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Metropolitan isolation segregation and Black-White disparities in very preterm birth: A test of mediating pathways and variance explained

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Abstract

Residential isolation segregation (a measure of residential inter-racial exposure) has been associated with rates of preterm birth (<37 weeks gestation) experienced by black women. Epidemiologic differences between very preterm (<32 weeks gestation) and moderately preterm births (32–36 weeks) raise questions about whether this association is similar across gestational ages, and through what pathways it might be mediated. Hierarchical Bayesian models were fit to answer three questions: is the isolation-prematurity association similar for very and moderately preterm birth; is this association mediated by maternal chronic disease, socioeconomic status, or metropolitan area crime and poverty rates; and how much of the geographic variation in black-white very preterm birth disparities is explained by isolation segregation? Singleton births to black and white women in 231 U.S. metropolitan statistical areas in 2000–2002 were analyzed and isolation segregation was calculated for each. We found that among black women, isolation is associated with very preterm birth and moderately preterm birth. The association may be partially mediated by individual level socioeconomic characteristics and metropolitan level violent crime rates. There is no association between segregation and prematurity among white women. Isolation segregation explains 28% of the geographic variation in black-white very preterm birth disparities. Our findings highlight the importance of isolation segregation for the high-burden outcome of very preterm birth, but unexplained excess risk for prematurity among black women is substantial.

Keywords

Preterm birth; Racial disparity; Residential segregation; Bayesian modeling; Social determinants; USA

Introduction

Preterm birth (<37 weeks gestation) is the leading cause of infant mortality in the US (Callaghan, MacDorman, Rasmussen, Qin, & Lackritz, 2006). However a small subset—approximately 2% of all US births (Martin, Hamilton, Sutton, Ventura, Menacker et al., 2009)— who are born less than 32 weeks (*very preterm birth*) account for 95% of preterm-

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associated infant mortality. Very preterm birth is also a leading risk factor for pediatric morbidities such as cerebral palsy and mental retardation (Yeargin-Allsopp, Van Naarden Braun, Doernberg, Benedict, Kirby et al., 2008), and substantial economic costs for neonatal care, rehabilitative and educational services through a child's life (Behrman & Butler, 2007). The stubborn persistence of racial disparities in prematurity further contributes to the urgency of this public health problem. Black women in the US have 43% greater risk than white women for delivering moderately preterm at 32–36 weeks gestation, and 2–3 times the risk of delivering at less than 32 weeks gestation (Martin, Kung, Mathews, Hoyert, Strobino et al., 2008).

It is common in epidemiologic studies to combine all births less than 37 weeks into a single group, but these births are not only heterogeneous with regards to the magnitude of the racial disparity; there may also be etiological differences related to gestational age. Infection and inflammation are leading causes of very preterm birth, particularly for black women, but less important in near-term births (Goldenberg, Hauth, & Andrews, 2000). On the other hand, assisted reproductive technology, increases in medical intervention in pregnancy, and societal changes in maternal age at conception may contribute to the recent increase in moderately preterm births (Branum & Schoendorf, 2002; Raju, 2006).

Further efforts to disentangle the causes of these profound racial disparities and minimize the burdens that result, requires greater attention to the heterogeneity within common epidemiologic definitions of disease.

Pathways to prematurity

Preterm birth is a complex health problem with social, environmental, behavioral and genetic determinants of individuals' risk (Kramer & Hogue, 2009a). While numerous risk factors for preterm birth have been reported, their failure to explain a significant portion of the racial disparity has lead investigators to look further upstream in the causal chain to place-based contextual effects and structural inequities in access to resources. For instance, controlling for individual-level risk factors, neighborhood violent crime and poverty rates have been associated with increased risk of preterm birth (Kaufman, Dole, Savitz, & Herring, 2003; Messer, Kaufman, Dole, Savitz, & Laraia, 2006; O'Campo, Burke, Culhane, Elo, Eyster et al., 2008).

The association between preterm birth and neighborhood context could be mediated by a combination of material and psychosocial pathways related to access to safe living conditions, maternal perceptions of neighborhood quality, and exposure to discrimination or stressful life events (Collins, David, Handler, Wall, & Andes, 2004; Collins, David, Symons, Handler, Wall et al., 1998). These acute or chronic stressors may interact with maternal neuroendocrine and immune function resulting in preterm birth, or could be mediated by the prevalence of poorly controlled chronic hypertension, or individual behavioral responses (e.g. smoking) to stressful environments, each of which could differentially pattern risk (Kramer & Hogue, 2009a).

While local residential neighborhoods have been useful constructs for conceptualizing contextual rather than individual determinants of health disparities, some exposures may operate differently at different geographic scales. Health-relevant policies and social context may be patterned at the scale of counties, urbanized areas or states (Bird, 1995; Centers for Disease Control and Prevention, 2002; Sims, Sims, & Bruce, 2007). The risk of very preterm birth varies substantially more among metropolitan statistical areas (MSA's) for black women than it does for white women (Kramer & Hogue, 2008). The neighborhoods within these MSA's may still be the proximally relevant exposure, but the propensity for

some MSA's to have more unhealthy neighborhoods than others suggests additional patterning forces.

