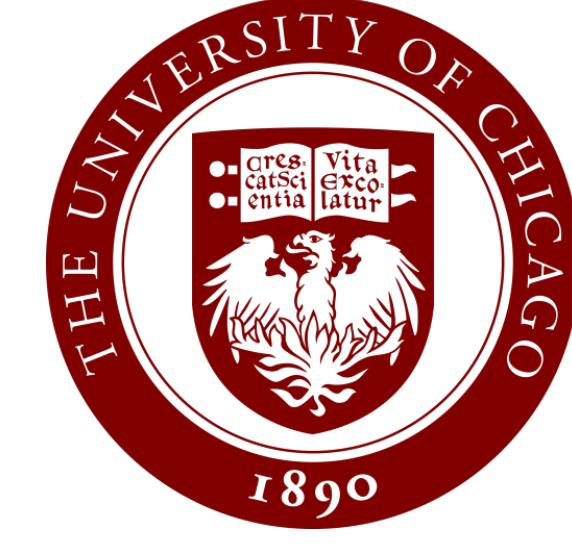


Bayesian Analysis of Latent Underdispersion Using Discrete Order Statistics

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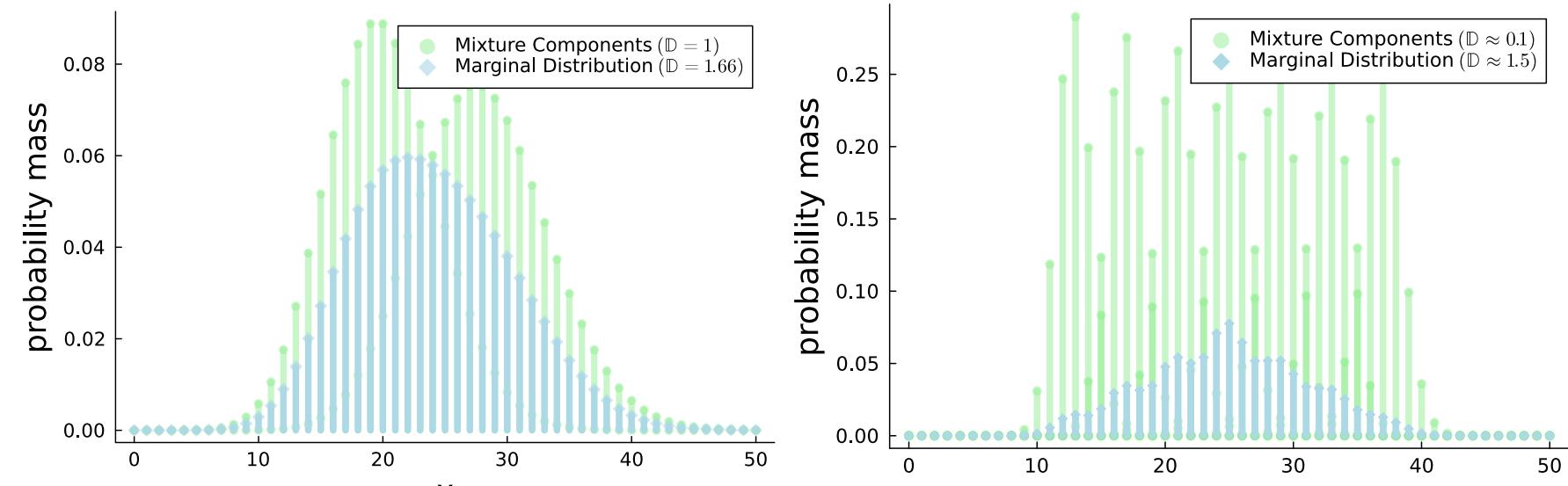
Conditional Underdispersion

Problem: no flexible tools exist to model underdispersed count data in probabilistic modeling frameworks.

Definition: A discrete random variable X is underdispersed with respect to the Poisson distribution if

$$\mathbb{D}[X] = \mathbb{V}[X]/\mathbb{E}[X] < 1$$

Count data which is *marginally* overdispersed may be consistent with a model that is underdispersed *conditionally*, once covariates or latent variables are observed.



