TLBW. These kinds of probability effect maps are commonly used in Bayesian modeling of pollutant data. Summary statistics for each pollutant overall, and pre-clustered exposure profiles are in Table 1. These data summaries are color coded to help indicate which exposure profile clusters have either elevated (red), typical (green), or lowered (blue) pollutant concentrations compared to the overall concentrations. According to pollutant summaries in Table 1, the elevated NO3 clusters are pollutant clusters P1, P7, P9, P10, and P13. The elevated NO pollutant clusters are the same as for NO2, plus P12. The elevated PM2.5 pollutant clusters are clusters P6, P7, P9, P10 and P13. Four of the pollutant clusters show elevated levels for all pollutants, including P7, P9, P10, and P13. Whereas pollutant profile clusters P1, P2, P4, P5, and P11 show low levels for all pollutants. We mapped the spatial distributions of pollutant clusters in Fig. 2. Clusters with high concentrations for all pollutants are mostly within the downtown/central area of LA, and a relatively fewer number of census block groups in the eastern section of the county (P13).

2.6. Covariates

Individual-level covariates adjusted for were maternal factors including age at delivery (<20 years, 20–24 years, 25–29 years, 30–34 years, ≥35 years), race-ethnicity (non-Hispanic White, non-Hispanic Black, Hispanic, Asian, and Other race), highest education level attained (<9 years, 9–12 years, 13–15 years, and ≥16 years), parity, along with infant factors such as gestational days, gestational days squared, and infant sex (male/female).

2.7. Characterization of pollutant clusters

In order to infer which clusters are most likely affected by near-highway traffic emission (or primary emissions) we characterized each pollutant cluster in terms of their respective pollutant ratios, between-pollutant correlations, and maternal residential distance to major highways. Such metrics have shown to be helpful in terms of assessing sources of emissions related to near road vehicle traffic (Austin et al., 2012; Janhäll et al., 2012; Laurent et al., 2014; Levy et al., 2013). Also, since both NO and NO2 could be highly correlated with PM2.5 under certain emissions scenarios, we normalized NO and NO2 concentration to PM2.5 as described in Austin et al. (2012). This normalization helps to indicate which pollutant clusters have elevated NO and NO2 concentrations, after accounting for their overall relationship with PM2.5. We obtained the normalized concentrations by calculating the cluster-specific ratio of NO or NO2 to PM2.5 and dividing by the overall study area ratio of NO or NO2 to PM2.5.

3. Results

We observed an overall TLBW prevalence of 2.07% (95% CI: 2.04–2.11, n = 16,694) for the study population. Our data also showed spatial autocorrelation at the census tract level with respect to prevalence of TLBW (Fig. S5, Supplemental materials). Average census block group concentrations of NO2, NO, and PM2.5 were 22.49 ppb (interquartile range [IQR]: 19.68, 25.30 ppb), 21.84 ppb (IQR: 16.05, 26.11), and 16.94 µg/m3 (IQR: 15.96, 18.18), respectively (Table 1).

3.1. Pollutant and contextual variable correlations

As indicated in Fig. 1, there is evidence of strong correlation (Pearson’s r) for between pollutant concentrations, most notably for NO2 and NO. The positive correlations between PM2.5 and NO2 and between PM2.5 and NO are considerably weaker (Fig. 1). However, we did find that the between-pollutant correlations for PM2.5 and the NOx species varied widely across LA County (Fig. S7, Supplemental materials). Correlations between our contextual variables and the study pollutants ranged from only weak to moderate (range: −0.53 to 0.64) (Fig. 1). Correlations between the contextual variables on the other hand were stronger (range: −0.66 to 0.64).

3.1.1. Pollutant clusters

The profile regression identified 13 pollutant profile clusters (P1–P13) from the 6280 census block groups from which we had complete air pollution data. Summary statistics for each pollutant overall, and stratified by cluster, are in Table 1. These data summaries are color coded to help indicate which exposure profile clusters have either elevated (red), typical (green), or lowered (blue) pollutant concentrations compared to the overall concentrations. According to pollutant summaries in Table 1, the elevated NO3 clusters are pollutant clusters P3, P7, P9, P10, and P13. The elevated NO pollutant clusters are the same as for NO2, plus P12. The elevated PM2.5 pollutant clusters are clusters P6, P7, P9, P10 and P13. Four of the pollutant clusters show elevated levels for all pollutants, including P7, P9, P10, and P13. Whereas pollutant profile clusters P1, P2, P4, P5, and P11 show low levels for all pollutants. We mapped the spatial distributions of pollutant clusters in Fig. 2. Clusters with high concentrations for all pollutants are mostly within the downtown/central area of LA and South-Central LA (clusters P7, P9, and P10), and a relatively fewer number of census block groups in the eastern section of the county (P13).

