



Posteriors, conjugacy, and exponential families for completely random measures

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- Gamma process, Poisson likelihood process (DP, CRP)

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$$p(x|\theta) = \theta^x (1+\theta)^{-1} \qquad x \in \{0,1\}$$
 $\theta > 0$

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$$p(x|\theta) = \theta^x (1+\theta)^{-1} \qquad x \in \{0,1\}$$

$$\theta > 0$$

$$p(\theta) \propto \theta^{\alpha} (1+\theta)^{-\alpha-\beta} = \text{BetaPrime}(\theta|\alpha,\beta)$$

$$\alpha > 0, \beta > 0$$

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$$p(x|\theta) = \theta^{x} (1+\theta)^{-1} \qquad x \in \{0,1\} \qquad \theta > 0$$
$$p(\theta) \propto \theta^{\alpha} (1+\theta)^{-\alpha-\beta} = \text{BetaPrime}(\theta|\alpha,\beta) \qquad \alpha > 0, \beta > 0$$
$$p(\theta|x) \propto \theta^{\alpha+x} (1+\theta)^{-(\alpha+x)-(\beta-x+1)}$$

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Background

- Parametric exponential family conjugacy [Diaconis & Ylvisaker 1979]
 - Likelihood → conjugate prior, straightforward inference

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Background

- Parametric exponential family conjugacy [Diaconis & Ylvisaker 1979]
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 - Integration → addition

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Want: One framework

• For Bayesian *nonparametric* models:

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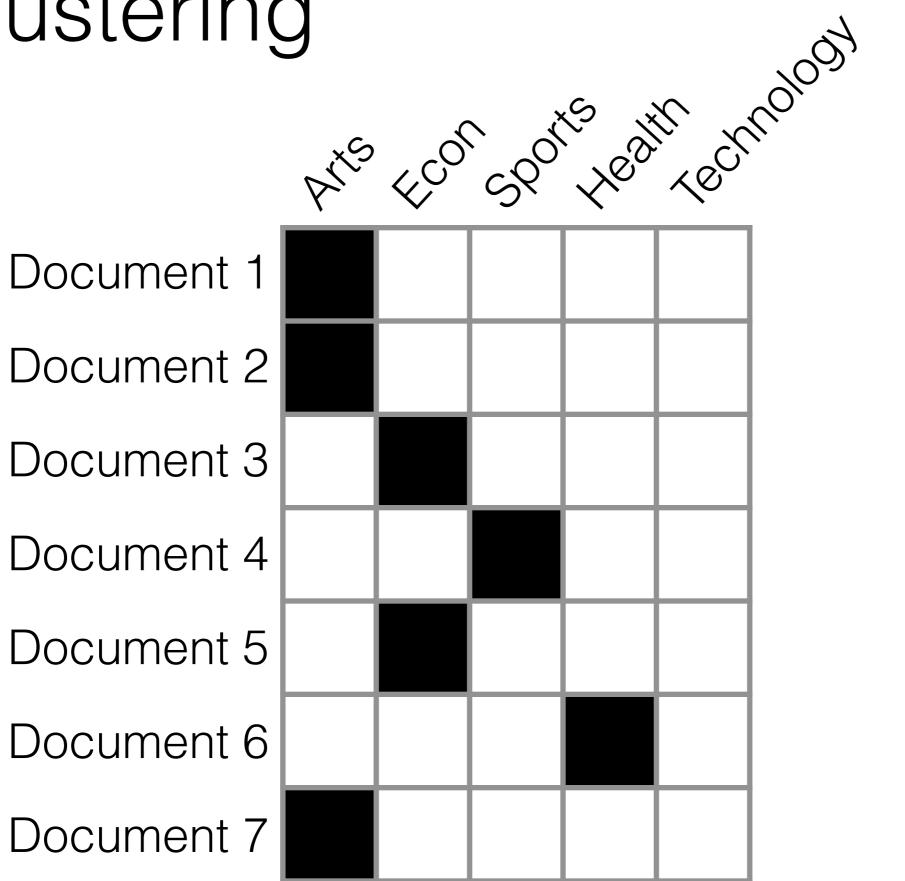
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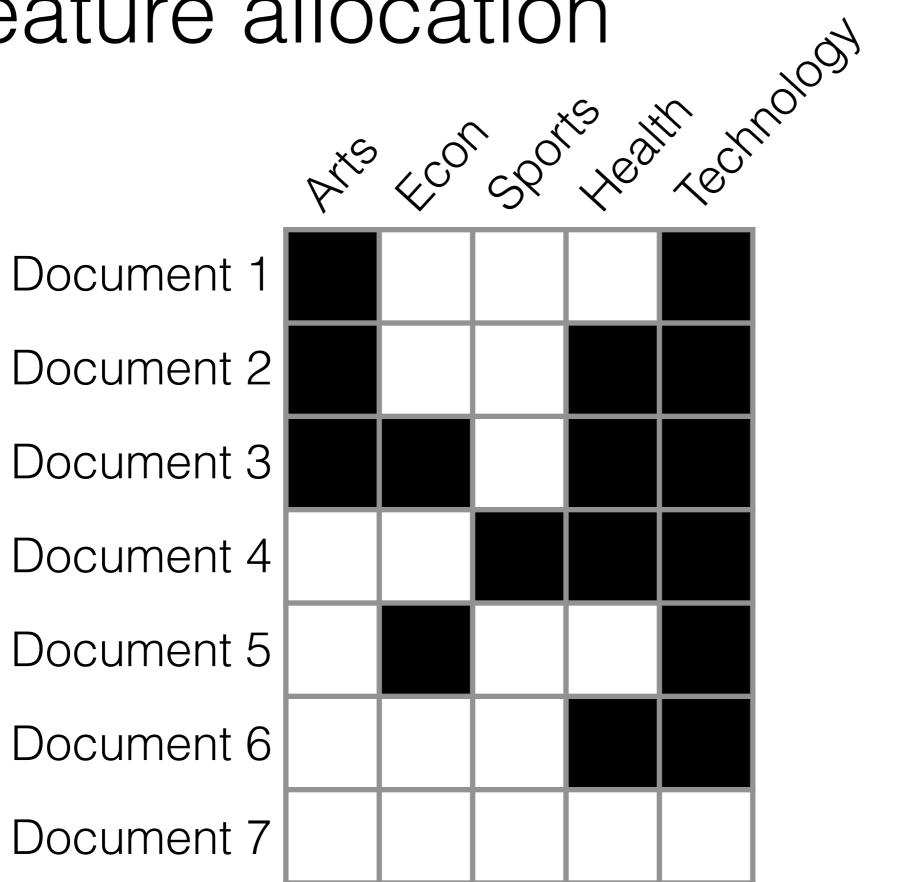
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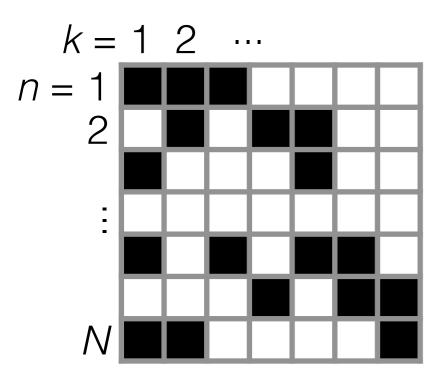
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Clustering



Feature allocation

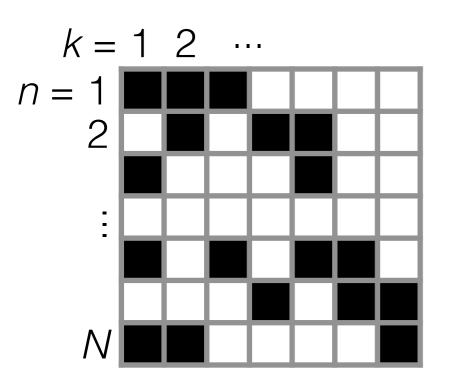




For
$$n = 1, 2, ..., N$$

1.