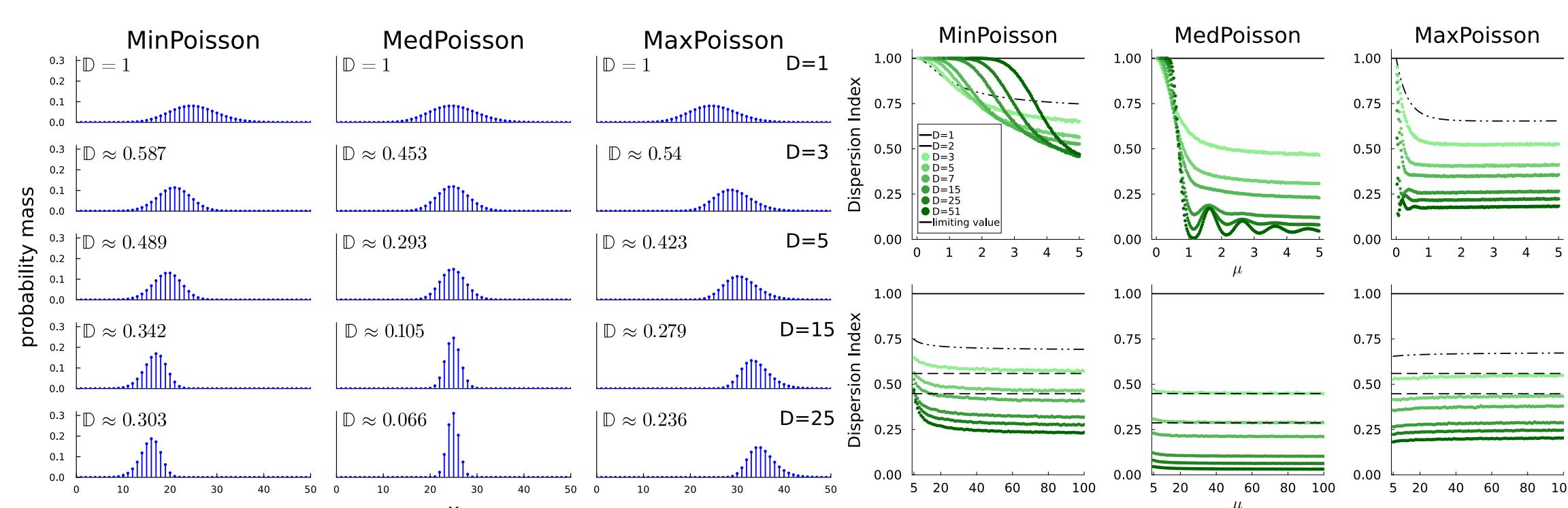
Key Benefit: Modeling conditional underdispersion allows for more precise probabilistic predictions than are possible with a Poisson likelihood.

Poisson Order Statistics are Underdispersed

We introduce discrete order statistics as a modular building block for probabilistic modeling. In particular, we build models with likelihoods of the form:

$$Y = Z_{(j,D)} \text{ for } j \in \{1, \dots, D\}, \text{ where } Z_d \stackrel{\text{iid}}{\sim} \text{Poisson}(\mu)$$

This family of likelihoods gives rise to underdispersed count distributions.



D , the number of latent Poissons, controls the index of dispersion $\mathbb{D}[X]$.

As D grows larger, $\mathbb{D}[X]$ decreases as long as μ is sufficiently large.

Data Augmentation Connects Inference to the Poisson

We derive efficient MCMC-based inference for models with Poisson order statistic likelihoods via a data augmentation scheme with updates of the following form

$$Z_1, \dots, Z_D \sim P(Z_1, \dots, Z_D | \mu, Y) \quad (1)$$

$$\mu \sim P(\mu | Z_*, Y), \text{ where } Z_* = \sum_{k=1}^D Z_k \quad (2)$$

With the appropriate prior on μ , sampling from (2) is standard. Sampling from (1) is cumbersome due to the positive probability of exact ties.

Generalizing the Poisson, say $Z_k \sim f_\mu$ and write a truncated distribution as $tf_\mu(a, b)$.