Between pollutant correlations for each pollutant cluster are indicated in Table 2 (see Supplemental materials for the spatial pattern of pollutant correlations). In Fig. 3 we present the distributions of NO/NO2 ratios and the normalized NOx and normalized NO values for each pollutant cluster, while the median residential distances to major highways throughout the county are shown in Fig. 4. According to these pollutant and near-highway exposure metrics, mothers residing in pollutant clusters P9 and P10 are most likely exposed to higher levels of primary traffic pollution since these two clusters are characterized by high NO3 ratios and high normalized NO and NO2 concentration. Additionally, P9 and P10 are characterized by low correlations between NO and NO2 (suggestive of near roadway emissions) and elevated PM2.5.
concentrations. Moreover, clusters P9 and P10 have the shortest median residential distance to major highways (478 m and 230 m respectively). Even though pollutant cluster P13 exhibited the highest average PM$_{2.5}$ in addition to elevated NO$_2$ and NO, this cluster is emblematic of combined high levels of regional sources of PM$_{2.5}$ and secondary particulate formation from traffic emissions. For instance, compared to all other clusters, P13 shows some of the strongest positive correlation between NO$_2$ and PM$_{2.5}$, a low NO/NO$_2$ ratio and low normalized NO and NO$_2$ levels. The pollutant metrics including maternal distance to highways (Fig. 4), between-pollutant correlations (Table 2), normalized NO concentration, and NO/NO$_2$ ratio (Fig. 3), indicate that all other pollutant profile clusters are more characteristic of secondary traffic-related pollution and other non-local (or regional transport) PM$_{2.5}$ sources instead of primary traffic-related.

3.1.2. Pollutant cluster random effects

For each pollutant cluster, Table 3 presents the total number of births, the number of TLBW cases, percent prevalence of TLBW, and the posterior means and 95% credible intervals for the pollutant cluster random effects. The cluster-specific effects represent the variation in the baseline log odds of TLBW, after adjusting for individual-level covariates, contextual cluster random effects, and accounting for spatially structured and unstructured residual error. In Table 3 we also present the posterior probability that a pollutant cluster effect is above the overall baseline log odds (probability = 99.5%). The income mean effect (0.124) and the highest probability for a baseline log odds of TLBW (probability = 99.5%). The income distribution for cluster C6 is significantly below the overall median income for LA County and consists of a significantly lower percentage of non-Hispanic Whites. Additionally, we find that cluster C6 has above average percentage of homes older than 1950, percentage of Blacks, and percentage of Hispanics. The next two highest probability contextual clusters are clusters C11 and C14, with elevated baseline TLBW log odds probabilities of 90.2% and 81.6%, respectively. While contextual clusters C11 and C14 have elevated percentages for homes older than 1950 and elevated percentages for Black populations, only C11 has significantly lower median income levels. All other clusters fell below 80% for elevated baseline TLBW log odds probabilities.

We also mapped the spatial distribution of contextual profile cluster effect probabilities in Fig. 7. This map indicates that contextual profile clusters with the highest probabilities for an elevated baseline log odds of TLBW are mostly in the urban core of LA County (central LA and south central LA). As anticipated, we find a large degree of spatial overlap between pollutant profile clusters with elevated effects on TLBW and contextual profile clusters with elevated effects on TLBW. Despite this spatial overlap, the two types high risk clusters take on distinct spatial patterns from one another.

3.3. Fixed effects results

In Table 5 we summarize each fixed effects estimate and corresponding 95% credible intervals from the multilevel spatial model.

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Table 1
Summary statistics of mean census block group-level pollutant concentrations for pollutant exposure profile clusters.

