

Beta diffusion trees

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Joint work with

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Outline

We are interested in random tree structures on overlapping clusters
(aka feature allocations).

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1. Briefly review feature allocations.

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2. Describe a stochastic process (the beta diffusion tree) that leads to trees over feature allocations.

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3. Describe a hierarchically-clustered factor analysis model.

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1. Briefly review feature allocations.
2. Describe a stochastic process (the beta diffusion tree) that leads to trees over feature allocations.
3. Describe a hierarchically-clustered factor analysis model.
4. Present some inference techniques over the tree structure and some experimental results.

Feature allocations

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Feature allocations = Overlapping subsets = Overlapping clusters

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Consider an image labelling application:



Feature allocations

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Consider an image labelling application:



“image contains
a chair”

“image contains
a table”

“image is in
a kitchen”

Feature allocations

Feature allocations = Overlapping subsets = Overlapping clusters

Consider an image labelling application:



“image contains
a chair”

{1}

“image contains
a table”

{1}

“image is in
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{1}

Feature allocations

Feature allocations = Overlapping subsets = Overlapping clusters

Consider an image labelling application:



“image contains
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$\{1\}$

“image contains
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$\{1\}$

“image is in
a kitchen”

$\{1\}$

Feature allocations

Feature allocations = Overlapping subsets = Overlapping clusters

Consider an image labelling application:



"image contains
a chair"

$\{1, 2\}$

"image contains
a table"

$\{1, 2\}$

"image is in
a kitchen"

$\{1\}$

Feature allocations

Feature allocations = Overlapping subsets = Overlapping clusters

Consider an image labelling application:

Feature allocation:

$\{\{1, 2\}, \{1, 2\}, \{1\}\}$



Feature allocations

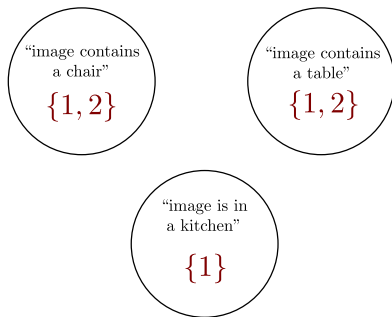
Feature allocations = Overlapping subsets = Overlapping clusters

Consider an image labelling application:

Feature allocation:

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The clusters/subsets are called
features



Hierarchies on features

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Hierarchies of features are a natural assumption:

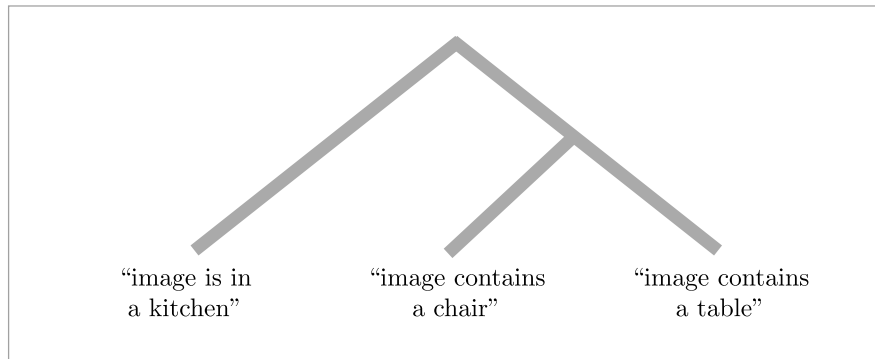
“image is in
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Hierarchies on features

Hierarchies of features are a natural assumption:



The beta diffusion tree

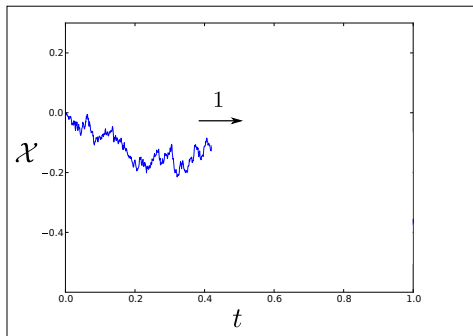
The beta diffusion tree

Consider a sequence of particles, labeled with one of N objects, diffusing in a continuous space \mathcal{X} over the unit time interval $[0, 1]$.