Algorithm 4.1 Sampling from $P(Z_1, \dots, Z_D | \mu, Y_{(j,D)} = y)$

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1: Input:  $y \in \mathbb{R}$ ,  $D \in \mathbb{N}$ ,  $j \in [D]$ ,  $f_\mu$ 
2: Initialize:  $r_1 = r_2 = r_3 = 0$ 
3: for  $k \in \{1, \dots, D\}$  do
4:   Compute  $p_1 = P(Z_k < y | \mu, r_1, r_2, r_3, Y_{(j,D)} = y)$ 
5:    $p_2 = P(Z_k = y | \mu, r_1, r_2, r_3, Y_{(j,D)} = y)$ 
6:    $p_3 = P(Z_k > y | \mu, r_1, r_2, r_3, Y_{(j,D)} = y)$ 
7:   Sample  $c_k \sim Cat(p_1, p_2, p_3)$ .
8:   Set  $r_{c_k} = r_{c_k} + 1$ 
9:   if  $r_2 \geq 1$  and  $j - r_1 = 1$  and  $k < D$  then
10:     $Z_{k+1}, \dots, Z_D \sim tf_\mu(y, \infty)$  and break
11:   end if
12:   if  $r_2 \geq 1$  and  $D - r_3 = j$  and  $k < D$  then
13:     $Z_{k+1}, \dots, Z_D \sim tf_\mu(0, y)$  and break