<table>
<thead>
<tr>
<th>Pollutant cluster (number of census block groups)</th>
<th>PM$_{2.5}$ µg/m$^3$ (IQR)</th>
<th>NO$_2$ ppb (IQR)</th>
<th>NO ppb (IQR)</th>
</tr>
</thead>
<tbody>
<tr>
<td>P1 (329)</td>
<td>13.20 (11.95, 14.55)</td>
<td>18.19 (15.58, 20.79)</td>
<td>10.99 (9.30, 12.87)</td>
</tr>
<tr>
<td>P2 (242)</td>
<td>12.83 (12.03, 13.69)</td>
<td>21.34 (19.28, 23.2)</td>
<td>19.17 (16.07, 21.60)</td>
</tr>
<tr>
<td>P3 (633)</td>
<td>14.93 (14.58, 15.30)</td>
<td>28.14 (26.12, 29.64)</td>
<td>29.10 (23.46, 33.90)</td>
</tr>
<tr>
<td>P5 (624)</td>
<td>15.96 (15.27, 16.52)</td>
<td>21.94 (20.61, 23.33)</td>
<td>22.50 (21.64, 25.21)</td>
</tr>
<tr>
<td>P6 (500)</td>
<td>18.87 (18.31, 19.41)</td>
<td>17.84 (16.12, 19.66)</td>
<td>15.40 (13.24, 18.20)</td>
</tr>
<tr>
<td>P7 (3715)</td>
<td>18.16 (17.79, 18.56)</td>
<td>25.63 (24.06, 27.31)</td>
<td>20.36 (26.11, 33.27)</td>
</tr>
<tr>
<td>P8 (96)</td>
<td>16.00 (16.40, 17.18)</td>
<td>16.46 (13.23, 18.78)</td>
<td>22.34 (16.29, 30.91)</td>
</tr>
<tr>
<td>P9 (513)</td>
<td>17.66 (17.12, 18.15)</td>
<td>31.40 (30.14, 33.27)</td>
<td>42.31 (37.75, 46.37)</td>
</tr>
<tr>
<td>P10 (52)</td>
<td>17.23 (16.84, 17.67)</td>
<td>42.51 (39.91, 44.29)</td>
<td>74.08 (63.40, 80.79)</td>
</tr>
<tr>
<td>P11 (28)</td>
<td>10.01 (8.93, 10.38)</td>
<td>14.44 (7.44, 16.80)</td>
<td>18.67 (10.86, 23.07)</td>
</tr>
<tr>
<td>P12 (52)</td>
<td>16.87 (16.28, 17.47)</td>
<td>16.51 (15.78, 17.42)</td>
<td>59.62 (53.45, 63.88)</td>
</tr>
<tr>
<td>P13 (96)</td>
<td>22.77 (21.89, 23.75)</td>
<td>31.39 (29.55, 32.85)</td>
<td>50.69 (28.13, 33.10)</td>
</tr>
<tr>
<td>Overall (5280)</td>
<td>16.84 (15.84, 18.13)</td>
<td>24.41 (21.33, 27.45)</td>
<td>25.99 (19.12, 30.94)</td>
</tr>
</tbody>
</table>

---

3.2. Contextual clusters

The profile regression determined 14 distinct contextual profiles clusters (C1–C14) and are summarized in Fig. 6. Since the contextual clusters were developed separately from the pollutant clusters, these contextual clusters are distinct from the pollutant clusters. Table 4 presents a summary of the contextual profile cluster random effects. Again, these random effects represent the variation in the baseline log odds of TLBW across contextual clusters in our multilevel model. Compared to all other clusters, contextual cluster C6 showed the highest posterior mean effect (0.124) and the highest probability for a baseline log odds above the overall baseline log odds (probability = 99.5%). The income distribution for cluster C6 is significantly below the overall median income for LA County and consists of a significantly lower percentage of non-Hispanic Whites. Additionally, we find that cluster C6 has above average percentage of homes older than 1950, percentage of Blacks, and percentage of Hispanics. The next two highest probability contextual clusters are clusters C11 and C14, with elevated baseline TLBW log odds probabilities of 90.2% and 81.6%, respectively. While contextual clusters C11 and C14 have elevated percentages for homes older than 1950 and elevated percentages for Black populations, only C11 has significantly lower median income levels. All other clusters fell below 80% for elevated baseline TLBW log odds probabilities.

We also mapped the spatial distribution of contextual profile cluster effect probabilities in Fig. 7. This map indicates that contextual profile clusters with the highest probabilities for an elevated baseline log odds of TLBW are mostly in the urban core of LA County (central LA and south central LA). As anticipated, we find a large degree of spatial overlap between pollutant profile clusters with elevated effects on TLBW and contextual profile clusters with elevated effects on TLBW. Despite this spatial overlap, the two types high risk clusters take on distinct spatial patterns from one another.

3.3. Fixed effects results

In Table 5 we summarize each fixed effects estimate and corresponding 95% credible intervals from the multilevel spatial model.

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Fig. 1. Pearson correlation matrix of census block group level averages for air pollutants, contextual variables, and pollutant metrics.
Individual-level factors associated with odds of TLBW were female infant sex, gestational days, and gestational days squared, as well as maternal factors such as parity, age, race-ethnicity, and education level.