Residential segregation and prematurity

The spatial segregation of blacks and whites across residential neighborhoods may indicate the extent of regional structural inequality and institutionalized racism which could explain MSA-level disparities in infant mortality and low birthweight (Ellen, 2000; Laveist, 1989; Polednak, 1996). More recently, segregation has been associated with other indicators of perinatal health including preterm birth (Bell, Zimmerman, Almgren, Mayer, & Huebner, 2006; Osypuk & Acevedo-Garcia, 2008), teen pregnancy (Sucoff & Upchurch, 1998), and smoking during pregnancy (Bell, Zimmerman, Mayer, Almgren, & Huebner, 2007). While studies of small area (e.g. neighborhood) segregation find poor health outcomes for both black and white women as segregation increases (Grady & Ramirez, 2008; Mason, Messer, Laraia, & Mendola, 2009), studies focusing on MSA's suggest a cross-level interaction where segregation is associated with increased risk for black but not white women (Ellen, 2000; Kramer & Hogue, 2008). This is likely due to the fact that very few white women live in predominantly black neighborhoods in highly segregated MSA's.

Residential segregation is often distinguished along several 'dimensions' describing unique spatial patterns of residential settlement in urban and suburban areas (Massey & Denton, 1988). Three dimensions commonly employed in research on segregation and health outcomes are isolation, evenness, and clustering. Isolation is the probability of interaction between blacks and whites within neighborhoods; evenness describes the racial distribution within neighborhoods to the overall MSA composition; clustering is the aggregation of racially homogenous neighborhoods in sub-regions of a metropolitan area.

Although correlated, these patterns may capture distinct aspects of health-relevant exposures (Kramer & Hogue, 2009b). For example Bell, et al (2006) report a reduced risk for preterm birth when clustering of predominantly black neighborhoods is present, but a elevated risk of prematurity in the presence of isolation segregation. The authors suggest that conditional on the degree of isolation, clustering enhances social support and networks for black women and families, which may reduce risk by buffering against psychosocial stressors, and facilitating health protective behaviors. Isolation on the other hand may capture economic disenfranchisement and poverty concentration, and is associated with elevated violent crime rates, diminished access to healthy food options, and reduced access to preventive healthcare (Collins, 1999; Morland & Filomena, 2007; Shihadeh & Flynn, 1996). These factors may result in racial differences in risk factors for preterm birth including exposure to psychosocial stress, and prevalence of chronic diseases such as diabetes and hypertension.

Study Questions

Thus three observations motivate this study. First, prematurity may best be understood as a heterogeneous outcome, resulting from complex health processes, with at least one domain of difference being *very* preterm versus *moderately* preterm births. Second, the magnitude of black-white racial disparities is not constant but varies geographically and at different geographic scales. This variation in relative risk may offer clues as to some causes of the disparity. Finally, residential segregation has been hypothesized to be a distal determinant of racial disparities in prematurity, but this relationship has never been looked at for very preterm birth, and is incompletely understood in terms of mediating pathways.

From these three observations we pose three questions.

1. Is isolation segregation associated similarly with very versus moderate preterm birth?

2. Which individual or metropolitan characteristics mediate the association between isolation and preterm birth?
3. Does isolation segregation explain the geographic variation in black-white disparities of very preterm birth?

Methods

Data sources

Individual level variables—At the individual level, each singleton live births born in 2000–2002 to non-Hispanic white or black mothers living in US metropolitan statistical areas was abstracted from National Center for Health Statistics natality files. Individuals are geo-located within MSA's but not within specific neighborhoods. Births were categorized as very preterm (from 20 to less than 32 weeks), moderately preterm (from 32 to less than 37 weeks), or term (37 to 44 weeks).

Variables including maternal education, marital status, smoking, and chronic disease status were also abstracted from birth records. These variables are hypothesized to be mediators of a segregation effect on prematurity. Residential segregation may influence family social structure, area school quality and adult educational attainment (Card & Rothstein, 2007; Howell-Moroney, 2005); maternal education and marital status are each associated with preterm birth, and thus to the degree that segregation influences metropolitan-level patterns they could mediate a health effect. Pre-conceptional chronic diseases such as hypertension and diabetes are also associated with preterm birth (Ehrenthal, Jurkowitz, Hoffman, Kroelinger, & Weintraub, 2007), and there is some evidence that prevalent chronic disease mediates some of the segregation association with low birthweight for black women (Grady & Ramirez, 2008). Smoking during pregnancy is also a risk factor for preterm birth, and the rate of smoking may vary by degree of segregation (Bell et al., 2007). Because smoking is not measured on California birth records, this variable is coded with a level to indicate missing values to retain California births in all models including smoking.

Other variables captured for individual births include maternal age, parity, and history of a prior preterm birth.

Metropolitan level variables—Metropolitan statistical areas were chosen as the contextual unit of analysis. MSA's represent contiguous counties surrounding a core city which are deemed by the federal government to be economically and socially integrated (OMB, 2000). The geographic scale of MSA's are particularly well suited for this study because residential segregation is conceptualized as a process of sorting individuals into living environments on the basis of race and class. This process happens across regional residential housing markets; therefore simultaneously recognizing the housing choices of economically and socially linked urban and suburban communities is beneficial. We analyzed 231 MSA's which had a population of at least 100,000 and had a non-Hispanic black population of at least 5,000 in the 2000 Census.

Metropolitan population size (categorized as <500,000, 500,000–1 million, or >1 million), and Census region (Northeast, Southeast, Midwest, West) were obtained from the 2000 Census. Because there may be variation in both segregation and very preterm birth risk across these variables, we control for them in all models to better describe the independent association of segregation.