14:   end if
15:   if  $r_2 \geq j - r_1$  and  $r_2 \geq D - r_3 - j + 1$  and  $k < D$  then
16:     $Z_{k+1}, \dots, Z_D \sim f_\mu$  and break
17:   end if
18:   end for
19:  $Z_1, \dots, Z_{r_1} \sim tf_\mu(0, y - 1)$ ,  $Z_{r_1+1}, \dots, Z_{r_1+r_2} = y$ ,  $Z_{r_1+r_2+1}, \dots, Z_k \sim tf_\mu(y + 1, \infty)$ 
20: Output:  $\{Z_1, \dots, Z_D\}$ 
```

Using this algorithm, we can exactly sample from (1) for any discrete order statistic Y , including, for example, negative binomial order statistics.

Choosing which Order Statistic

While all Poisson order statistics can be used to model underdispersion, there are differences:

Maximum: computationally efficient, especially under **sparsity**. ($Y = 0 \implies Z_1, \dots, Z_D = 0$)

Median: approximately mean-parameterized by latent μ , but more expensive computationally.

Minimum: computationally efficient, but not under sparsity.

Tailored Models for Applications



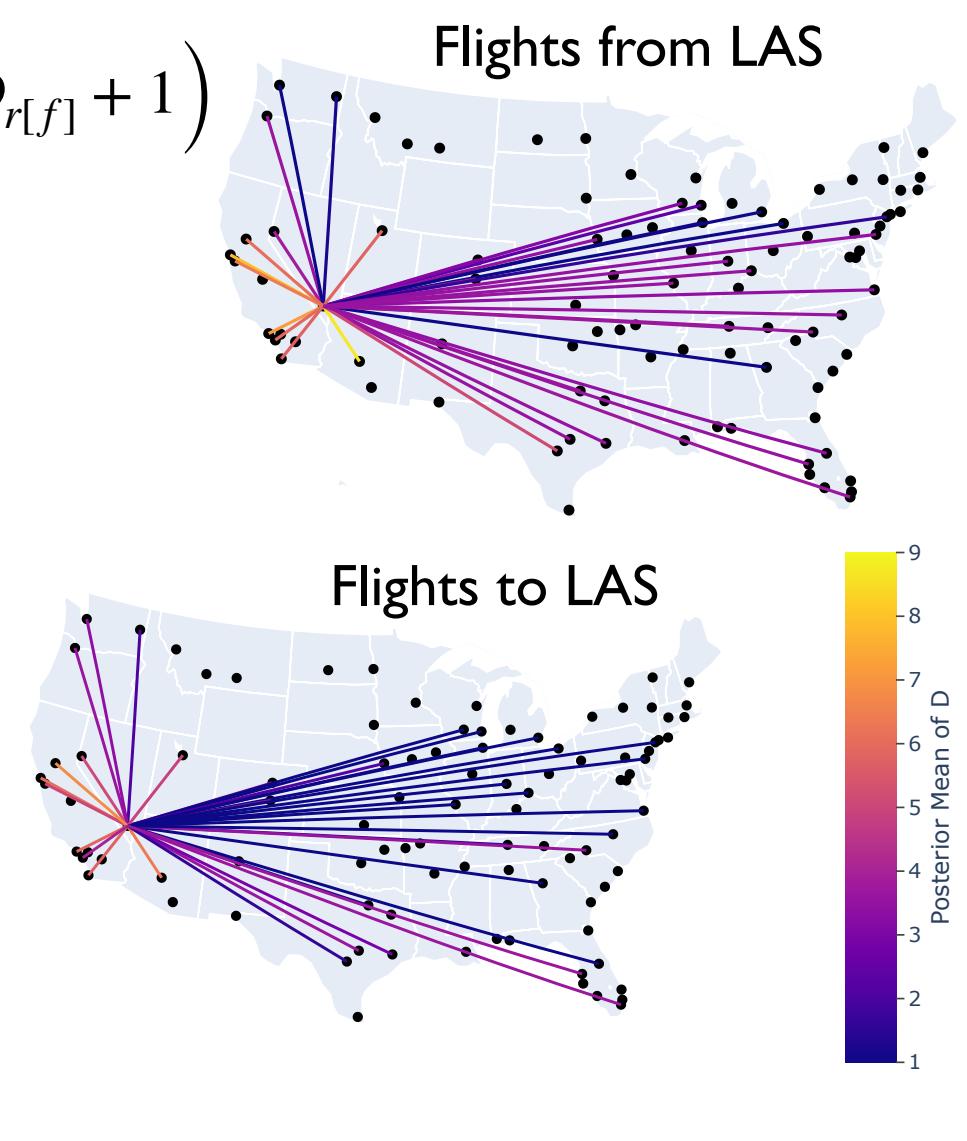
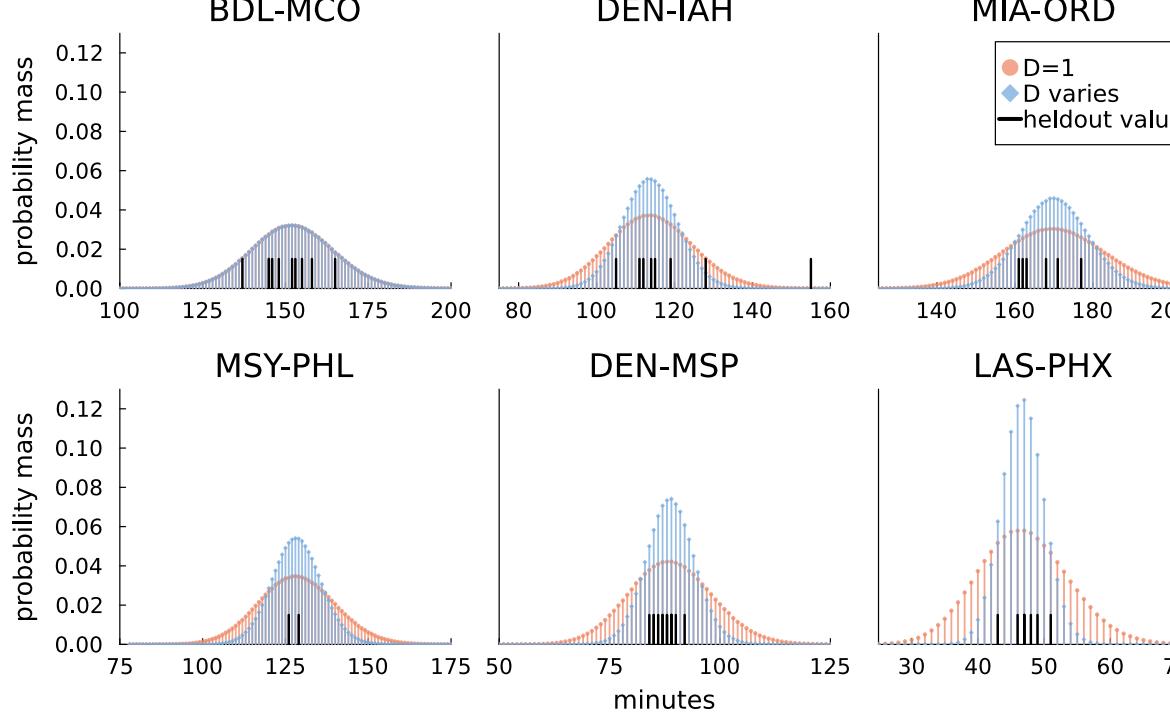
Predicting Flight Times

We model the number of minutes Y_f of flight f with origin $o[f]$, destination $d[f]$, and route $r[f] = (o[f], d[f])$ as

$$Y_f \sim \text{MedPoisson} \left(a_{o[f]} + b_{d[f]} + dist_{r[f]} \mu_{r[f]}, 2D_{r[f]} + 1 \right)$$

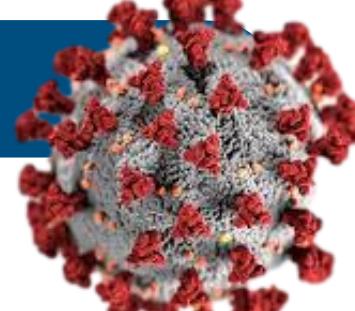
$$D_r \sim \text{Binomial} \left(\frac{D_{\max} - 1}{2}, p \right)$$

Some Example Posterior Predictive Distributions



A flexible dispersion parameter D gives varied uncertainty and learned latent structure.

COVID-19 Forecasting

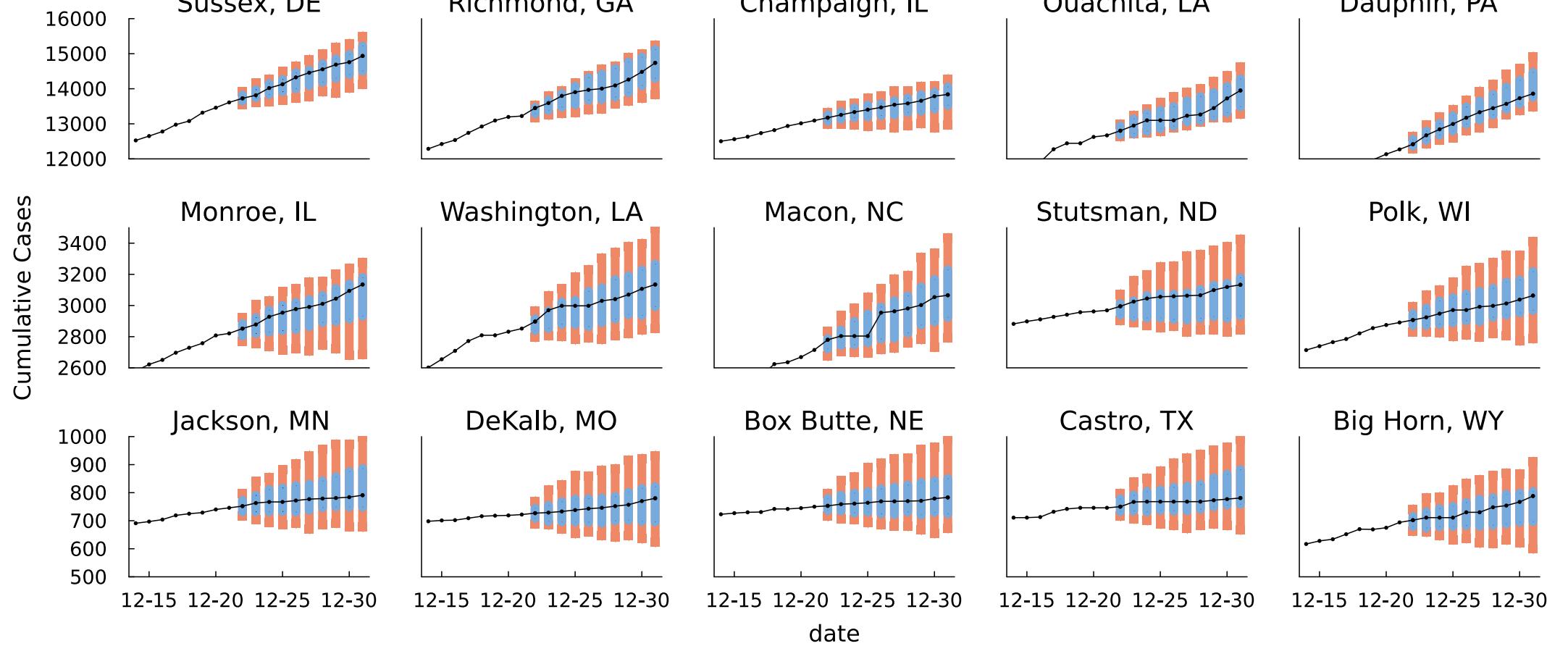


