# Discovering signals in fMRI data; a Bayesian nonparametric approach

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#### **Outline**

#### Introduction

Method

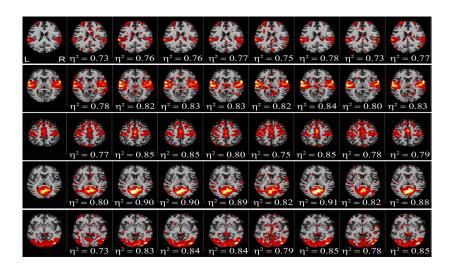
Example: simulated data

Example: fMRI data

#### What is fMRI?

- ▶ fMRI: functional magnetic resonance imaging
- measures the change in brain blood flow while subject is stimulated
- data tends to look like (voxel, time, intensity of reading)
- e.g., to identify regions of the brain associated with hunger, fMRI readings can be taken while hungry subjects are shown pictures of food.

#### **Example**



#### **Problems**

- multiple comparison problem thousands of readings over time and space
- want to incorporate location and time information want continuous significant regions, not voxels
- want to determine these significant regions adaptively

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#### The idea

- ▶ focus on data with p-values rather than intensities
- assume that data is generated according to a prior which induces clustering of significant p-values
- get the posterior distribution (somehow)
- p-values which are in significant clusters are discoveries

## The prior we assumed

- ▶ number of signal clusters:  $k \sim \text{Truncated Poisson}(\lambda, 1, k_{max})$
- lacktriangle signal centers:  $c_j \sim \mathsf{Uniform}$  over location-time space for j=1,...,k
- ▶ signal radius:  $r_j \sim \mathsf{Truncated} \; \mathsf{Normal}(\mu, \sigma, r_{min}, r_{max}) \; \mathsf{for} \; \mathsf{j} = \mathsf{1}, \ldots \mathsf{k}.$
- ▶ signal strength:  $\beta_j \sim \mathsf{Uniform}(\beta_{min}, \beta_{max})$  for  $j = 1, \ldots, k$ .
- ▶ p-values in signal clusters:  $p_i \sim \text{Beta}(\frac{1}{\beta_i}, \beta_j)$ , when  $x_i$  is in cluster j.
- ightharpoonup p-values not in signal clusters:  $p_i \sim \mathsf{Uniform}(0,1)$ .

## How do we write down the posterior??

- ▶ we don't we sample from it
- ▶ we invented a Markov chain, inspired by Stephens (2000), with posterior distribution given by the likelihood of the data
- the type of chain is a birth-death chain (over an infinite dimensional state space!)

## The rough algorithm

- ▶ initialize clusters and labels randomly by sampling from the prior
- ▶ repeat the following 10,000 times
  - flip a weighted coin to determine if a birth or death occurs
  - if birth, add a new cluster randomly
  - if death, delete a cluster which doesn't explain the data well
- ▶ take the average of the last 5,000 labels to determine if a p-value is signal or null.

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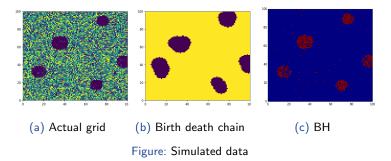
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## **Toy Data: Preliminaries**

- ▶ 100-by-100 grid with k=5 clusters of signals and the rest is null.
- ▶ the centers  $C_k \sim$  uniform from the grid while being distinct for k=1,2,...,5
- ▶ the radius  $R_k \sim TN(7, 2, 5, 10)$  for k = 1, 2, ..., 5
- ▶ signals in clusters  $p_{ki} \sim Beta(1, \beta_k)$  where  $\beta_k \sim TN(8, 5, 2, 200)$  for k = 1, 2, ..., 5

# Toy Data (continued): Performance



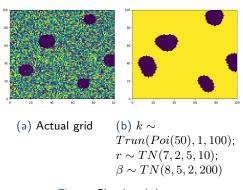


Figure: Simulated data

First, let's compare different priors on beta

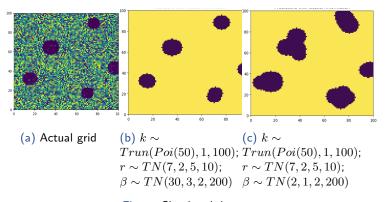


Figure: Simulated data

Now, let's compare different priors on radius.