Potential mediators between isolation and prematurity at the metropolitan level were chosen based on evidence from neighborhood-level contextual determinants reviewed above, and

include violent crime and poverty rates. The mechanisms for an effect of these variables on pregnancy health could result from chronic exposure to psychosocial stressors and subsequent 'weathering' or premature aging of maternal neuroendocrine and vascular function (Geronimus, 1996). Alternatively effects could be material in nature, related to access to health promoting resources including health and dental care (Haas, Phillips, Sonneborn, McCulloch, Baker et al., 2004; Kushel, Gupta, Gee, & Haas, 2006).

The murder rate per 100,000 persons in each metropolitan area was obtained from federal statistics for 2000 (Federal Bureau of Investigation, 2000). The black poverty rate as well as the ratio of black to white poverty rates in each MSA were calculated from Census 2000 data to represent indicators of area-based racial inequity (US Census Bureau, 2000). We used measures of the exposure of poor children to high poverty neighborhoods as calculated by Acevedo-Garcia, et al (2007) to approximate spatial poverty concentration. This variable is the proportion in each MSA of black children under 18 years of age whose families are below the poverty line *and* live in census tracts with median household incomes less than 80% of the median household income for the MSA as a whole in 2000.

Measuring residential segregation—The primary segregation dimension of interest is isolation, although unevenness/clustering was also measured (Massey & Denton, 1988). We use explicitly spatial adaptations of common census-tract derived segregation indices to measure these dimensions (Reardon & O'Sullivan, 2004). The strength of these spatial measures is that they minimize measurement bias introduced by reliance on the arbitrary size and shapes of census tracts by applying a consistent definition of residential neighborhoods within MSAs in the calculation segregation indices (Kramer, Cooper, Drews-Botsch, Waller, & Hogue, 2010). The spatial segregation indices also allow a flexible definition the size or scale of neighborhoods, recognizing that the granularity of segregation may be an attribute of the segregation pattern in and of itself (Lee, Reardon, Firebaugh, Farrell, Matthews et al., 2008). The spatial isolation index is calculated using the formula:

$$x \tilde{P} *_x = \sum_{p \in R} \frac{\tau_p}{T} \tilde{\pi}_p$$

Eq. 1 Spatial isolation index

τ_p is the total population density for each point p in region R (the MSA), and T is the total MSA population. $\tilde{\pi}_p$ is the proportion black in the spatial area or residential environment of point p . We defined the residential environment as a 500-meter radius circle around each point p in the MSA. This definition of neighborhood is based on exploratory analysis of best model fit at different spatial scales (Kramer et al., 2010).

Reardon & O'Sullivan argue that the clustering dimension of segregation is simply an uneven distribution of predominantly black neighborhoods, and thus an index of evenness such as the dissimilarity index would measure clustering if the spatial scale of the local area were sufficiently broad. We operationalized clustering (or sub-regional unevenness) using a spatial adaptation of the dissimilarity index:

$$\tilde{D} = \sum_{p \in R} \frac{\tau_p |\tilde{\pi}_p - \pi|}{2T\pi_p(1 - \pi)}$$

Eq. 2. Spatial dissimilarity index

Again, τ_p is the total population density at point p , while π and π_p are the proportion black in the MSA overall and at point p respectively. $\tilde{\pi}_p$ denotes the proportion black in the spatial

area around point p which for clustering is set as a 4000-meter radius circle. More detail on the calculation method and correlation between isolation and clustering is provided in the electronic technical appendix available with the online version of the paper [SUPPLEMENTARY FILE]. The resulting indices each range from 0 to 1; for isolation 0 represents highest inter-racial exposure and 1 is highest isolation. For the dissimilarity index 0 suggests least clustering or most even distribution across large sub-regions, while 1 suggests a high degree of clustering. Each segregation index was standardized to a mean of 0, standard deviation of 1, so that model interpretations are made in terms of a 1-standard deviation change in the relevant index.

Analysis

All analyses were conducted using hierarchical Bayesian logistic regression models (Gelman & Hill, 2007). The setup for each model follows this template:

$$\begin{aligned} P(y_i=1) &= \text{logit}^{-1}(\alpha_j + \beta X_i) \\ \alpha_j &\sim N(\gamma_0 + U_j \gamma_1, \sigma_\alpha^2) \\ \beta &\sim N(0, 10000) \\ \gamma_i &\sim N(0, 10000) \\ \sigma_\alpha &\sim \text{Unif}(0, 100) \end{aligned}$$

Eq. 3. Hierarchical Bayesian logistic model

In the first level y_i is the binary pregnancy outcome for the i^{th} woman, α_j is a random intercept for the j^{th} MSA, β is a vector of parameters for individual variables, and \mathbf{X} is a matrix of individual level covariates. Each parameter is assigned a prior probability distribution. Relatively uninformative priors are assigned to the β -parameters, while the α -intercept has an informative prior, in the form of the second level of the model. The random intercepts (alphas) are assumed to come from a normal distribution with a variance of σ_α^2 . The mean of the distribution is the sum of a global intercept, γ_0 , and the vector of γ -parameters corresponding to the MSA-level covariates in matrix \mathbf{U} , including segregation. All models reported control for census region and metropolitan size as MSA-level confounders and except for crude models adjust for maternal age, parity and history of prior preterm birth at the individual level.

For question one, concerning the association of segregation with very and moderately preterm births, separate models were fit for white and black women, and for very preterm and moderately preterm birth as the dependent variable.

