

Linking US registry data with geomarker data: longitudinal associations between environmental factors and lung function decline

European Cystic Fibrosis Conference

Eleni-Rosalina Andrinopoulou, Pedro Afonso, Rhonda Szczesniak, Grace C. Zhou, John P. Clancy, Anushka Palipana, Erika Rasnick, Cole Brokamp, Patrick Ryan, Ruth Keogh

10 June, 2022

Introduction: Motivation

A lot of information is available
→ Electronic medical records

Introduction: Motivation

A lot of information is available
→ Electronic medical records

Different types of information

- Baseline characteristics: Sex, F508del, SESlow, Enzymes
- Biomarkers: FEV₁ % pred
- Nutritional status: BMI percentile
- Geomarkers (environmental/community factors): Deprivation index



This research is supported by the National Institutes of Health / National Heart, Lung and Blood Institute (grant R01 HL141286)

Introduction: Research question

How can we appropriately connect registry data with geomarker data

Introduction: Descriptive statistics

Deprivation index (https://geomarker.io/dep_index/)

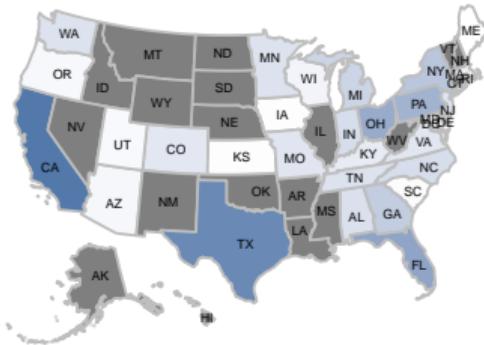
- Socioeconomic variables from the American Community Survey (ACS): capture “community deprivation”
 - ◊ Principal components analysis of six different 2015 ACS measures
 - ◊ “Deprivation Index”: the first component explains over 60% of the total variance
 - ◊ Rescaling and normalizing forces the index to range from 0 to 1, with a higher index being more deprived



Cole Brokamp, Andrew F. Beck, Neera K. Goyal, Patrick Ryan, James M. Greenberg, Eric S. Hall. Material Community Deprivation and Hospital Utilization During the First Year of Life: An Urban Population-Based Cohort Study. Annals of Epidemiology. 30. 37-43. 2019

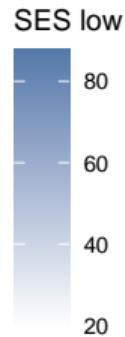
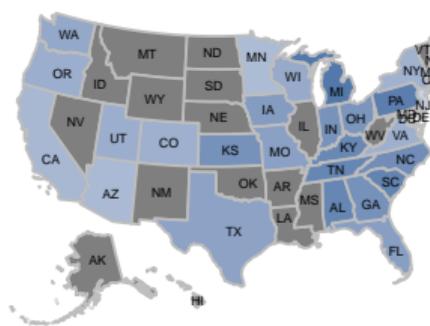
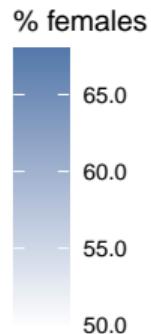
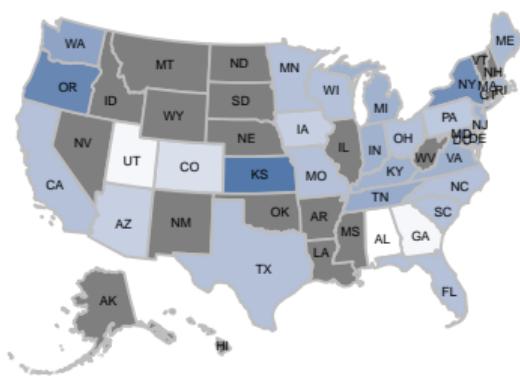
Introduction: Descriptive statistics