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For example:



- ▶ $\mathcal{X} = \mathbb{R}$.
- ▶ Diffusion is Brownian motion.

The beta diffusion tree

Consider a sequence of particles, labeled with one of N objects, diffusing in a continuous space \mathcal{X} over the unit time interval $[0, 1]$.

The beta diffusion tree

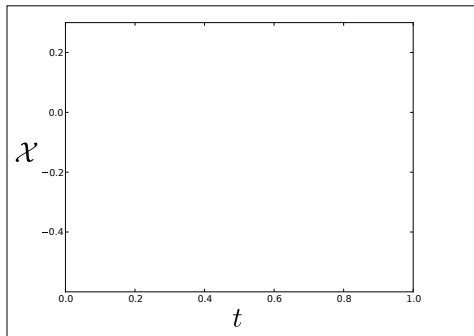
Consider a sequence of particles, labeled with one of N objects, diffusing in a continuous space \mathcal{X} over the unit time interval $[0, 1]$.

At random times:

- ▶ **Replicate**: Copies of particles may be created.
- ▶ **Stop**: Particles are destroyed.

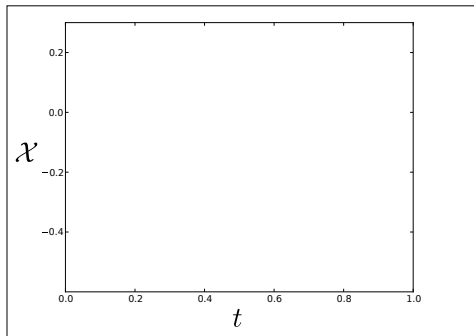
The beta diffusion tree

Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.



The beta diffusion tree

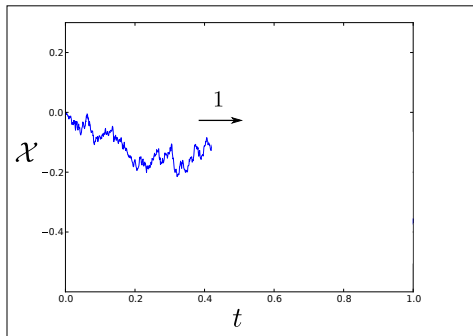
Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.



Consider the first object.

The beta diffusion tree

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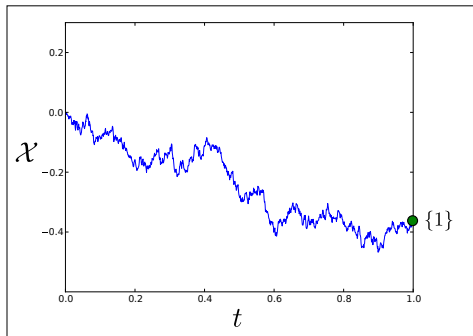


1. The first particle starts at the origin.
2. May replicate or stop after $\exp(\lambda_r)$ or $\exp(\lambda_s)$ time.

Consider the first object.

The beta diffusion tree

Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.

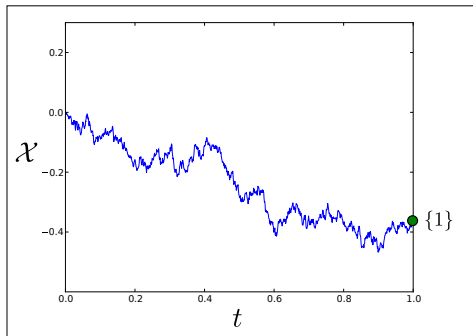


1. The first particle starts at the origin.
2. May replicate or stop after $\exp(\lambda_r)$ or $\exp(\lambda_s)$ time.
3. Diffuses as Brownian motion until $t = 1$.

Consider the first object.

The beta diffusion tree

Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.