Question two concerns possible mediation of a segregation-prematurity association by hypothesized mechanisms. Models of the crude association of individual covariates and segregation indices with preterm birth are adjusted only for MSA region and size. Baseline models controlling for possible individual and metropolitan level confounders are denoted M1. Mediation was evaluated by comparing the magnitude of the segregation-preterm birth association with and without the candidate mediator(s), with meaningful attenuation of the association suggesting the variable is a mediator in the pathway, or alternately a non-causal confounder of the crude association. Potential mediation by socioeconomic status is evaluated by adding variables for maternal education and marital status (model M2); likewise maternal health status is considered as a mediator by entering chronic hypertension, diabetes and smoking variables (model M3). Model M4 includes all of the above covariates. The primary segregation pattern of interest is spatial isolation; to consider whether spatial clustering segregation contributes to the model, a final model with both segregation indices is fit (M5).

Question three, concerning the geographic variation in the racial disparity of preterm birth attributable to segregation includes births to white and black mothers. The model structure is similar to the previous questions with the addition of a binary variable for race, and an associated random slope for this variable:

$$P(y_i=1) = \text{logit}^{-1}(\alpha_j + \delta_j * \text{Black}_i + \beta X_i)$$

$$\delta_j \sim N(\gamma_{\text{race}0} + U_j \gamma_{\text{race}1}, \sigma_{\text{race}}^2)$$

Eq. 4. Model for black-white disparity

The random intercept, α_j , and the priors remain the same as the previously described model (equation 3). δ_j is the MSA-specific relative black-white disparity accounting for individual and area covariates. $\gamma_{\text{race}0}$ is then the average excess risk across all MSA's for black as compared with white women, and $\gamma_{\text{race}1}$ is the vector of second-level parameters corresponding to the matrix, \mathbf{U} , of MSA-level covariates including isolation segregation. σ_{race}^2 is the variation in the disparity across MSA's.

Bayesian models were fit with WinBUGS 1.4 (Lunn, Thomas, Best, & Spiegelhalter, 2000) using R 2.7 (R Development Core Team, 2008) and the **R2WinBUGS** package (Sturtz, Ligges, & Gelman, 2005). All models were run with three chains, each for 50,000 iterations with the first half discarded; convergence was evaluated by visual inspection of the trace plots of the posterior parameter estimates from each chain, as well as an R-hat statistic of 1.1 or lower for each parameter (Gelman & Hill, 2007). Relative improvements in model fit were assessed using the deviance information criterion (DIC) (Spiegelhalter, Best, Carlin, & van der Linde, 2002).

Results

Of 6,180,544 eligible births during the study period, 23.5% were born to black mothers (Table 1). In this sample black women experienced three times the risk as white women for very preterm birth, and 60% greater risk for moderately preterm birth. Racial differences in the importance of individual risk factors result in variation in the racial disparity across covariates. For instance, higher maternal education and being married are more strongly protective for white women than for black women, so that the relative racial disparity is smaller among women without a high school degree, or among unmarried women.

For both black and white women, risk for preterm birth varied regionally, with the Western metropolitan areas having substantially lower risk than other areas for black women; for white women risk is higher in the Northeast and Midwest but lower in the West and Southeast.

Across the 231 metropolitan areas analyzed, isolation segregation ranged from 0.06 (Salt Lake City, Utah) to 0.86 (Gary, Indiana), with a median value of 0.51 and an interquartile range of 0.36–0.64. The measure of spatial clustering ranged from 0.22 (Lawton, Oklahoma) to 0.79 (also in Gary, Indian) with a median value of 0.51 and an interquartile range of 0.43–0.59. The Spearman rank correlation of MSA isolation and clustering was 0.47 ($p < .001$). After standardizing the indices, MSA's ranged from approximately -2 to +2 standard deviations for each index.

Table 2 reports odds ratios for very preterm birth among black women for models with isolation, individual covariates, and control for region and population size. In the crude model, each standard deviation increase in isolation is associated with an 11% increased odds of very preterm birth for black women (95% CI 1.08, 1.14). Models M1–M4 consider this association under different specifications of covariates. History of prior preterm birth,

chronic hypertension, or tobacco use is each an important predictor of very preterm birth risk; however the independent association of isolation with very preterm birth remains relatively unchanged with adjustment for such factors. A modest reduction in the association is seen in model M2 controlling for socioeconomic characteristics (OR 1.10, 95% CI 1.07, 1.14) compared with the baseline adjusted M1 model (OR 1.12, 95% CI 1.08, 1.15).

The independent association of clustering segregation (measured with spatial dissimilarity index) with very preterm birth in black women seen in the crude model results (OR 1.05, 95% CI 1.02–1.08) was similar across models M1 to M4 (data not shown). However in model M5, with both isolation and clustering segregation, the isolation-very preterm birth association strengthened (OR 1.15, 95% CI 1.10, 1.19), while the independent association of clustering appears to be modestly protective (OR 0.95, 95% CI 0.92, 0.99). Put another way, among MSA's with average clustering (e.g. standardized index=0), the odds ratio for very preterm birth in black women living in the MSA's with 1 standard deviation (SD) above the mean isolation level to those living in MSA's 1 SD below the mean isolation (2 SD change) was 1.32 (95% CI 1.21–1.42). This 2-SD change represents the range of observed isolation values conditional on clustering (online electronic technical appendix).

There was no association between clustering or isolation segregation and very preterm birth in any of the preceding models for white women (data not shown). Figure 1 plots the observed and model predicted risks using model M5 for black and white women. As MSA isolation increases the average very preterm birth risk for black women increases, while for white women there is no similar association.

