

Estimate of true COVID-19 infections in US Counties



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Background

- Goal
 - Predict the true number of people infected with the COVID-19 virus in each county in the US
- Data
 - COVID-19 infection data per county (4/20/20)
 - County-level predictors (2738 of 3007 counties)
 - Log population density, infection increase over previous 2 days, socio-economic vulnerability index, number of retirement homes
 - State-level predictors
 - Cook Partisan Voting Index, hospitals per person, number of tests administered per person

Model Structure

$$\text{Confirmed}_i \sim \text{NegBinom}(r, \frac{r}{m_i})$$

r = size, m_i = mean

$$m_i = \underbrace{\text{pop}_i * \lambda_i}_{\text{true \# infected}} * \theta_j$$

λ_i = true rate of infection
 θ_j = underestimation proportion

State level

$$\theta_j \sim \text{Beta}(rr, p_j, rr * (1 - p_j))$$

rr = size, p_j = mean

$$p_j = \text{inv_logit}(Z_j \gamma)$$

$$\gamma_p \sim \text{Double_Exp}(0, \sigma_\gamma)$$

LASSO prior

$$\gamma_1 \sim \text{Normal}(-2.28, 0.368)$$

Informative prior

County level

$$\lambda_i = \text{inv_logit}(X_i \beta)$$

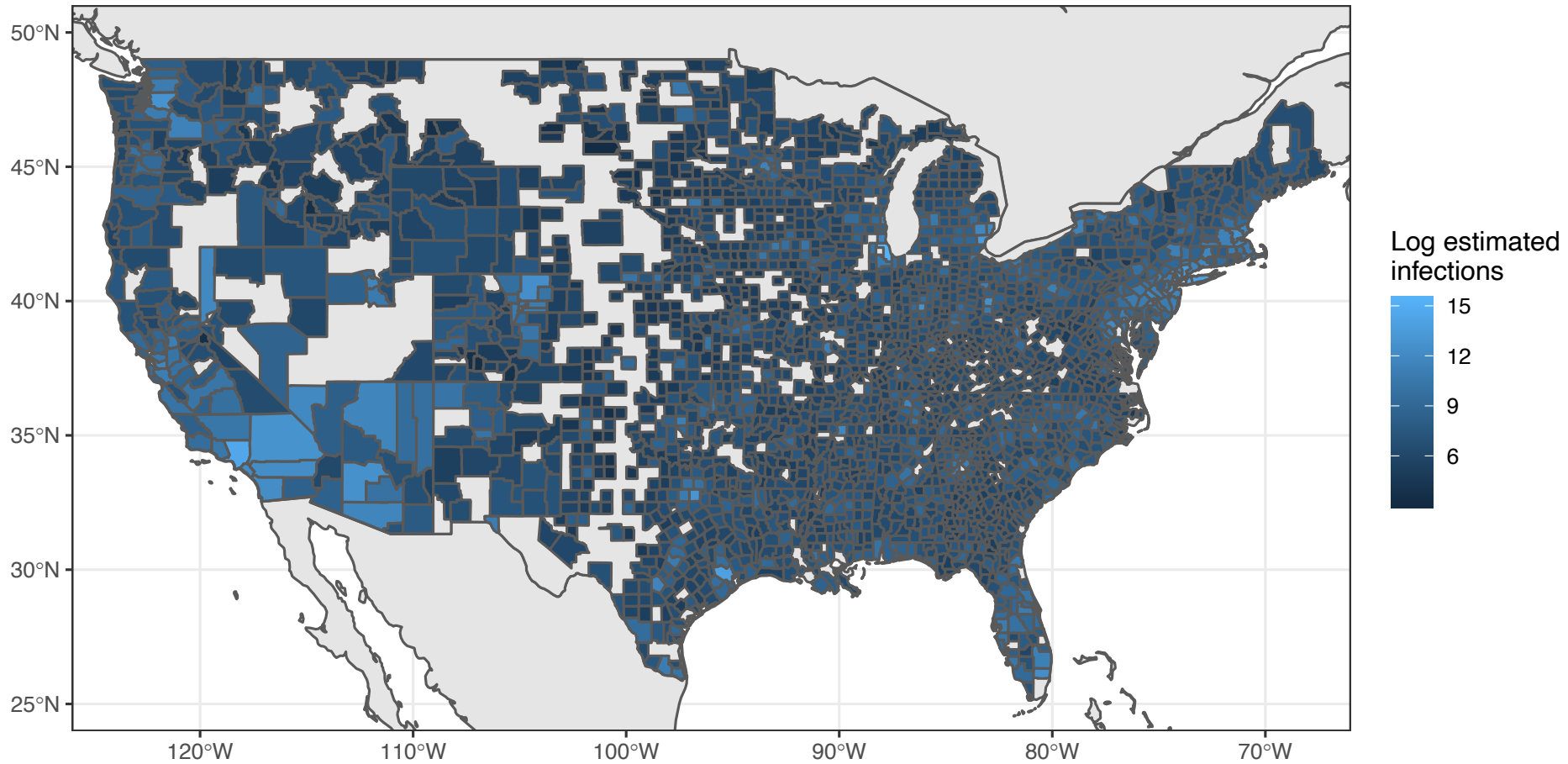
$$\beta_k \sim \text{Double_Exp}(0, \sigma_\beta)$$

LASSO prior

Details

- B-splines
 - Univariate b-splines with 5 df
 - Tensor product multivariate splines
 - Splines thinned based on max value
- Informative prior for underestimation proportion (θ)
 - Based on the range of antibody testing from New York State recently reported
 - Better estimate of true # of infections
- Problems estimating both true infection rate and underestimation proportion at county level
- Model fit with Stan
 - 20,000 sampling iterations

Results



Estimated number of true infections for select counties

County	Lower HDI	Upper HDI
AZ_Maricopa	17,692	906,079
CA_Los Angeles	98,301	8,669,920
GA_Fulton	52,901	124,507
IL_Cook	646,152	5,223,719
NC_Mecklenburg	31,716	72,620
NC_Wake	20,946	48,657
NY_Westchester	194,720	968,815
WA_King	32,534	1,035,048