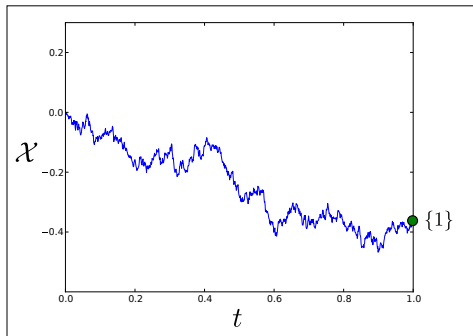


1. Follow previous particles.

Consider the second object.

The beta diffusion tree

Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.



1. Follow previous particles.
2. May replicate or stop after

$$\exp\left(\frac{\theta_r}{\theta_r + m} \lambda_r\right)$$

or

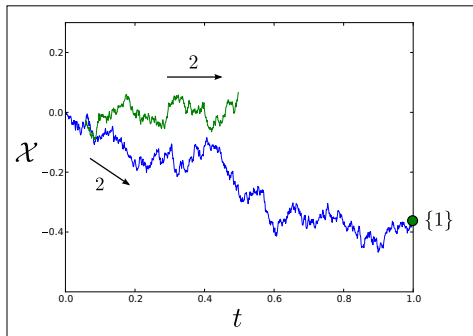
$$\exp\left(\frac{\theta_s}{\theta_s + m} \lambda_s\right)$$

amount of time.

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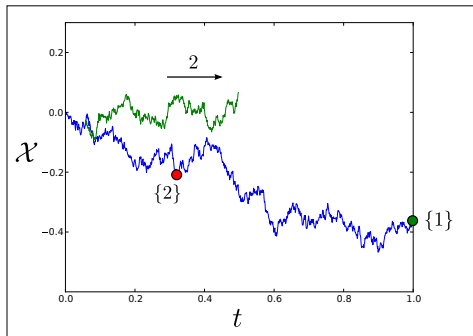
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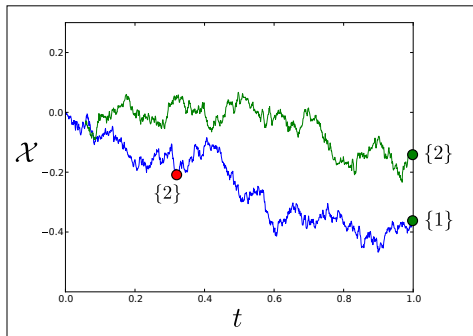
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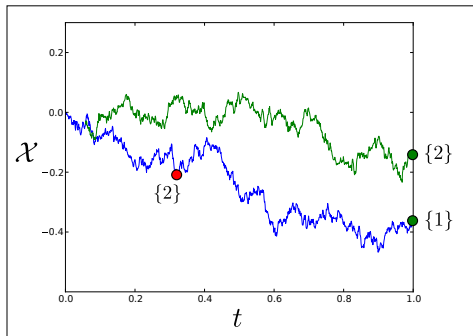
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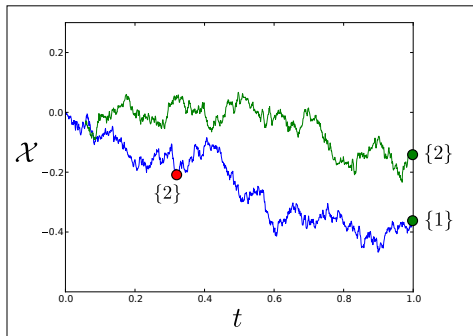
$$\exp\left(\frac{\theta_s}{\theta_s + m} \lambda_s\right)$$

amount of time.

Consider the third object.

The beta diffusion tree

Let $\lambda_s, \lambda_r, \theta_s, \theta_r > 0$ and $\mathcal{X} = \mathbb{R}$. Diffusion is Brownian motion.



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2. May replicate or stop after

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or

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amount of time.