Baseline information



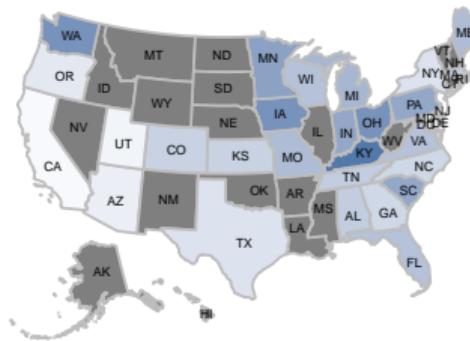
Introduction: Descriptive statistics

Baseline information

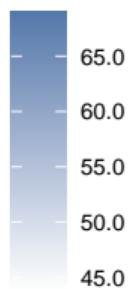


Introduction: Descriptive statistics

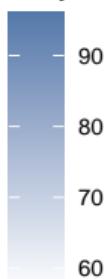
Baseline information



F508del Homozygous



Enzymes



Introduction: Descriptive statistics

Biomarker

Introduction: Descriptive statistics

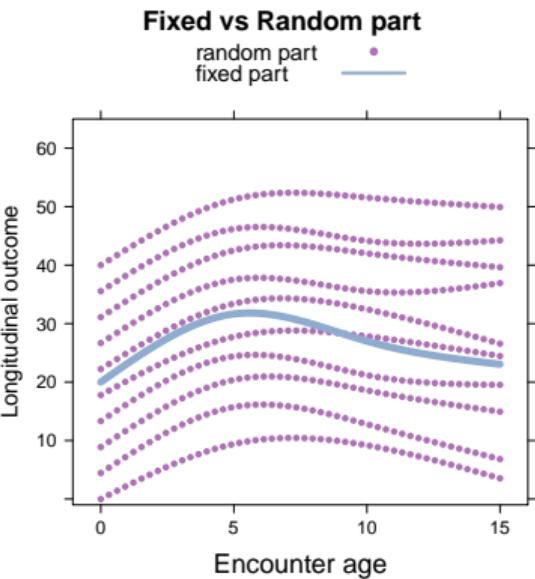
Geomarker

Introduction: Descriptive statistics