We model the cumulative COVID-19 cases Y_{ct} in each county c at time t as

$$Y_{ct} \sim \text{MedPoisson} \left(Y_{ct-1} + \log(pop_c) \left(\epsilon + \alpha \sum_{k=1}^K \theta_{ck} \phi_{kt} \right), 2D_{ct} + 1 \right)$$

$$D_{ct} \sim \text{Binomial} \left(\frac{D_{\max} - 1}{2}, \sigma \left(\sum_{q=1}^Q \beta_{cq} \tau_{qt} \right) \right)$$

Some Example Posterior Predictive 95% Credible Intervals



The underdispersed likelihood avoids artificially wide predictive intervals.

Modeling Abundance of Finnish Birds

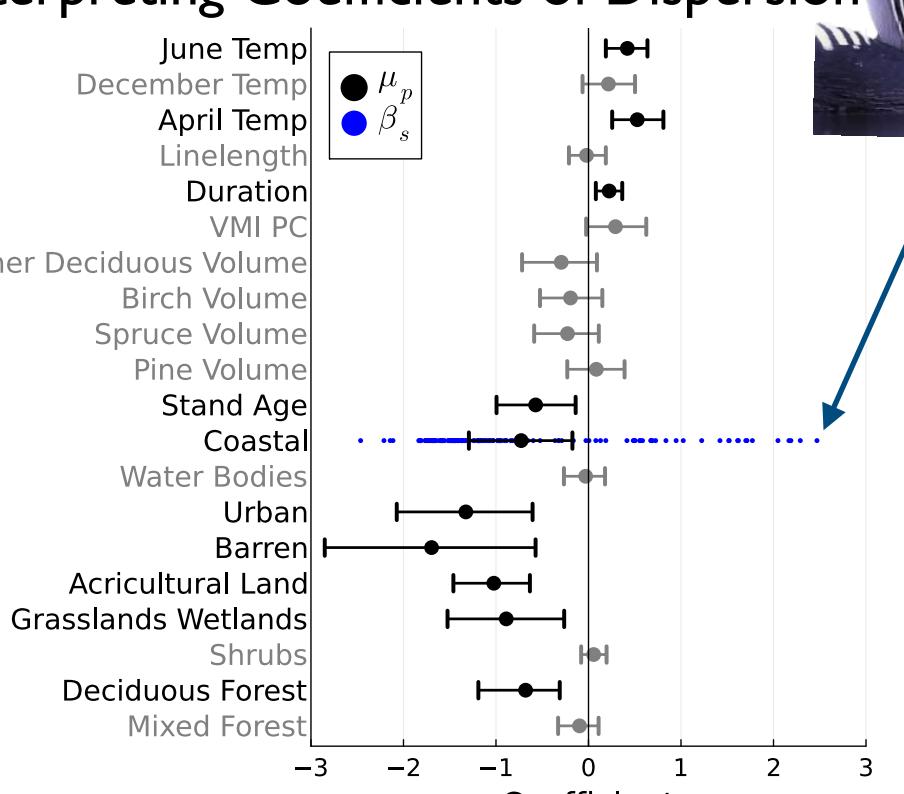


We model the number of birds Y_{ns} at sampling site n of species s , where each sampling site has covariate vector X_n as

$$Y_{ns} \sim \text{MaxPoisson} \left(\sum_{k=1}^K \theta_{ck} \phi_{ks}, D_{ns} + 1 \right)$$

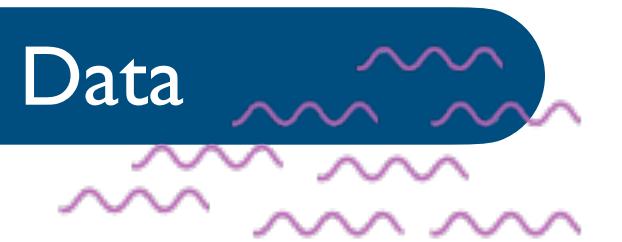