3. Replicate and stop w.p.

$$\frac{n_r}{\theta_r + m}$$

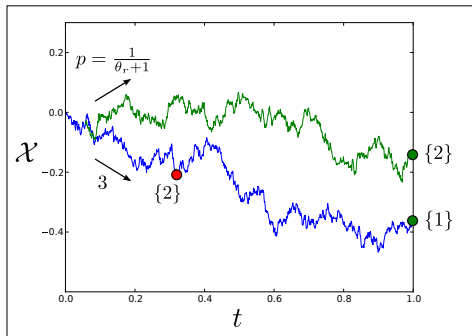
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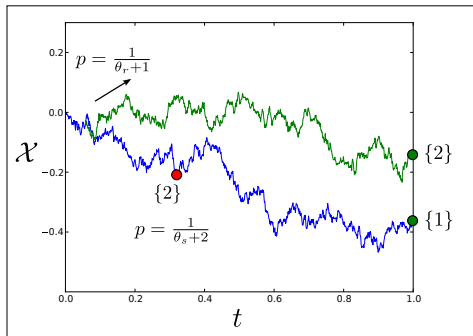
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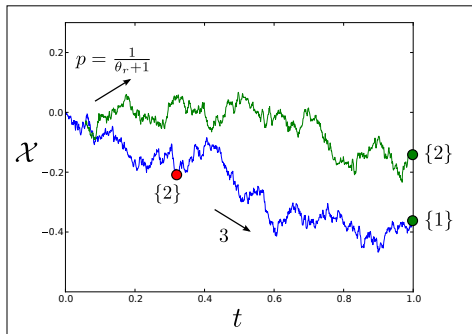
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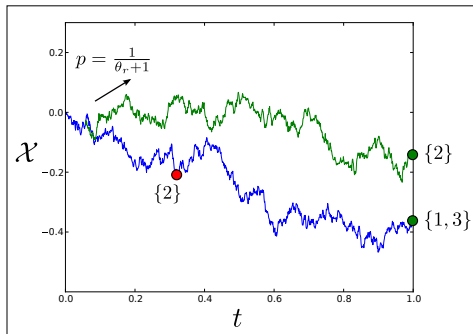
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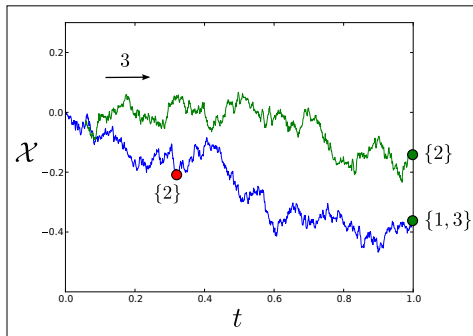
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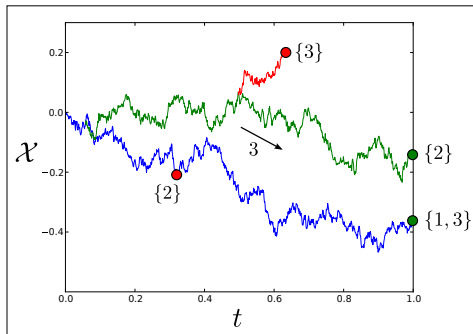
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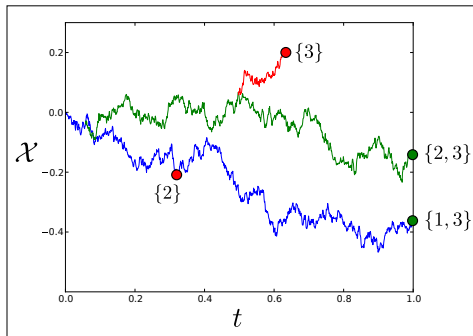
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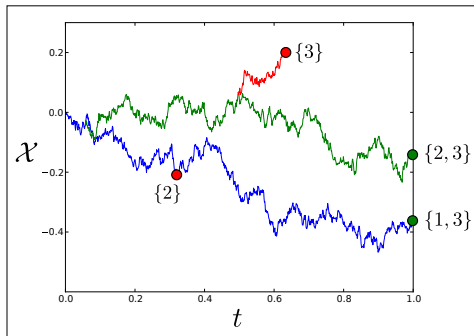
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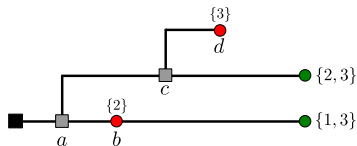
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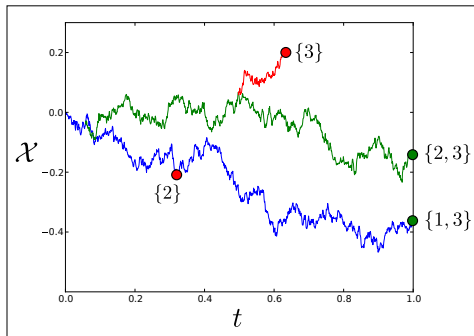