4. Discussion

Our Bayesian profiling approach highlights the varied and distinct spatial patterns of pollutant exposure profiles and how such exposures contribute to TLBW within the context of clustered indicators of socio-demographic and housing. While exposure profile clustering has been used in previous epidemiologic studies (Molitor et al., 2014b; Papathomas et al., 2010; Pirani et al., 2015; Zanobetti et al., 2014), no such studies have examined birth outcomes. Our clustering procedure and multilevel analysis provided concentration estimates for pollutants and TLBW risk estimates for place-based air pollution mixtures across LA County that take on a strong spatial structure.

4.1. Pollutant profile clusters and effects on TLBW

We find that census block groups adjacent to major highways in the downtown/metro LA, South-Central LA, and parts of eastern LA County show elevated PM$_{2.5}$, NO$_2$, and NO concentrations. We also find that census block groups in downtown/metro LA, South-Central LA, and east LA County, consistently had the most hazardous air pollution mixtures in relation to prevalence of TLBW.

We also identified profiles of clustered neighborhood contextual factors to show that sub-populations previously shown to be vulnerable to TLBW are concentrated in the downtown/metro area of LA and South-Central LA County. These high risk contextual clusters partially overlapped spatially with the most hazardous air pollution mixtures. Such co-occurrence of clustered indicators of disadvantage and hazardous air pollution mixtures reinforced the validity in our approach of adjusting for such correlated factors in examining multiple pollutant health effects.

<table>
<thead>
<tr>
<th>Pollutants</th>
<th>Pollutant clusters</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>P1</td>
</tr>
<tr>
<td>NO$_2$–NO</td>
<td>0.86</td>
</tr>
<tr>
<td>NO$<em>2$–PM$</em>{2.5}$</td>
<td>0.49</td>
</tr>
<tr>
<td>NO–PM$_{2.5}$</td>
<td>0.45</td>
</tr>
<tr>
<td>PM$_{2.5}$ – NO/NO$_2$</td>
<td>0.16</td>
</tr>
</tbody>
</table>

*a Overall correlations: NO$_2$–NO = 0.76, NO$_2$–PM$_{2.5}$ = 0.25, NO–PM$_{2.5}$ = 0.26, PM$_{2.5}$–NO/NO$_2$ ratio = 0.19.

*b Correlation between PM$_{2.5}$ concentration and the ratio of NO concentration to NO$_2$ concentration.
Spatially clustered emissions related to residential proximity to emissions sources may explain why our multilevel analysis of pollutant profile TLBW risks reveals such distinct spatial patterning. Spatial clustering inducing factors may include localized circumstances, such as traffic volumes and congestion, the vehicle fleet (e.g. heavy-duty trucks), and higher exposures to specific primary PM components that result from proximity to PM emissions sources. In our study, the spatial patterning across pollutant profiles of maternal distance to major highways, between pollutant correlations, elevated concentrations of NO and NO₂, and elevated NO/NO₂ ratios suggest that clusters reflective of primary traffic emissions tended to impart the greatest risk of TLBW (i.e. pollutant clusters P9 and P10). For instance our spatial proximity data show that the highest risk pollutant clusters (P9 and P10) are characterized by the smallest median maternal residential distances to major roadways, compared to all other cluster-specific residential distances to highways. In terms of primary traffic emission above background levels, the residential distances for P9 and P10 are consistent with the literature that suggest primary traffic emissions decay to background levels between 115 m to 570 m (Karner et al., 2010). In addition, there was clear spatial clustering of higher risk pollutant profiles (P9 and P10) at interchanges where LA’s major interstate highways (I-5, I-110, I-710, and I-10) converge in central and south-central areas of LA County. This is suggestive of highly localized traffic and emission patterns that are germane to these major highways and their intersections.

Findings from our study are notably consistent with our earlier studies and other’s conducted in LA County that found variation in estimated effects on birth outcomes between traffic-related sources of air pollution (e.g. traffic-related versus natural background sources) and proximity to major roadways (Laurent et al., 2014; Ritz et al., 2007; Wilhelm et al., 2011a; Wilhelm and Ritz, 2005). When juxtaposed with other recent studies (Bell et al., 2011; Coker et al., 2015; Laurent et al., 2014; Pirani et al., 2015; Wilhelm et al., 2011a; Zanobetti et al., 2014), our findings carry the implication that TLBW risks related to spatial patterns in exposure combined with the physical and chemical properties of PM₂.₅ requires further investigation, and further suggests important spatially derived hypotheses. For instance, recent findings by our group showed that the exposure response relationship of PM₂.₅ on TLBW varied spatially across LA County (Coker et al., 2015). Spatially varying effects suggests greater than additive health impacts influenced by (1) the sources of localized emissions, (2) proximity to PM sources (Buonocore et al., 2009; Cho et al., 2009; Greco et al., 2007; Kuhn et al., 2005; Laurent et al., 2014; Wagner et al., 2012), and (3) the varied pollutant profiles associated with proximity to different emissions sources of PM₂.₅ (Laurent et al., 2014).