$$D_{ns} \sim \text{Binomial} \left(D_{\max} - 1, \sigma(X_n^T \beta_s) \right)$$

$$\beta_{sp} \sim N(\mu_p, \sigma_p^2) \quad \mu_p \sim N(0, 1)$$



Our model yields more precise predictions for waterbirds in coastal areas.

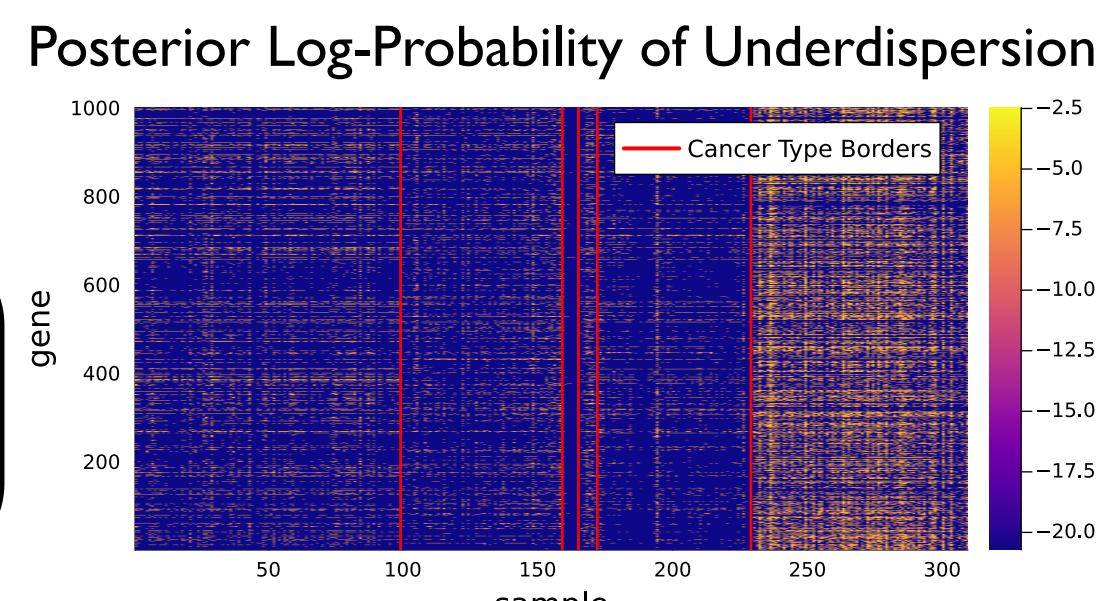
Investigating Dispersion in RNA-Seq Data



We model the RNA-sequencing count Y_{gs} for gene g at sample s as

$$Y_{gs} \sim \text{MedNB} \left(\sum_{k=1}^K \theta_{gk} \phi_{ks}, p_g, 2D_{gs} + 1 \right)$$

$$D_{gs} \sim \text{Binomial} \left(\frac{D_{\max} - 1}{2}, \sigma \left(c_g + \sum_{q=1}^Q \beta_{gq} \tau_{sq} \right) \right)$$



We find little evidence of underdispersion in RNA sequencing data, but there are different patterns across cancer types.

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