

Understanding the Opioid Syndemic in North Carolina: A Novel Approach to Modeling and Identifying Factors

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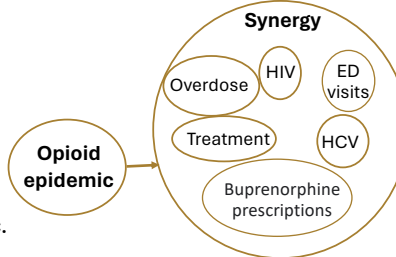


Motivation

- **Opioid epidemic** is a critical public health problem, e.g. in 2021 4041 NC residence died from opioid overdose

Challenge: no data to fully capture this complex problem.

Need of an alternative approach



We know that:

- Opioid epidemic is associated with HIV and HCV infections.
- Fatal and nonfatal overdose, ED admissions and treatment admissions are also linked to opioid misuse.
- These epidemics interact synergistically, forming a syndemic.

Identify underlying factors that explain dependencies in the observed data.

Identifiability of $\log(\lambda_{ij}) = \Gamma F_{ij} + \epsilon_{ij}$

- A well-known challenge in factor analysis modeling is the identifiability¹ of model parameters, i.e., The likelihood $f(Y_{ij} | \Gamma, F_{ij}, \epsilon_{ij})$ is equivalent to $f(Y_{ij} | \Gamma H, H' F_{ij}, \epsilon_{ij})$ for any $m \times m$ orthogonal H .
- **Common solution:** specify the loadings matrix to be lower triangular² \Rightarrow factors lacking meaningful interpretation.
- **Proposed solution:** specify the loadings matrix Γ as needed to quantify meaningful relationships among subsets of outcomes and employ LQ decomposition³ on Γ to decompose it into a lower triangular matrix with **positive diagonal elements**, L , and an orthogonal matrix, Q , i.e., $\log(\lambda_{ij}) = \Gamma F_{ij} + \epsilon_{ij} = LQF_{ij} + \epsilon_{ij} = L \tilde{F}_{ij} + \epsilon_{ij}$, ensuring that the new model adheres to the identifiability constraints⁴ of a factor model.

Illustration:

$\log(\lambda_{ijk}) = \Gamma_k' F_{ij} + \epsilon_{ijk}, k = 1, \dots, 4$, written in matrix form:

$$\log(\lambda_{ij}) \equiv \begin{bmatrix} \log(\lambda_{ijD}) \\ \log(\lambda_{ijE}) \\ \log(\lambda_{ijT}) \\ \log(\lambda_{ijB}) \\ \log(\lambda_{ijC}) \\ \log(\lambda_{ijI}) \end{bmatrix} = \begin{bmatrix} 1 & \gamma_{12} & 0 & 0 \\ \gamma_{21} & 1 & 0 & 0 \\ \gamma_{31} & 0 & 1 & 0 \\ \gamma_{41} & 0 & \gamma_{43} & 0 \\ \gamma_{51} & 0 & 0 & 1 \\ \gamma_{61} & 0 & 0 & \gamma_{64} \end{bmatrix} \cdot \begin{bmatrix} F_1 \\ F_2 \\ F_3 \\ F_4 \end{bmatrix} + \begin{bmatrix} \epsilon_D \\ \epsilon_E \\ \epsilon_T \\ \epsilon_C \\ \epsilon_I \end{bmatrix}$$

Must be identifiable \equiv lower triangular with positive diagonal elements

L factorization Q

Algorithm: Metropolis-Hastings Update of Loadings

- Let Γ denote the current value of the loadings with $\Gamma = LQ$.
- Propose a new value γ_{km}^* using a normal random walk, centered at the current value γ_{km} . Let Γ^* be the loadings matrix containing the proposed γ_{km}^* .
- Perform the LQ-decomposition of Γ^* :
 - Input (Γ^*)**
 - Let $\Gamma^{*'} = [\Gamma_2' | \Gamma_2^{*'}]$.
 - Apply the QR Gram-Schmidt process to get Q^* and R_1^* such that $\Gamma_1^{*'} = Q^* R_1^*$.
 - Compute $R_2^* = [Q^* \cdot \Gamma_2^{*'}]$.
 - Let $R^* = [R_1^* | R_2^*]$.
- Compute the Metropolis-Hastings acceptance probability: $\tau = \min\left(1, \frac{f(y|L(\Gamma^*), Q(\Gamma^*), F, \epsilon)\pi(\Gamma^*)}{f(y|L(\Gamma), Q(\Gamma), F, \epsilon)\pi(\Gamma)}\right)$.
- Set $\Gamma = \Gamma^*$ with probability τ .

Methodology

Bayesian Hierarchical Model:

- Observed count for outcome k in year j and in county i :

$$Y_{ijk} \sim \text{Poisson}(E_{ijk}\lambda_{ijk})$$

$$\log(\lambda_{ijk}) = \Gamma_k' F_{ij} + \epsilon_{ijk}$$

$$F_{ij} \sim \text{ICAR} - \text{AR}(1)$$

$$\epsilon_{ijk} \sim N(0, \sigma_k^2)$$

Factor Analysis

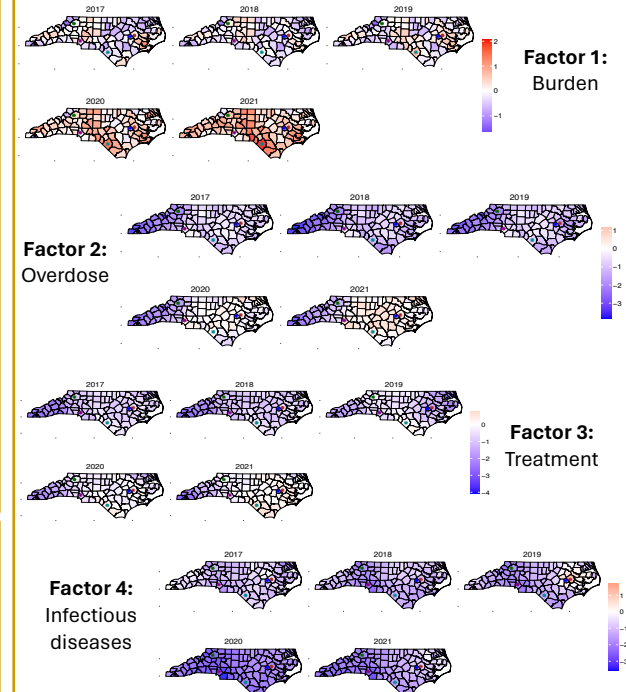
- allows us to design factors that yield **meaningful interactions** among predefined subsets of the the observed outcome.

Describe the opioid syndemic through 4 factors:

- Factor 1 to characterize **variation** shared by all **six outcomes: overall burden**.
- Factor 2 to characterize **variation** shared by **Death (D) and ED visits (E)**, after accounting for F1: **overdose aspect** of the epidemic.
- Factor 3 to characterize **variation** shared by **Treatment (T) and Buprenorphine (B)**, after accounting for F1: **treatment aspect** of the epidemic.
- Factor 4 to characterize the remaining shared **variation** between **HCV (C) and HIV (I)**, after accounting for F1: **infectious disease** elements of the epidemic.

Results

Posterior mean estimates of the four latent factors from 2017 to 2021. **Brown** star represents Clay county, **blue** star represents Green county, **dark green** star represents Mecklenburg county, **dark orange** star represents Pitt county, **dark magenta** star represents Robeson county, **dark cyan** star represents Wilkes county.



Next steps

- Combine county-level data with zip code-level data to understand the opioid epidemic at a more granular level.
- Use loadings that vary geographically.

References

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