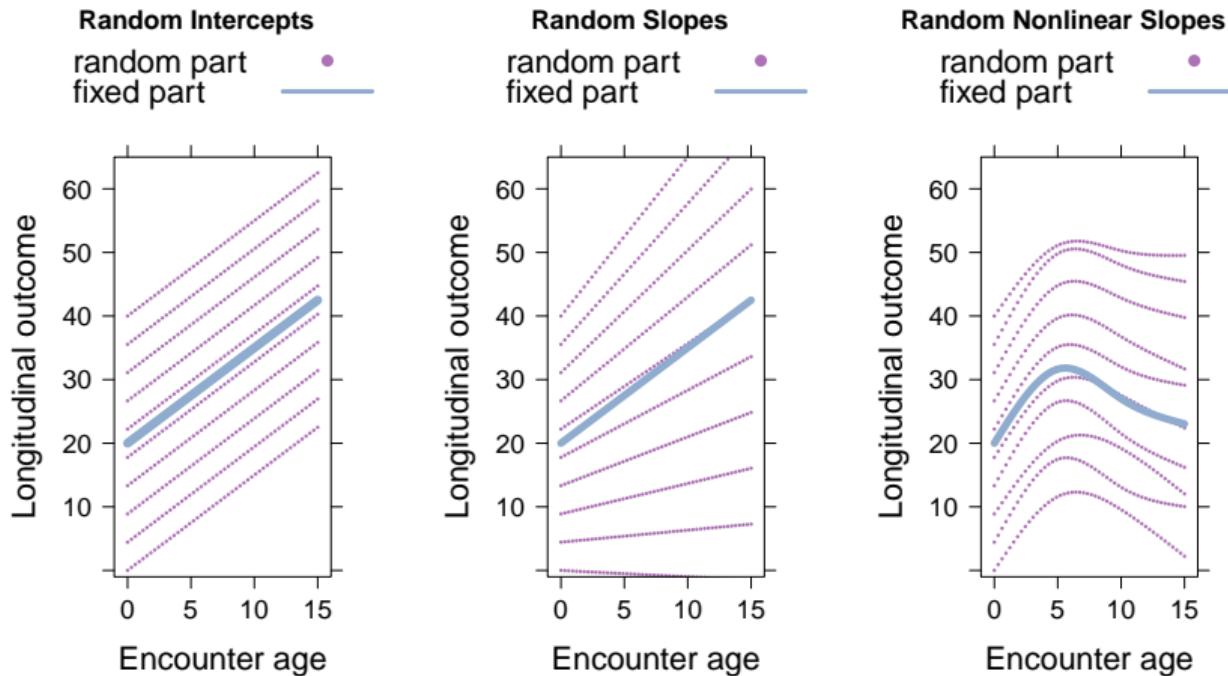
Nutritional status

Methods: Univariate Mixed Model

$$y_i(t) = \underbrace{x_i^\top(t)\beta_1}_{\text{fixed effects part}} + \underbrace{z_i(t)^\top b_i}_{\text{random effects part}} + \epsilon_i$$



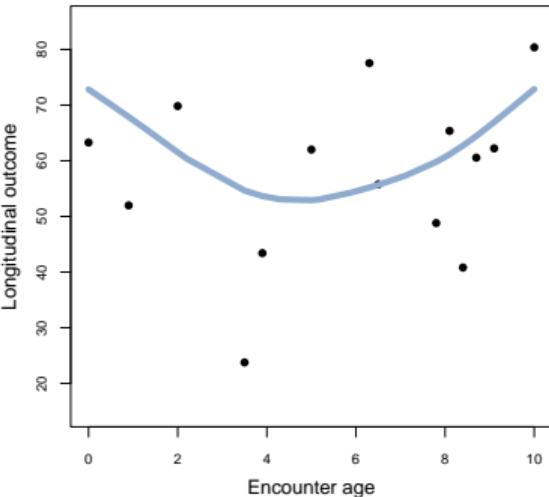
Methods: Univariate Mixed Model



Methods: Univariate Mixed Model

$$y_i(t) = \underbrace{x_i^\top(t)\beta_1}_{\text{fixed effects part}} + \underbrace{z_i(t)^\top b_i}_{\text{random effects part}} + \epsilon_i$$