An important limitation of previous studies that attempted to find gradients in TLBW risk associated with various PM₂.₅ components is the inability to pinpoint major sources or components contributing to TLBW risks. Instead, nearly all sources and components imparted a risk of exposure and are correlated. For example, the inherent dependencies between PM₂.₅, PM₁₀, and various carbonaceous particulates (e.g. organic carbon, black carbon, and elemental carbon), or between PM₂.₅ and sulfates in the exposure model used by Laurent et al. (2014), made it impossible to parse out which fraction sizes, components within PM₂.₅- or combination thereof - are most likely to impart the greatest TLBW risk (Laurent et al., 2014). A single major source combined with certain spatially determined factors may produce a particular air pollution mixture that is more hazardous, yet multiple regression techniques struggle to distinguish between them.

The results from ours and a recent study by Pirani et al. (2015), show that Bayesian profile regression provides a tangible clustering procedure to develop profiles of exposure to multiple pollutants and simultaneously provide visualization tools. For instance, Pirani et al. (2015) studied variations in respiratory mortality across exposure profile clusters using a similar Bayesian profile regression. They found that days with high levels of secondary particulates (e.g. nitrates and sulfates) imparted the highest mortality risk in comparison to all other PM₂.₅ component exposure profiles. Thus, rather than multiple regression models with pollutants and sources being highly correlated (Hampel et al., 2015; Laurent et al., 2014), our clustering approach could be applied to develop PM-exposure profiles using data on PM₂.₅ components, sources, and size fractions. Furthermore, our spatially-based clustering approach enables identification and mapping of sub-regions that are characterized by the most hazardous PM-source components.

Rather than simply examining gradients in multipollutant health outcome risks devoid of spatial information, our study illustrates the importance in examining the spatial patterning of multipollutant health effects to help bring out the likely causes of apparent non-linear effects. For example, pollutant cluster P9 - reflective of primary traffic PM₂.₅ pollution - was not the only cluster with elevated concentrations for all

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**Fig. 3.** Boxplots of (a) NO/NO₂ ratios, (b) normalized NO, and (c) normalized NO₂ for each pollutant cluster. The red line indicates the overall average for each of these pollutant metrics.

**Fig. 4.** Distribution of residential distance to highways among mothers within each pollutant cluster. Redline indicates the overall median for all maternal residential distances (median = 867 m).
study pollutants. Cluster P9, however, displayed the highest probability for an estimated effect above zero and the second highest effect estimate size, despite other pollutant clusters displaying higher PM$_{2.5}$ or higher NO and NO$_2$ concentrations. The only other pollutant cluster with a larger estimated effect size was cluster P10, which was also characteristic of primary traffic pollution. Whereas pollutant cluster P13—reflective of mostly secondary traffic PM$_{2.5}$—has a PM$_{2.5}$ concentration 30% higher than P9 and P10. Despite this, the estimated effect size for cluster P13 is lower than P9 and P10. Our approach thus identifies patterns that help explain apparent non-linear effects, such as: (1) mapping of cluster effects that exhibited strong spatial patterns related to major roadways, (2) the variations in pollutant metrics such as correlations and pollutant ratios across clusters combined with the spatial patterning of these pollutant metrics, and (3) the influence of residential distance to major roadways. These spatial data provide strong evidence that primary traffic emissions uniquely impart the largest effect on TLBW.

4.2. Correlated pollutants and health effects research

Our approach to examining health effects of correlated exposures via exposure profile clusters offers several advantages compared to co-pollutant regression methods. Problems with collinearity within a multivariate regression include inflated variance in regression coefficients, unstable effect estimates, and causal inference challenges (Dormann et al., 2013; Lin, 2008; Schmidt and Muller, 1978). Several outdoor air pollution studies find that pollutants contribute to a health outcome in single pollutant models; however, mutual adjustment for correlated pollutants can result in no pollutant showing an association (Ebisu and Bell, 2012) or coefficients flipping sides i.e. opposite in direction