Table 3 reports parallel model results with moderately preterm birth as the outcome. The point estimate for the association of isolation with moderately preterm birth is half as strong as it is with very preterm birth. While a one standard deviation increase in isolation conditional on clustering and all individual covariates increased very preterm birth odds 15% among black women, it increased moderately preterm birth odds only 8%. There was no association between segregation and moderately preterm birth in white women (data not shown).

Table 4 builds on model M5, by further considering metropolitan characteristics which could mediate the association with segregation. Statistical control for MSA murder rate modestly attenuates the isolation-very preterm birth association from 1.15 to 1.12 (95% CI 1.07, 1.17). There is also a substantial reduction in the DIC (suggesting improved model fit) in the models including murder rates or poverty concentration as compared to model M5.

In Table 5, models with both black and white women assess the change in the adjusted racial disparity for very preterm birth under various model specifications. Change in the disparity is calculated as a percentage excess risk explained (Lynch, Kaplan, Cohen, Tuomilehto, & Salonen, 1996): $(OR_{\text{crude}} - OR_{\text{adjusted}}) / (OR_{\text{crude}} - 1)$.

Inclusion of all individual covariates reduced the crude black-white odds ratio by approximately 25%. The addition of isolation segregation reduced the disparity an additional 8%, and there was a concomitant 28% decrease in the inter-MSA variation in the disparity (σ^2_{race}) from a variance of 0.107 to 0.077 with control for isolation. Addition of clustering added little to the model in terms of model fit or explanation of the disparity.

Valid measurement of gestational age is notoriously hard to capture even in prospective clinical studies and differential misclassification by race and class are particular concerns when using birth certificate data. Very low birthweight (VLBW) is highly correlated with very preterm birth and more reliably measured. To assess sensitivity of our findings to misclassification of gestational age, models were fit with VLBW as an alternate outcome.

Parameter magnitude and direction were similar in these models, suggesting misclassification of gestational age is not a significant source of bias.

To assess the sensitivity of our findings to the choice of spatial segregation measures, models were repeated using traditional census-tract-derived isolation, dissimilarity, and spatial proximity indices (Massey & Denton, 1988). Odds ratios for the association of these indices with preterm birth were similar in direction, but smaller in magnitude than with spatial indices. The difference in magnitude of association may be due to our use of a highly granular and consistent neighborhood definition as compared with the coarser and more variably sized pattern identified with census tracts (Kramer et al., 2010).

In models which included segregation categorized into quintiles, there was no evidence of non-linearities in the segregation-preterm birth association.

Discussion

The black-white racial disparity in preterm birth in the US is a stubborn problem which defies simple explanations. Residential segregation has been proposed as a fundamental cause of racial disparities in health because of the manner in which segregation may constrain some individuals' economic attainment, health, and welfare (Williams & Collins, 2001). Consistent with prior work (Bell et al., 2006; Osypuk & Acevedo-Garcia, 2008) this study finds evidence that for black women, independent of measured individual and area level risk factors, living in a metropolitan area characterized by high isolation segregation significantly increases risk for preterm birth. We further demonstrate that the association is nearly twice as strong for the outcome of very preterm birth as compared to near-term births, an important finding because of the substantial public health burden and increased racial gap for very preterm birth risk.

The different associations by gestational age may also hint at mechanisms by which segregation affects individual health. For example bacterial vaginosis is much more strongly associated with very preterm birth as compared to moderately preterm birth (Goldenberg, Iams, Mercer, Meis, Moawad et al., 1998). The higher prevalence of bacterial vaginosis among black women may be due to experiences of stress, discrimination or poor social support (Culhane, Rauh, & Goldenberg, 2006; Paul, Boutain, Manhart, & Hitti, 2008). Women living in neighborhoods characterized by social stressors such as high violent crime rates and economic disadvantage have both increased overall risk for poor pregnancy outcomes, as well as accelerated age-associated risk for poor outcomes consistent with Geronimus' weathering hypothesis (Cerdeira, Buka, & Rich-Edwards, 2008; Geronimus, 1996; Masi, Hawkey, Piotrowski, & Pickett, 2007). However living in MSA's characterized by high racial isolation means different things for white and black women. Poor whites tend to live in mixed-income neighborhoods, while blacks—and particularly poor blacks—living in highly segregated cities tend to live in neighborhoods with high poverty rates and lower economic opportunity (Osypuk, Galea, McArdle, & Acevedo-Garcia, 2009). High MSA isolation segregation may therefore be a risk marker of racial differences in exposures to chronic stress, lack of social support, and discrimination thereby disproportionately affecting black women's immune status and infection prevalence (Hogue & Bremner, 2005). There was a modest attenuation of the segregation-very preterm birth association with inclusion of MSA murder rate, a finding consistent with prior research (Masi et al., 2007; Messer et al., 2006). Violent crime could be a direct source of chronic stress, or it could be a marker of another process such as poor social support, low economic opportunity, or infrastructure decay.

We found, as have other investigators (Bell et al., 2006), that isolation and clustering segregation patterns have different independent associations with prematurity. Clustering and isolation are modestly correlated variables and each separately is associated with increased risk for prematurity in black women. However in models considering the joint relationship it appears that the isolation construct best captures the negative effect of segregation, while clustering is conditionally protective (very preterm birth) to null (moderately preterm birth). MSA's which are divergent on these two dimensions tend to be smaller in size and have smaller black populations (electronic technical appendix and Kramer et al., 2010). Possible explanations for a different association of clustering with prematurity conditional on isolation include buffering aspects of social networks and black political empowerment (Laveist, 1993).