$$= m_i(t) + \epsilon_i$$



Methods: Multivariate Mixed Models

DepIndex = encounterage

FEV1%pred = encounterage + sex + SESlow +

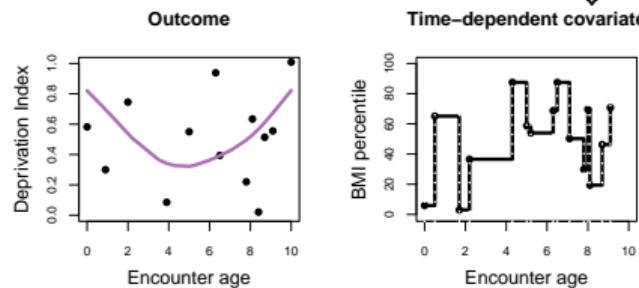
F508del + BMIpercentile + Enzymes

Methods: Multivariate Mixed Models

DepIndex = encounterage

FEV1%pred = encounterage + sex + SESlow +

F508del + BMIpercentile + Enzymes

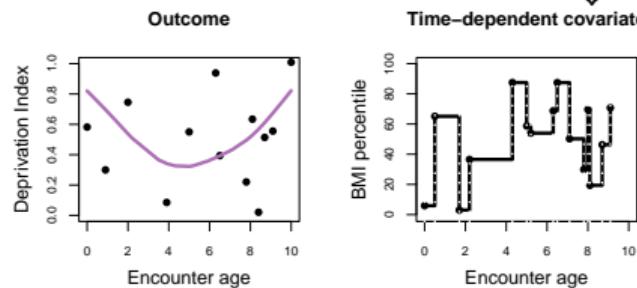


Methods: Multivariate Mixed Models

DepIndex = encounterage

FEV1%pred = encounterage + sex + SESlow +

F508del + BMIpercentile + Enzymes + DepIndex



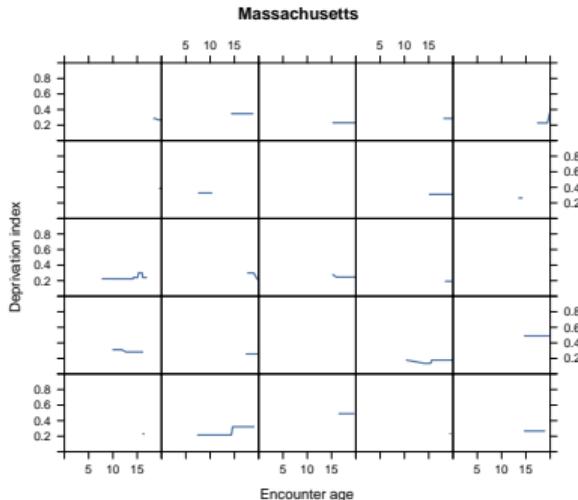
Methods: Multivariate Mixed Models

$$\text{DepIndex: } y_{2i}(t) = m_{2i}(t) + \epsilon_{2i}(t) = x_{2i}^\top(t)\beta_1 + z_{2i}(t)^\top b_{2i} + \epsilon_{2i}(t)$$

$$\text{FEV1\%pred: } y_{1i}(t) = x_{1i}^\top(t)\beta_1 + z_{1i}(t)^\top b_{1i} + \alpha_{S2} m_{2i}(t) + \epsilon_{1i}(t)$$

where

$$\diamond \quad b_i^\top = (b_{1i}^\top, b_{2i}^\top) \sim N(0, D)$$



Methods: Multivariate Mixed Models

$$\text{DepIndex: } y_{2i}(t) = m_{2i}(t) + \epsilon_{2i}(t) = x_{2i}^\top(t)\beta_1 + z_{2i}(t)^\top b_{2i} + \epsilon_{2i}(t)$$