The corresponding
tree structure:

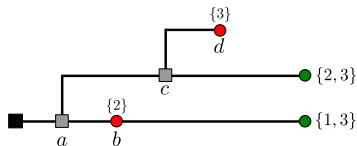


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The corresponding
tree structure:



The leaves form a feature allocation of $[3]$:

$$\{\{1, 3\}, \{2, 3\}\}$$

Related work

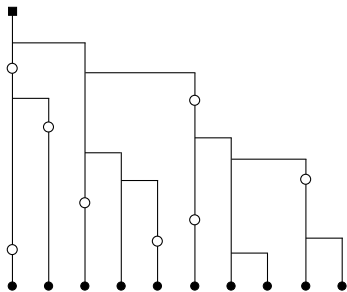
As opposed to...

Random tree structures over **non-overlapping clusters**
(aka partitions):

- ▶ Dirichlet diffusion tree (Neal, 2003 [6])
- ▶ Pitman–Yor diffusion tree (Knowles & Ghahramani, 2011 [5])
- ▶ Kingman's coalescent (Kingman, 1982 [4])

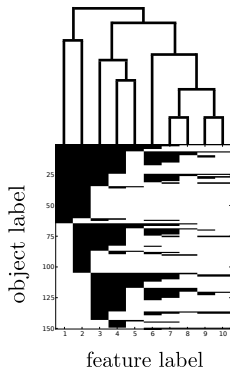
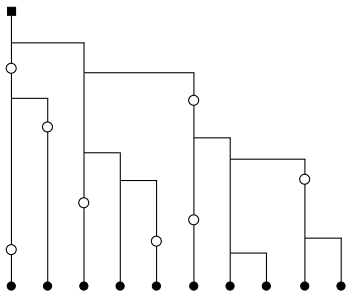
The beta diffusion tree

Consider the following larger example:



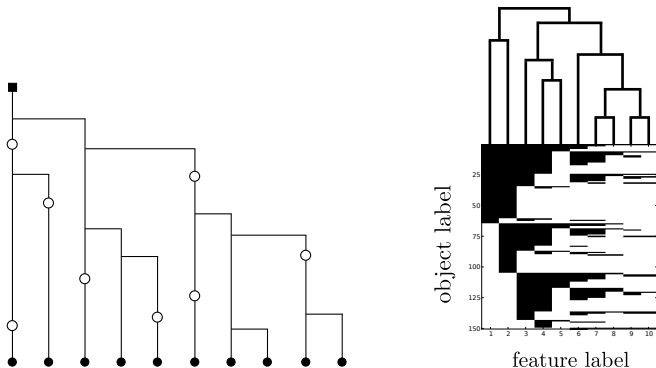
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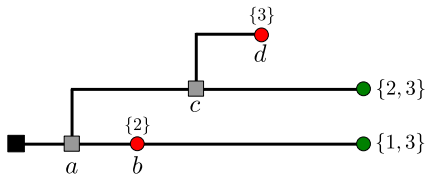
Consider the following larger example:



A tree structure over the columns of a binary matrix.