Our findings can also be viewed in the context of research on neighborhood-level ethnic density and pregnancy outcomes. One study in Chicago neighborhoods reported lower risk for low birthweight among black women living in predominantly black neighborhoods after controlling for area deprivation and individual socioeconomic status (Roberts, 1997), while studies in New York City and North Carolina found increased risk for low birth weight and preterm birth for black women in predominantly black neighborhoods (Grady & McLafferty, 2007; Mason et al., 2009). Ethnic enclave effects may also differ by maternal nativity (foreign born versus US born) (Grady & McLafferty, 2007), and by ethnicity (Hispanic versus non-Hispanic) (Osypuk, Bates, & Acevedo-Garcia, 2010). Such conflicting or complex associations may be attributable to regionally specific forces which affect individual selection into neighborhoods (Oakes, 2004) or makeup the broader regional social context. In a study which simultaneously measured neighborhood racial composition and city-level segregation, it was in fact city-level segregation which was found to have the independent (and deleterious) association with pregnancy health of black women, with no remaining significant association of neighborhood racial composition (Reichman, Teitler, & Hamilton, 2009).

Residential segregation has a statistically significant association with excess risk of preterm birth in black women, yet it independently accounts for only 8% of the black-white racial disparity, and in fact adjustment for all measured covariates leaves two-thirds of the disparity unexplained! Modeled inter-MSA variance decreases by about 28% when isolation is added to the models, suggesting that segregation accounts for a portion of this geographic variation. The persisting disparity results from unmeasured risk factors which may include racial differences in lifecourse health behaviors (Lu & Halfon, 2003), residual confounding by socioeconomic status (Kaufman & Cooper, 2008), pervasive exposures to chronic stress, or genetic and epigenetic interactions with any of the above (Hogue & Bremner, 2005).

Because of the history of slavery and racial inequality following the Civil War and into the current century, the patterns and consequences of black-white segregation we describe may be uniquely American. While our results may not be fully generalizable to other countries, ethnic and economic residential segregation likely occurs in all urbanized areas, and comparable tools can be applied to understanding the consequences of such segregation, although relationships may be similarly complex (Pickett, Shaw, Atkin, Kiernan, & Wilkinson, 2009).

Limitations

This cross-sectional, vital records-based observational study is limited by lack of information on important variables. Such variables include detailed individual biological and social exposures, and measures of neighborhoods in addition to metropolitan areas. Because birth records are not nationally available with neighborhood-level geocodes it is not possible

to distinguish whether the contextual effect of segregation or poverty is primarily exerted at the neighborhood or the metropolitan level.

Conclusions

Isolation segregation, previously associated with risk for low birthweight and preterm birth, is most strongly associated with *very* preterm birth, an outcome where the racial gap is also largest and the burden of mortality and morbidity is most severe. The association between racial isolation segregation and preterm birth persists under numerous model specifications, and is only attenuated modestly with control for individual socioeconomic variables and metropolitan murder rate. While important, this association remains small in magnitude and, like other known risk factors, explains only a fraction of the racial disparity. Future work should continue to explore the manner in which structural processes in urban areas influence pregnancy outcomes. These efforts will likely be most effective if combined with improved measurement at multiple scales of study from individual clinical and biological information, to neighborhood environment, and metropolitan characteristics (Kramer & Hogue, 2009a).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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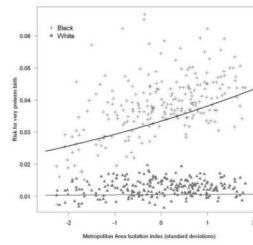


Figure 1. Metropolitan area very preterm birth risk by isolation segregation for black and white women

Symbols (cross and triangles) represent observed risk for VPT birth in each of 231 MSA's for black and white women respectively; lines represent model predicted risk for VPT birth from Table 2, Model M5.

Table 1

Distribution of singleton live births, 231 metropolitan statistical areas, 2000–2002

	Non-Hispanic black mothers			Non-Hispanic white mothers			B/W Risk Ratio		
	N births	%VPT	%MPT	N births	%VPT	%MPT	VPT	MPT	MPT
TOTAL	1,452,943	3.45	12.38	4,727,601	1.09	7.62	3.17	3.17	1.62
Maternal age									
<15	8,036	6.25	17.26	2,875	3.90	14.50	1.60	1.60	1.19
15–19	259,919	3.71	13.30	331,585	1.88	9.36	1.97	1.97	1.42
20–24	462,992	3.05	11.89	931,881	1.26	8.06	2.42	2.42	1.48
25–29	335,450	3.16	11.41	1,260,933	0.94	7.28	3.36	3.36	1.57
30–34	235,487	3.69	12.18	1,377,871	0.89	7.01	4.15	4.15	1.74
35–39	121,912	4.17	14.15	681,430	1.08	7.66	3.86	3.86	1.85
40+	29,147	4.85	16.04	141,026	1.36	9.34	3.57	3.57	1.72
Maternal education ^d									
<12 years	346,198	3.92	14.30	485,564	1.83	9.99	2.14	2.14	1.43
12 years	555,945	3.44	12.53	1,322,573	1.31	8.30	2.63	2.63	1.51
>12 years	525,107	3.01	10.93	2,879,627	0.84	6.91	3.58	3.58	1.58
Marital status									
Married	467,200	2.83	10.94	3,736,839	0.89	7.16	3.18	3.18	1.53
Unmarried	985,743	3.74	13.06	990,762	1.85	9.37	2.02	2.02	1.39
Parity									
Primiparous	432,811	3.42	11.49	1,609,656	1.24	7.80	2.76	2.76	1.47
Multiparous	1,020,132	3.46	12.76	3,117,945	1.01	7.53	3.43	3.43	1.69
Prior preterm or SGA birth ^d									
Yes	20,637	11.18	26.25	54,813	3.84	22.22	2.91	2.91	1.18
No	1,420,965	3.32	12.18	4,618,748	1.05	7.44	3.16	3.16	1.64
Chronic hypertension ^d									
Yes	20,519	8.54	21.34	35,256	3.46	16.31	2.47	2.47	1.31
No	1,421,083	3.36	12.25	4,628,305	1.07	7.55	3.14	3.14	1.62
Diabetes ^d									