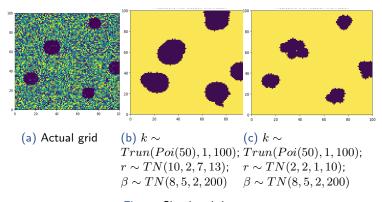


Figure: Simulated data

What if we make signal stronger?

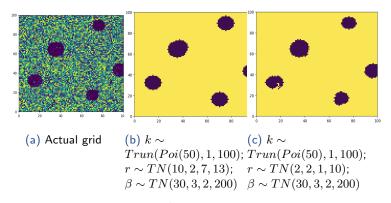


Figure: Simulated data

Next, we change the priors on number of clusters.

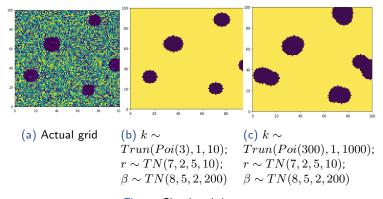


Figure: Simulated data

Again, let's make signal stronger

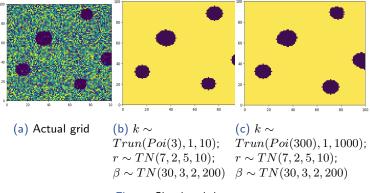


Figure: Simulated data

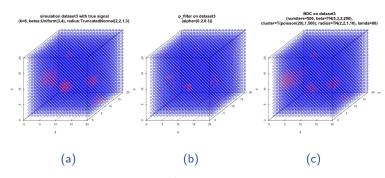


Figure: Simulated data

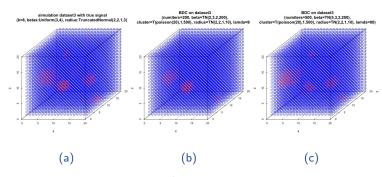


Figure: Simulated data

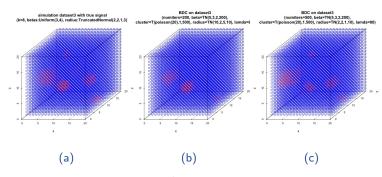


Figure: Simulated data

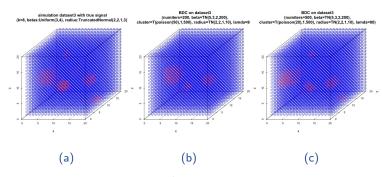


Figure: Simulated data

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#### Performance on real fMRI data

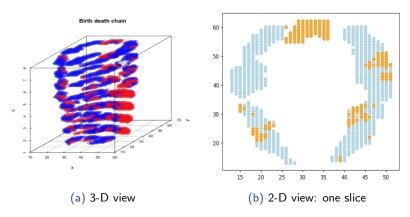
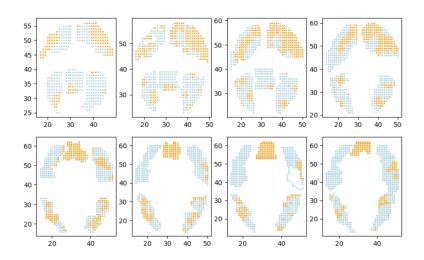


Figure: fMRI data

# Performance on real fMRI data (continued)



# Performance on real fMRI data (continued): Comparison to p-filter

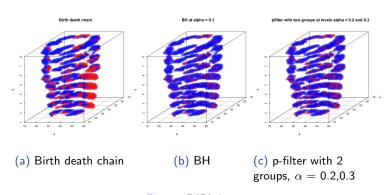


Figure: fMRI data

#### **Conclusion and Future Work**

- presented a nonparametric-bayesian method to adaptively identify clusters of signals.
- ▶ it showed promising results on both simulation and real fMRI data.
- possible extensions
  - work with intensities directly by specifying appropriate priors for null distributions and for signal distributions.
  - put priors on the hyper-parameters and maximize these priors using EM

#### Thanks!