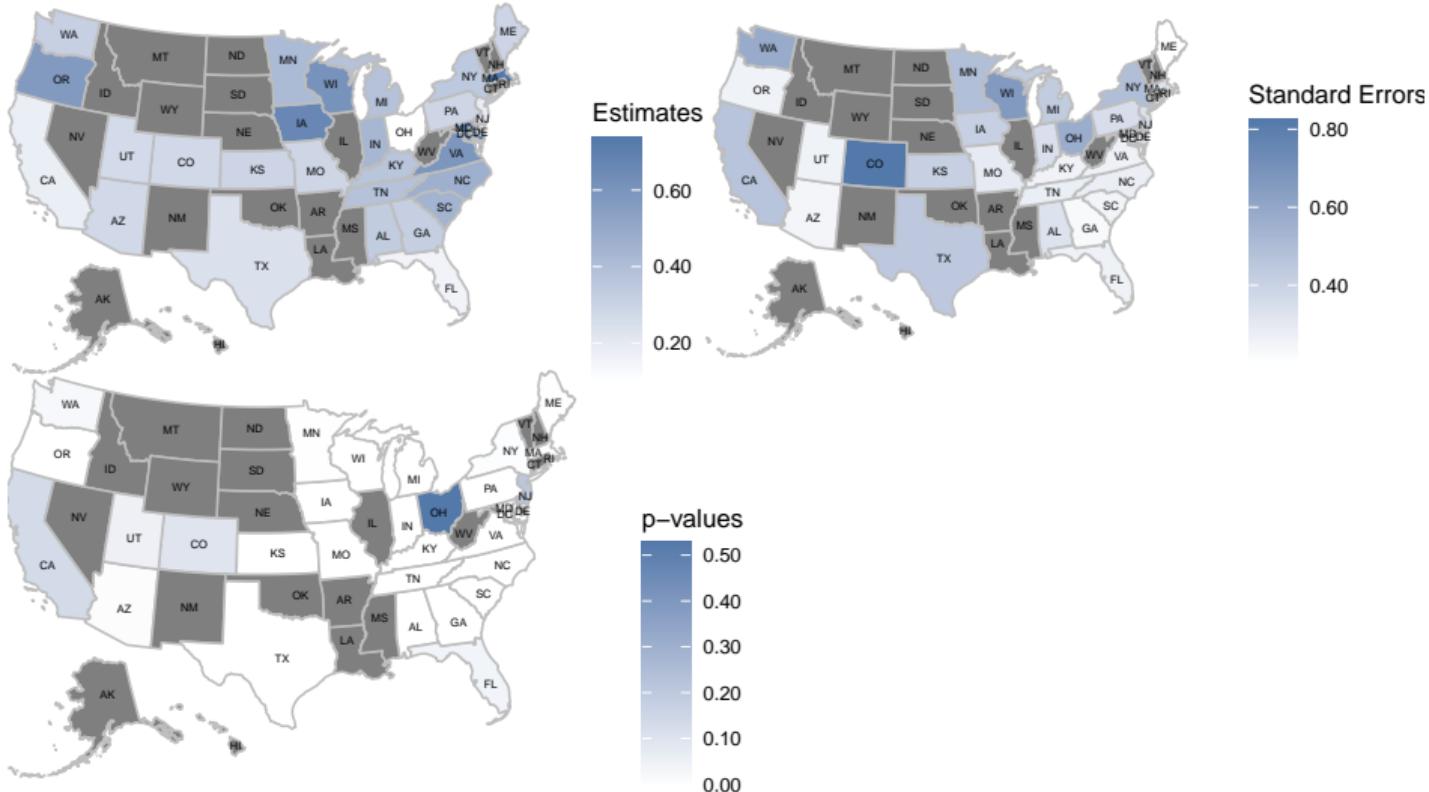
$$\text{FEV1%pred: } y_{1i}(t) = x_{1i}^\top(t)\beta_1 + z_{1i}(t)^\top b_{1i} + \alpha_{S2} \int_0^t m_{2i}(s)ds + \epsilon_{1i}(t)$$