	Non-Hispanic black mothers			Non-Hispanic white mothers			B/W Risk Ratio		
	N births	%VPT	%MPT	N births	%VPT	%MPT	VPT	MPT	MPT
Yes	41,956	3.42	16.92	135,907	1.17	11.94	2.92	1.42	
No	1,399,646	3.44	12.25	4,537,654	1.08	7.49	3.19	1.64	
Tobacco use ^d									
Yes	122,275	4.94	15.85	576,214	1.84	10.02	2.68	1.58	
No	1,237,533	3.33	12.15	3,707,273	0.99	7.38	3.36	1.65	
Region									
Northeast	275,059	3.36	11.42	1,017,475	1.01	6.68	3.33	1.71	
Southeast	741,877	3.52	12.84	1,639,918	1.19	8.48	2.96	1.51	
Midwest	316,917	3.60	12.68	1,213,167	1.13	7.65	3.19	1.66	
West	119,090	2.77	10.94	857,041	0.94	7.07	2.95	1.55	

Source: National Center for Health Statistics Natality Files, 2000–2002

Abbreviations: VPT: Very preterm birth (greater than 20 but less than 32 weeks gestation); MPT: Moderately preterm birth (greater than or equal to 32 weeks but less than 37 weeks gestation)

^dDemographic factor does not sum to total due to missing values in natality files

Table 2

Parameter estimates for very preterm birth among black women

Individual level	Crude ^a		M1 -- Age, parity and prior preterm birth		M2 -- Socioeconomic condition pathway		M3 -- Maternal health pathway		M4 -- All covariate model		M5 -- Dual Index model						
	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI					
Maternal age																	
<15	2.12	1.93 2.34	2.23	2.03	2.45	1.66	1.51	1.83	2.29	2.08	2.50	1.62	1.80	1.97	1.80	1.64	1.99
15-19	1.18	1.15 1.21	1.22	1.18	1.25	0.97	0.94	1.00	1.23	1.19	1.27	0.98	1.05	1.02	1.02	0.98	1.05
20-24	0.96	0.93 0.98	0.96	0.94	0.99	0.86	0.84	0.89	0.97	0.94	0.99	0.85	0.88	0.90	0.88	0.86	0.90
25-29	<i>ref</i>		<i>ref</i>		<i>ref</i>				<i>ref</i>				<i>ref</i>		<i>ref</i>		<i>ref</i>
30-34	1.21	1.17 1.24	1.21	1.17	1.24	1.30	1.26	1.34	1.19	1.16	1.23	1.23	1.26	1.30	1.27	1.23	1.31
35-39	1.42	1.37 1.47	1.41	1.37	1.46	1.54	1.49	1.60	1.35	1.31	1.40	1.41	1.46	1.51	1.46	1.41	1.51
40+	1.71	1.62 1.82	1.71	1.61	1.81	1.85	1.74	1.96	1.58	1.49	1.68	1.60	1.70	1.80	1.70	1.61	1.80
Parity																	
Primiparous	<i>ref</i>		<i>ref</i>		<i>ref</i>				<i>ref</i>				<i>ref</i>		<i>ref</i>		<i>ref</i>
Multiparous	1.07	1.05 1.09	1.00	0.97	1.02	0.97	0.95	0.99	0.97	0.95	1.00	0.94	0.96	0.98	0.96	0.94	0.98
Prior preterm or SGA birth																	
Yes	4.75	4.54 4.98	4.77	4.57	5.00	4.69	4.47	4.91	4.56	4.34	4.79	4.31	4.53	4.74	4.54	4.34	4.75
No	<i>ref</i>		<i>ref</i>		<i>ref</i>				<i>ref</i>				<i>ref</i>		<i>ref</i>		<i>ref</i>
Maternal education																	
<12 years	1.27	1.25 1.30				1.21	1.18	1.24				1.11	1.15	1.18	1.15	1.12	1.18
12 years	1.10	1.08 1.12				1.08	1.06	1.11				1.04	1.06	1.09	1.06	1.04	1.09
>12 years	<i>ref</i>				<i>ref</i>				<i>ref</i>				<i>ref</i>		<i>ref</i>		<i>ref</i>
Marital status																	
Married	<i>ref</i>				<i>ref</i>				<i>ref</i>				<i>ref</i>		<i>ref</i>		<i>ref</i>
Unmarried	1.36	1.33 1.39				1.47	1.44	1.51				1.40	1.43	1.47	1.43	1.40	1.47
Chronic hypertension																	
Yes vs no	3.03	2.87 3.18							2.77	2.64	2.93	2.61	2.75	2.90	2.75	2.61	2.90
Diabetes																	
Yes vs. no	0.95	0.90 1.00							0.90	0.85	0.95	0.87	0.91	0.97	0.91	0.87	0.96