<table>
<thead>
<tr>
<th>Cluster</th>
<th>No. births*</th>
<th>No. TLBW</th>
<th>% TLBW (95% CI)</th>
<th>Cluster effect† (95% CI)</th>
<th>Probability effect &gt; 0‡</th>
</tr>
</thead>
<tbody>
<tr>
<td>P1</td>
<td>29394</td>
<td>431</td>
<td>1.47* (1.33, 1.61)</td>
<td>−0.122 (−0.254, 0.008)</td>
<td>0.033</td>
</tr>
<tr>
<td>P2</td>
<td>34263</td>
<td>589</td>
<td>1.72* (1.58, 1.86)</td>
<td>−0.108 (−0.232, −0.015)</td>
<td>0.042</td>
</tr>
<tr>
<td>P3</td>
<td>79199</td>
<td>1555</td>
<td>1.96 (1.87, 2.06)</td>
<td>−0.026 (−0.133, 0.081)</td>
<td>0.315</td>
</tr>
<tr>
<td>P4</td>
<td>168145</td>
<td>3305</td>
<td>1.97 (1.89, 2.03)</td>
<td>0.006 (−0.093, 0.106)</td>
<td>0.548</td>
</tr>
<tr>
<td>P5</td>
<td>66035</td>
<td>1595</td>
<td>2.42† (2.30, 2.53)</td>
<td>0.017 (−0.092, 0.126)</td>
<td>0.619</td>
</tr>
<tr>
<td>P6</td>
<td>58164</td>
<td>1067</td>
<td>1.83 (1.73, 1.95)</td>
<td>0.003 (−0.109, 0.115)</td>
<td>0.519</td>
</tr>
<tr>
<td>P7</td>
<td>256817</td>
<td>5612</td>
<td>2.19† (2.13, 2.24)</td>
<td>0.019 (−0.080, 0.117)</td>
<td>0.642</td>
</tr>
<tr>
<td>P8</td>
<td>7452</td>
<td>113</td>
<td>1.52 (1.25, 1.82)</td>
<td>−0.096 (−0.275, 0.079)</td>
<td>0.141</td>
</tr>
<tr>
<td>P9</td>
<td>77336</td>
<td>1824</td>
<td>2.36† (2.25, 2.47)</td>
<td>0.089 (−0.018, 0.195)</td>
<td>0.949</td>
</tr>
<tr>
<td>P10</td>
<td>3171</td>
<td>82</td>
<td>2.59 (2.06, 3.20)</td>
<td>0.141 (−0.060, 0.338)</td>
<td>0.916</td>
</tr>
<tr>
<td>P11</td>
<td>3491</td>
<td>74</td>
<td>2.12 (1.67, 2.65)</td>
<td>0.043 (−0.168, 0.252)</td>
<td>0.656</td>
</tr>
<tr>
<td>P12</td>
<td>3537</td>
<td>51</td>
<td>1.52 (1.13, 1.91)</td>
<td>−0.012 (−0.232, 0.205)</td>
<td>0.459</td>
</tr>
<tr>
<td>P13</td>
<td>17902</td>
<td>396</td>
<td>2.21 (2.00, 2.44)</td>
<td>0.055 (−0.087, 0.196)</td>
<td>0.774</td>
</tr>
<tr>
<td>Overall</td>
<td>804,726</td>
<td>16,694</td>
<td>2.07 (2.04, 2.11)</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

* Full term births (>259 days gestation).
† Percent prevalence of TLBW without model adjustment.
‡ Adjusted for maternal age, race, education, and parity, infant sex, gestation (days), gestation-squared, and contextual random effect clusters.
§ Random effect presented on the log-odds scale.
¶ Probabilities were calculated utilizing the "inla.pmarginal" function in INLA.
* TLBW prevalence is significantly (p-value < 0.01, two-sided) different from the overall proportion of TLBW (unadjusted).

Fig. 5. Spatial distribution of pollutant profile cluster random effect posterior probabilities.
from single pollutant models (Kelsall et al., 1997). Furthermore, whether mutual adjustment is necessary – i.e. whether multiple pollutants actually confound each other – cannot be determined in models with highly correlated pollutants. When we analyze our data using a co-pollutant model that includes all pollutants in a multivariate regression model, we find instability of specific pollutant coefficient estimates and increased standard errors (see Table S2 in the Supplemental materials). Also, while we find suggestive evidence for a statistically significant interaction between NO and NO₂ in a co-pollutant model (Table S1, Supplemental materials), the interpretation is challenging since both pollutants tend to co-vary and are linked through conversion into one-another by atmospheric chemistry involving ozone. Bayesian profile regression on the other hand harnesses the collinearity of air pollutants to find meaningful patterns of joint exposure that are relevant for determining different health risks across pollutant clusters.