where

$$\diamond \quad b_i^\top = (b_{1i}^\top, b_{2i}^\top) \sim N(0, D)$$

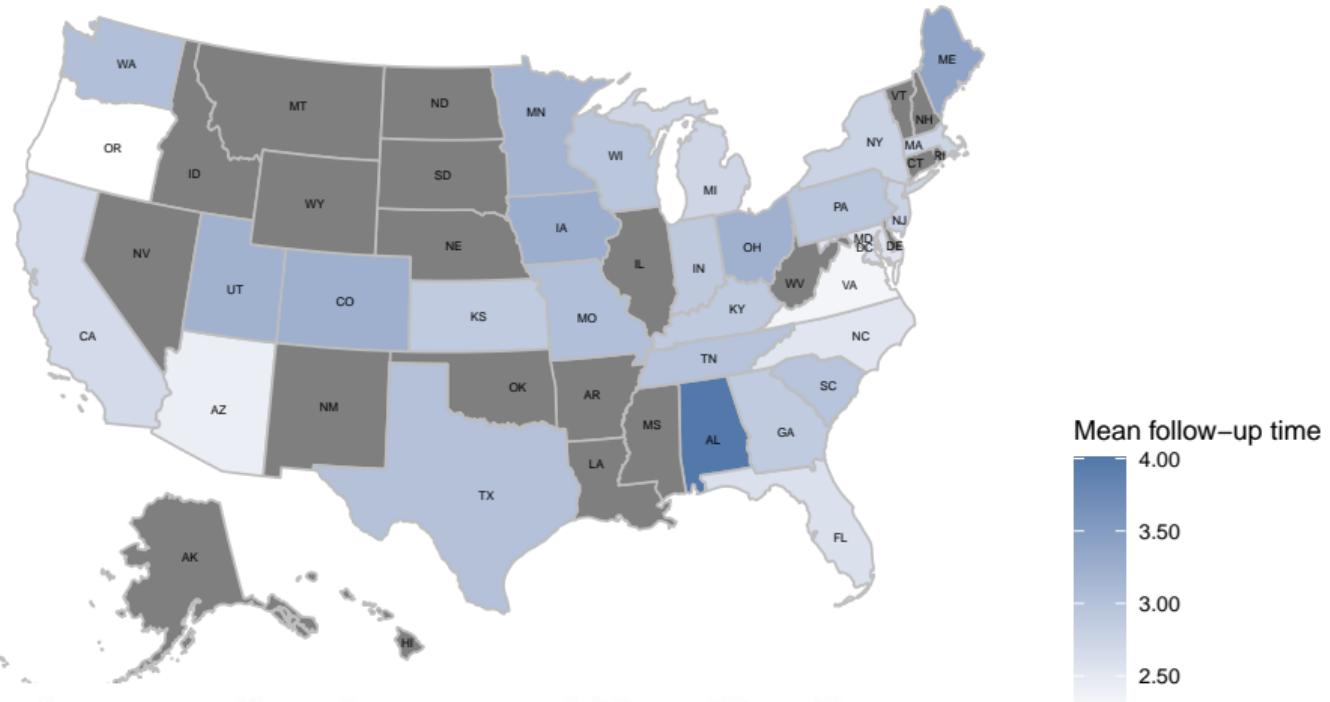
Results: Multivariate Mixed Models

Estimate: for 0.1 unit increase in the area under the curve of the deprivation index



Introduction: Descriptive statistics

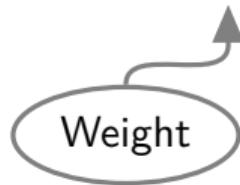
Baseline information



Methods: Multivariate Mixed Models

DepIndex: $y_{2i}(t) = m_{2i}(t) + \epsilon_{2i}(t) = x_{2i}^\top(t)\beta_1 + z_{2i}(t)^\top b_{2i} + \epsilon_{2i}(t)$

FEV1%pred: $y_{1i}(t) = x_{1i}^\top(t)\beta_1 + z_{1i}(t)^\top b_{1i} + \alpha_{S2} \frac{1}{t} \int_0^t m_{2i}(s)ds + \epsilon_{1i}(t)$