	Crude ^d		M1 -- Age, parity and prior preterm birth		M2 -- Socioeconomic condition pathway		M3 -- Maternal health pathway		M4 -- All covariate model		M5 -- Dual Index model	
	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI
Tobacco use ^c												
Yes vs no	1.58	1.54 - 1.63					1.53	1.48 - 1.57	1.38	1.34 - 1.42	1.38	1.34 - 1.42
MSA-level												
Isolation ^b	1.11	1.08 - 1.14	1.12	1.08 - 1.15	1.10	1.07 - 1.14	1.13	1.10 - 1.17	1.11	1.08 - 1.15	1.15	1.10 - 1.19
Clustering ^b	1.05	1.02 - 1.08									0.95	0.92 - 0.99
Population size												
<500,000	1.09	1.04 - 1.16	1.10	1.04 - 1.16	1.08	1.02 - 1.14	1.08	1.03 - 1.14	1.07	1.01 - 1.13	1.05	0.99 - 1.11
500k - 1 million	1.08	1.01 - 1.15	1.09	1.02 - 1.16	1.09	1.01 - 1.16	1.08	1.01 - 1.15	1.07	1.00 - 1.14	1.07	1.00 - 1.14
1 million +	<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>	
Region												
Northeast	<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>		<i>ref</i>	
Southeast	0.98	0.92 - 1.05	1.04	0.96 - 1.12	1.07	0.99 - 1.15	1.07	1.00 - 1.15	1.09	1.02 - 1.17	1.04	0.97 - 1.13
Midwest	0.98	0.91 - 1.06	1.01	0.93 - 1.09	1.00	0.92 - 1.08	1.01	0.93 - 1.09	1.00	0.93 - 1.09	1.01	0.94 - 1.09
West	0.84	0.76 - 0.92	0.86	0.78 - 0.95	0.88	0.80 - 0.98	0.93	0.84 - 1.03	0.94	0.85 - 1.03	0.91	0.83 - 1.01
σ^2_α			0.134		0.135		0.126		0.129		0.125	
DIC			416696		415134		414758		413557		413559	

Abbreviations: MSA: metropolitan statistical area

^aCrude models for individual covariates all include random intercepts for MSA, and adjustment for region and population size

^bThe odds ratios for segregation indices refer to the change in risk of very preterm birth for a 1-SD change in segregation index

^cCalifornia does not collect smoking on birth certificates so smoking parameter estimates exclude births in California. The variable was coded such that California births are not omitted from the model so all other parameters are inclusive.

Table 3

Parameter estimates for moderately preterm birth among black women

	Crude ^a		M1 -- Age, parity and prior preterm birth		M2 -- Socioeconomic condition pathway		M3 -- Maternal health pathway		M4 -- All covariate model		M5 -- Dual Index model	
	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI
MSA-level												
Isolation ^b	1.06	1.04 1.09	1.07	1.04 1.10	1.06	1.03 1.09	1.08	1.05 1.11	1.07	1.04 1.09	1.08	1.04 1.11
Clustering ^b	1.04	1.02 1.07									0.99	0.96 1.02
σ^2_{α}			0.134		0.134		0.131		0.132		0.132	
DIC			1065390		1062570		1062380		1060170		1060180	

Abbreviations: MSA: metropolitan statistical area

^aCrude models for individual covariates all include random intercepts for MSA, and adjustment for region and population size

^bThe odds ratios for segregation indices refer to the change in risk of very preterm birth for a 1-SD change in segregation index

Table 4

Candidate metropolitan level mediating variables for association between segregation and very preterm birth among black women

	Murder		Poverty		Poverty rate ratio		Poverty concentration	
	OR	95% CI	OR	95% CI	OR	95% CI	OR	95% CI
Isolation ^d	1.12	1.07 1.17	1.15	1.10 1.20	1.15	1.10 1.20	1.16	1.11 1.21
Clustering ^a	0.95	0.91 0.98	0.95	0.91 0.99	0.96	0.92 0.99	0.95	0.91 1.00
Murder rate ^b	1.12	1.03 1.21						
Black poverty rate ^c			1.03	0.98 1.07				
Black-white poverty rate ratio					1.00	0.97 1.03		
Black poverty concentration ^d							0.98	0.96 1.01
σ^2	0.120			0.125		0.127		0.122
DIC	410126			413558		413558		406547

NOTE: All models are also adjusted for maternal age, education, parity, marital status, smoking, history of prior preterm birth, chronic hypertension or diabetes, MSA region and MSA population size.

^aOdds ratios for isolation and clustering indices correspond to the change in the outcome for a 1-SD increase in segregation

^bMurder rate is scaled so that a 1-unit change in murder rate is equivalent to 10 murders/100,000 persons

^cBlack poverty rate is scaled so that a 1-unit change is equivalent to 10% change in poverty rate

^dBlack poverty concentration is the proportion of children in poor families who also live in high poverty neighborhoods. It is scaled so that a 1-unit change is equivalent to 10% change in this proportion.

Table 5
Black-white disparities in very preterm birth under different model specifications

	OR	95%CI	% disparity explained	σ^2_{race}	DIC
Crude	3.24	3.20 3.28			
All individual variables	2.68	2.37 3.02	25.0	0.107	960703
Individual + isolation	2.49	2.25 2.77	33.4	0.077	960681
Individual + isolation + clustering	2.51	2.25 2.80	32.5	0.075	960677