Another important problem with multipollutant modeling is that correlations between pollutants can vary over space (Dionisio et al., 2014; Levy et al., 2013; Snowden et al., 2015). Spatial variability in pollutant correlations between and within urban communities is challenging because it can lead to exposure measurement error and further calls into question estimating co-pollutant effects reliably. It is also unclear whether results from studies in one particular region are generalizable to others that have different spatial patterns of pollutant correlations. Consistent with previous studies (Dionisio et al., 2014; Levy et al., 2013), we found substantial within county variability in pollutant correlations between our pollutant clusters (Figs. S6 and S7, Supplemental materials). To some extent our approach overcomes spatial variation in pollutant correlations because we characterized the heterogeneity in pollutant relationships across space, and further relate these exposure profiles to a health outcome. Hence, employing our approach in environmental health studies may better inform policies designed to protect public health since policies can be tailored towards pollution mixtures relevant to a specific area.

### 4.3. Contextual neighborhood effects on TLBW

An important aspect to our pollutant clustering approach lies in simultaneously adjusting for clustered neighborhood indicators of disadvantage (i.e. income and race) and older housing. While other contextual factors related to TLBW could have been included in our clustering procedure, it is clear that our clustering variables are highly correlated with other adverse contextual factors (i.e. education, housing values, low social support, neighborhood greenness, violent crimes, etc.) in southern California, and thus likely account for these other contextual factors (Boggess and Hipp, 2010; Conway et al., 2010; English et al., 2003; Ghosh et al., 2010). Furthermore, since our contextual variables correlate with one another and correlate with pollutants, it was important to separate out the contextual area-level effects from the

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### Table 4

Prevalence of TLBW for contextual clusters and model results for contextual exposure profile cluster random effects (N = 804,726).

<table>
<thead>
<tr>
<th>Cluster</th>
<th>No. birthsa</th>
<th>No. TLBWb</th>
<th>% TLBW CI</th>
<th>Cluster effectc,d (95% CI)</th>
<th>Probability effect &gt; 0e</th>
</tr>
</thead>
<tbody>
<tr>
<td>C1</td>
<td>64018</td>
<td>965</td>
<td>1.51 (1.42, 1.60)</td>
<td>−0.09 (−0.19, 0.02)</td>
<td>0.05</td>
</tr>
<tr>
<td>C2</td>
<td>95559</td>
<td>1869</td>
<td>1.96 (1.87, 2.05)</td>
<td>−0.03 (−0.12, 0.07)</td>
<td>0.30</td>
</tr>
<tr>
<td>C3</td>
<td>76163</td>
<td>1218</td>
<td>1.60 (1.51, 1.69)</td>
<td>−0.08 (−0.18, 0.02)</td>
<td>0.07</td>
</tr>
<tr>
<td>C4</td>
<td>39105</td>
<td>787</td>
<td>2.01 (1.88, 2.16)</td>
<td>−0.03 (−0.14, 0.08)</td>
<td>0.31</td>
</tr>
<tr>
<td>C5</td>
<td>38084</td>
<td>809</td>
<td>2.12 (1.98, 2.27)</td>
<td>−0.02 (−0.11, 0.11)</td>
<td>0.49</td>
</tr>
<tr>
<td>C6</td>
<td>112919</td>
<td>2970</td>
<td>2.63 (2.54, 2.73)</td>
<td>0.12 (0.03, 0.22)</td>
<td>0.99</td>
</tr>
<tr>
<td>C7</td>
<td>39738</td>
<td>912</td>
<td>2.30 (2.15, 2.45)</td>
<td>0.03 (−0.08, 0.14)</td>
<td>0.71</td>
</tr>
<tr>
<td>C8</td>
<td>165328</td>
<td>3384</td>
<td>2.05 (1.98, 2.12)</td>
<td>0.03 (−0.07, 0.12)</td>
<td>0.71</td>
</tr>
<tr>
<td>C9</td>
<td>101455</td>
<td>2040</td>
<td>2.01 (1.93, 2.10)</td>
<td>0.02 (−0.07, 0.14)</td>
<td>0.67</td>
</tr>
<tr>
<td>C10</td>
<td>2624</td>
<td>42</td>
<td>1.60 (1.18, 2.16)</td>
<td>−0.08 (0.30, 0.13)</td>
<td>0.22</td>
</tr>
<tr>
<td>C11</td>
<td>43403</td>
<td>975</td>
<td>2.25 (2.11, 2.39)</td>
<td>0.07 (−0.04, 0.17)</td>
<td>0.90</td>
</tr>
<tr>
<td>C12</td>
<td>963</td>
<td>18</td>
<td>1.87 (1.18, 2.95)</td>
<td>0.07 (−0.31, 0.18)</td>
<td>0.30</td>
</tr>
<tr>
<td>C13</td>
<td>18784</td>
<td>538</td>
<td>2.86 (2.64, 3.11)</td>
<td>0.03 (−0.09, 0.15)</td>
<td>0.68</td>
</tr>
<tr>
<td>C14</td>
<td>6583</td>
<td>167</td>
<td>2.54 (2.18, 2.95)</td>
<td>0.07 (−0.09, 0.23)</td>
<td>0.82</td>
</tr>
</tbody>
</table>

---

a Full term births (≥259 days gestation).

b Percent prevalence of TLBW without model adjustment.

c Adjusted for maternal age, race, education, and parity, infant sex, gestation (days), gestation-squared, and pollutant random effect clusters.

d Random effect presented on the log-odds scale.

e Probabilities were calculated utilizing the “inla.pmarginal” function in INLA.