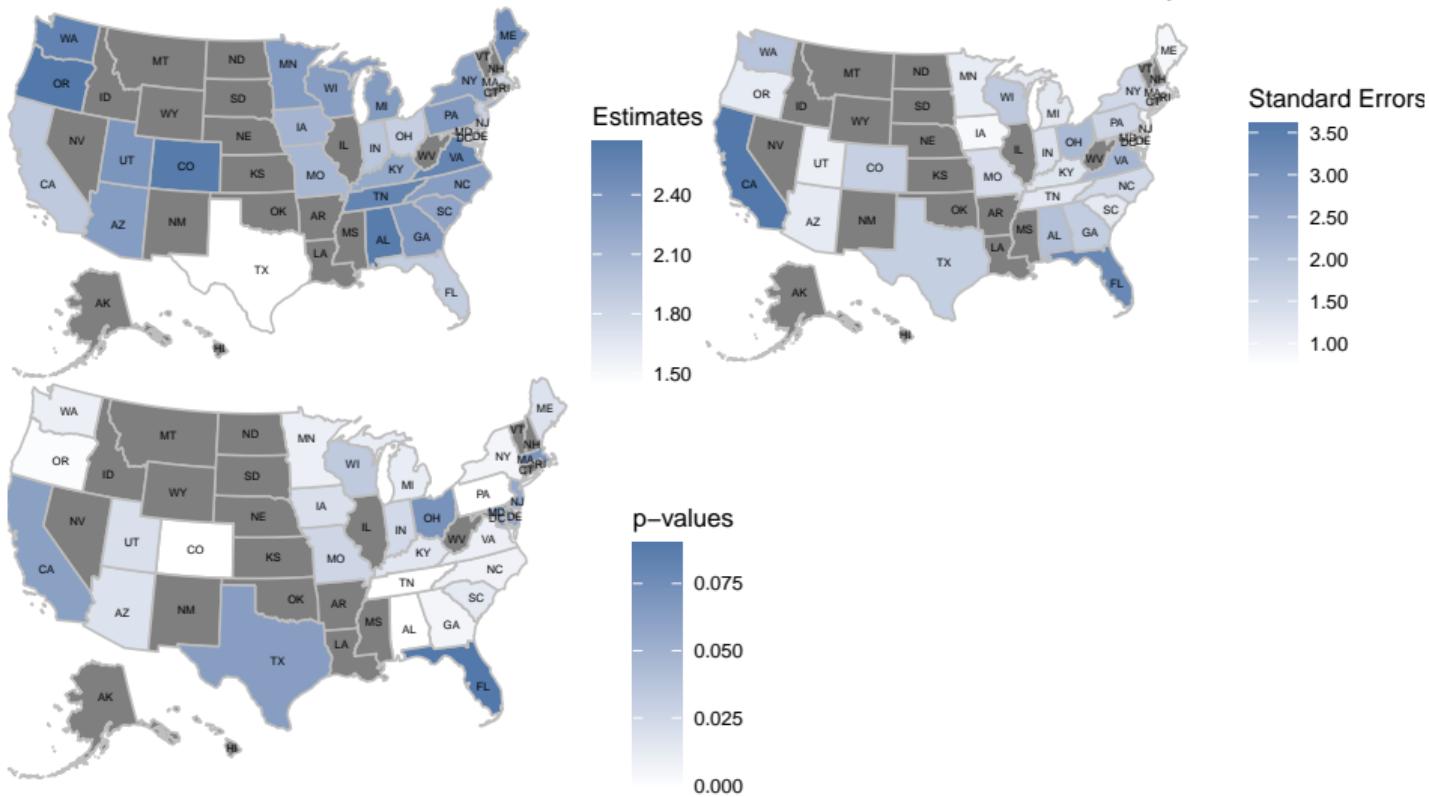


where

◇ $b_i^\top = (b_{1i}^\top, b_{2i}^\top) \sim N(0, D)$

Results: Multivariate Mixed Models

Estimate: for 0.1 unit increase in the normalized area under the curve of deprivation index



Methods: Multivariate Mixed Models

FEV1%pred = encounterage

DepIndex = encounterage + FEV1%pred

Methods: Multivariate Mixed Models

$$\text{FEV1\%pred: } y_{1i}(t) = m_{1i}(t) + \epsilon_{1i} = x_{1i}^\top(t)\beta_1 + z_{1i}(t)^\top b_{1i} + \epsilon_{1i}(t)$$

$$\text{DepIndex: } y_{2i}(t) = m_{2i}(t) + \epsilon_{2i} = x_{2i}^\top(t)\beta_1 + z_{2i}(t)^\top b_{2i} + \alpha_{S1} \frac{d}{dt}m_{1i}(t) + \epsilon_{2i}(t)$$

where

$$\diamond \quad b_i^\top = (b_{1i}^\top, b_{2i}^\top) \sim N(0, D)$$

Discussion

- A lot of data is available
- Better treatment and monitoring strategies if all information is used
- Challenge in combining different types of information
- Investigate other geomarker data

Thank you for your attention!