

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

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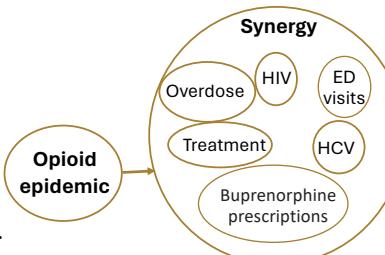


## Motivation

- 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 identifiability<sup>1</sup> 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 triangular<sup>2</sup>  $\Rightarrow$  factors lacking meaningful interpretation.
- Proposed solution:** specify the loadings matrix  $\Gamma$  as needed to quantify meaningful relationships among subsets of outcomes and employ LQ decomposition<sup>3</sup> on  $\Gamma$  to decompose it into a lower triangular matrix with **positive diagonal elements**,  $L$ , and an orthogonal matrix,  $Q$ , i.e.,  $\log(\lambda_{ij}) = \Gamma F_{ij} + \epsilon_{ij} = L Q F_{ij} + \epsilon_{ij} = L \tilde{F}_{ij} + \epsilon_{ij}$ , ensuring that the new model adheres to the identifiability constraints<sup>4</sup> of a factor model.

**Illustration:**

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

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

Must be identifiable  $\equiv$  lower triangular with positive diagonal elements  
L      Q  
factorization

## Algorithm: Metropolis-Hastings Update of Loadings

0. Let  $\Gamma$  denote the current value of the loadings with  $\Gamma = LQ$ .

1. Propose a new value  $\gamma_{km}^*$  using a normal random walk, centered at the current value  $\gamma_{km}$ . Let  $\Gamma^*$  be the loadings matrix containing the proposed  $\gamma_{km}^*$ .

2. Perform the LQ-decomposition of  $\Gamma^*$ :

**Input**( $\Gamma^*$ )

- Let  $\Gamma^{*\prime} = [\Gamma_2^* | \Gamma_2^*']$ .

- Apply the QR Gram-Schmidt process to get  $Q^*$  and  $R_1^*$  such that  $\Gamma_1^{*\prime} = Q^* R_1^*$ .

- Compute  $R_2^* = [Q^{*\prime} | \Gamma_2^*']$ .

- Let  $R^* = [R_1^* | R_2^*]$ .

**Output**( $L^* = R^{*\prime}, Q^* = Q^*$ )

3. Compute the Metropolis-Hastings acceptance probability:  $\tau = \min\left(1, \frac{f(y|L(\Gamma^*), Q(\Gamma^*), F, \epsilon)\pi(\Gamma^*)}{f(y|L(\Gamma), Q(\Gamma), F, \epsilon)\pi(\Gamma)}\right)$ .

4. Set  $\Gamma = \Gamma^*$  with probability  $\tau$ .

## 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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## Methodology

### Bayesian Hierarchical Model:

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

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

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

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

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

### 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.