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### Fig. 6

Contextual profile clusters: cluster size (number of census block groups) and posterior distributions for median income, percent older homes, percent black, percent white, and percent Hispanic (Nclusters = 14). Black lines indicate the overall average exposure for a given covariate.
pollutant profile cluster effects. Despite the spatial similarity between the two different types of clusters in our study, the high risk contextual clusters display a spatial pattern that is distinct from the high risk pollutant clusters, suggesting that these two separate exposure profiles measured different spatial patterns of risk related to their respective variables.

4.4. Study limitations

We lacked data on speciation and tracers for specific sources of PM2.5, which limits our ability to attribute a particular air pollution source to effects on TLBW. Despite this limitation, the spatial patterning of our results, supplemented with metrics such as cluster-specific NO/NO2 ratios and between pollutant correlations, offers strong evidence implicating primary traffic pollution. Another limitation lies in the lack of fine-scale spatial data for other air toxics (i.e. benzene, ozone and carbon monoxide). Since we lack data on other air toxics that correlate with the pollutants considered in our study (Fujita et al., 2011; Ghosh et al., 2012; Laurent et al., 2014; Protano et al., 2012; Salam et al., 2005; Wilhelm et al., 2011a), we cannot say whether and how these other pollutants may contribute to the observed spatial patterning of TLBW risks. Other limitations include a lack of information on indoor air pollution exposure and time-activity patterns that may influence air pollution exposures, such as information about whether the women worked outside the home (Ritz et al., 2007) or commuted daily (Zuurbier et al., 2010); all of which can contribute further to exposure misclassification. However, recent findings suggest that maternal outdoor air pollution estimates at the home address are unlikely affected by a lack of time-activity patterns during pregnancy (Ouidir et al., 2015). Finally, while our approach offers several important advantages over previous air pollution profile studies, our Bayesian approach is currently limited in regards to handling a dataset with high dimensionality, which can only be overcome via future gains in computational efficiencies.

4.5. Study strengths

The primary strength of our study is that we were able to examine exposures to multiple correlated air pollutants and a health outcome, mitigating some of the typical problems encountered with correlated exposures. Our study also had a large sample size and used population-wide data for exposure, thus avoiding selection of a study.
population based on proximity to major sources of air pollution or proximity to central site monitors (Dionisio et al., 2014; Kumar, 2012). Another strength is adjustment for individual-level covariates and contextual factors associated with TLBW, and adjustment for spatial residual confounding at the census tract-level. Another important strength is that we were able to fit a multilevel/hierarchical random effects model for the clustered pollution profiles and contextual profiles enabling us to look at multiple profile-specific risks, thus avoiding some of the issues related to multiple testing of myriad exposure/SES effects on health (Gelman and Hill, 2006).

5. Conclusion

Our Bayesian clustering procedures allowed us to go beyond simple one-at-a-time analyses usually employed to examine marginal effects of individual pollutants on health. Further, this spatially distributed mixture approach provides information on the spatial distribution of exposure/SES profiles that pertain to the levels of various pollutants and SES factors. Policy analysts can use this information to determine which exposure/SES profiles dominate a particular sub-region of LA County, as a starting point for regulatory considerations. In our analyses, we found that neighborhood-level PM$_{2.5}$, NO$_x$ and NO concentrations were correlated with census block group-level contextual factors throughout LA County; and the nature of these relationships was quite complex and highly spatially varying across the County. Moreover, the pollution profile clusters showed a strong spatial contrast with respect to exposure-related TLBW risks. LA County’s urban core, south-central urban region, and parts of the eastern-most region of the county exhibited the largest exposures for PM$_{2.5}$, NO$_x$ and NO, which decreased with distance from major highways. Moreover, the highest concentration pollution profile clusters imparted the greatest TLBW risks, especially those closest to major highways, which suggests near roadway emissions are more important in terms of risk of adverse birth outcomes related to these air pollution profiles.

Acknowledgments

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Appendix A. Supplementary data

Supplementary data for this article can be found online at http://dx.doi.org/10.1016/j.envint.2016.02.011.

References