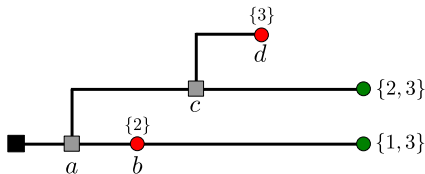
Connections to the Indian buffet process
(Ghahramani, Griffiths, & Sollich, 2007 [3]).

The beta diffusion tree



We defined the tree in terms of the ordered sequence of objects,
however:

The beta diffusion tree



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however:

Theorem (exchangeability)

The beta diffusion tree is exchangeable with respect to the ordering of the objects.

Other properties

- ▶ The beta diffusion tree is the continuum limit of a nested feature allocation scheme
- ▶ The beta diffusion tree is a continuous-time branching process

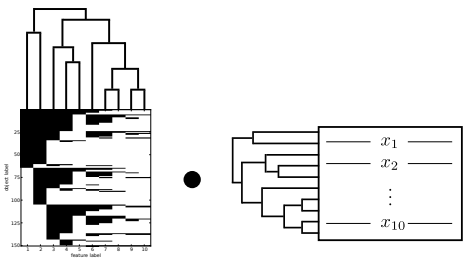
Example: A hierarchical factor analysis model

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$$\underbrace{Y}_{\substack{N \times D \\ \text{Observations}}} = \underbrace{Z}_{\substack{N \times K \\ \text{factors} \\ \text{= feature allocation}}} \bullet \underbrace{X}_{\substack{K \times D \\ \text{factor loadings} \\ \text{= leaves in } \mathcal{X}}} + \underbrace{\varepsilon}_{\substack{N \times D \\ \text{i.i.d. Gaussian} \\ \text{noise}}}$$

The diagram illustrates a hierarchical factor analysis model. The equation $Y = ZX + \varepsilon$ is shown, where Y is the matrix of observations ($N \times D$), Z is the matrix of factors ($N \times K$), X is the matrix of factor loadings ($K \times D$), and ε is the matrix of i.i.d. Gaussian noise ($N \times D$). The matrix Z is visualized as a heatmap with a dendrogram on top, representing feature allocation. The matrix X is visualized as a box with a dendrogram on the left and a list of leaves x_1, x_2, \dots, x_{10} on the right, representing leaves in \mathcal{X} .

Example: A hierarchical factor analysis model

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Both the factors and factor loadings are hierarchically structured

Inference

We perform a series of Metropolis–Hastings proposals to integrate over the random tree structures.

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1. Resample the paths of particles down the tree.

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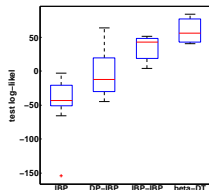
Inference

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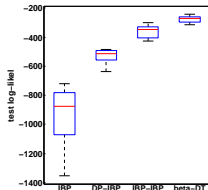
Each is accepted or rejected with a Metropolis–Hastings step.

Some numerical results

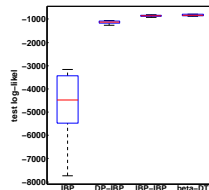
Test log-likelihoods:



(a) *E. Coli*



(b) UN



(c) India

- (a) A gene expression dataset
- (b) International development statistics
- (c) Household socioeconomic indicators

- [1] F. Doshi-Velez and Z. Ghahramani. Correlated non-parametric latent feature models. In Proc. UAI, 2009.
- [2] J. Geweke. Getting it right: Joint distribution tests of posterior simulators. Journal of the American Statistical Association, 99(467):799–804, 2004.
- [3] Z. Ghahramani, T. L. Griffiths, and P. Sollich. Bayesian nonparametric latent feature models. Bayesian Statistics, 8:201–226, 2007. See also the discussion and rejoinder.
- [4] J. F. C. Kingman. The coalescent. Stochastic Processes and Their Applications, 13(3):235–248, 1982.
- [5] D. Knowles and Z. Ghahramani. Pitman-Yor diffusion trees. In Proc. UAI, 2011.
- [6] R. M. Neal. Density modeling and clustering using Dirichlet diffusion trees. Bayesian Statistics, 7, 2003.