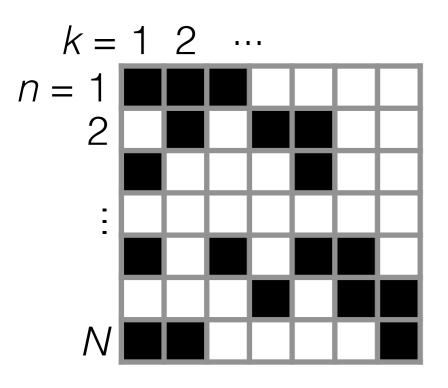
2.



For n = 1, 2, ..., N

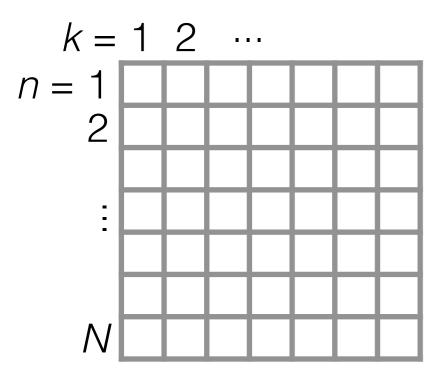
1. Data point n has an existing feature k that has occurred $S_{n-1,k}$ times with probability $S_{n-1,k}$ $\beta + n - 1$

2.



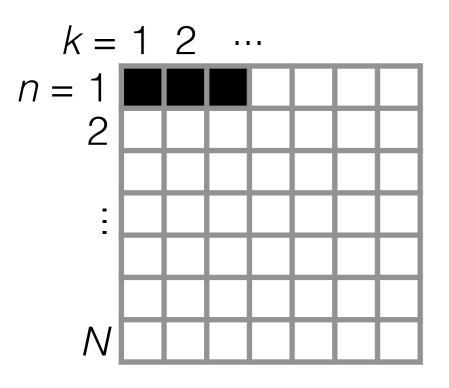
- 1. Data point n has an existing feature k that has occurred $S_{n-1,k}$ times with probability $S_{n-1,k}$ $\beta + n 1$
- 2. Number of new features for data point n:

$$K_n^+ = \text{Poisson}\left(\gamma \frac{\beta}{\beta + n - 1}\right)$$



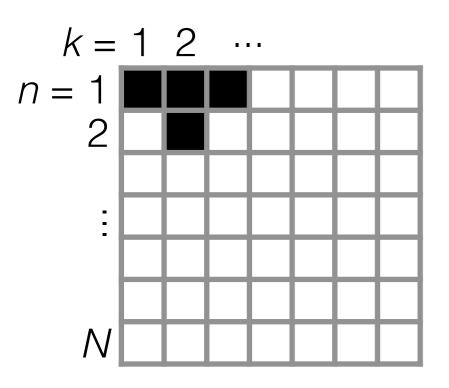
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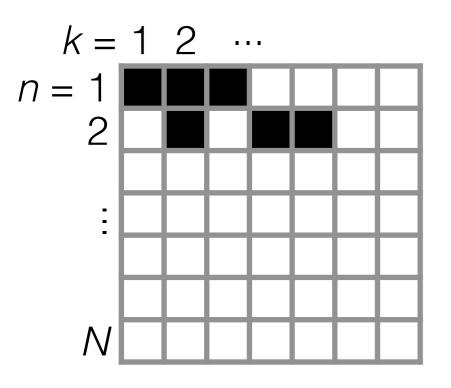
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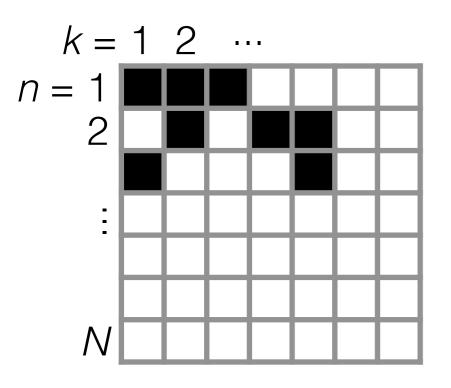
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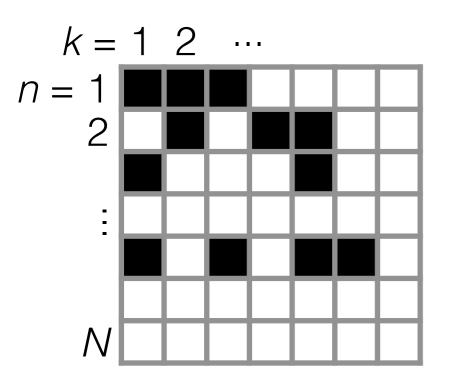
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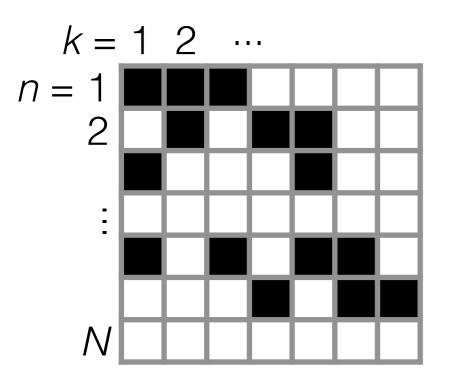
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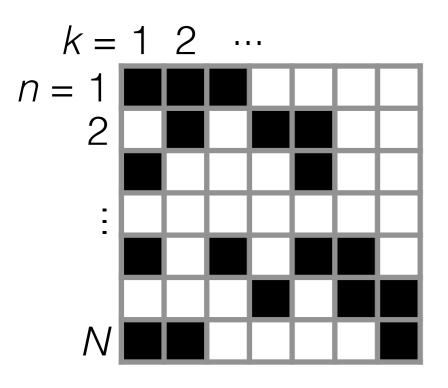
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Draw a frequency of size $\theta_k \sim \text{Beta}(1, \beta + m - 1)$

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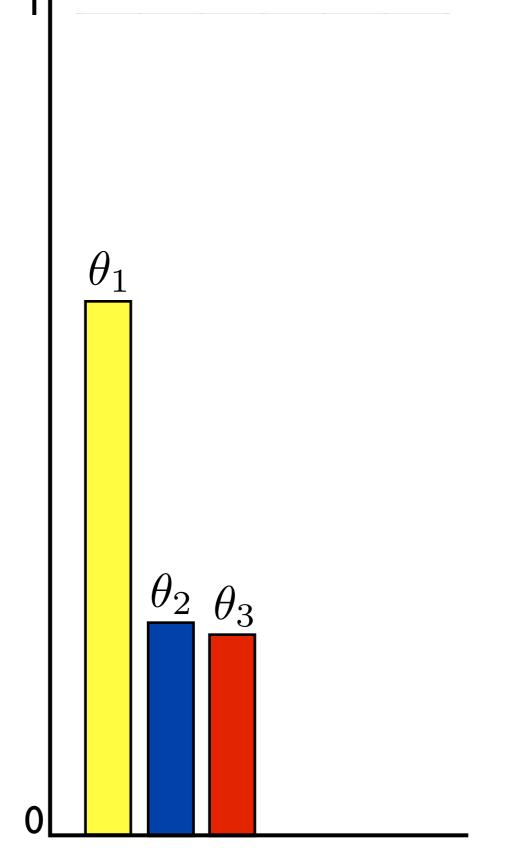
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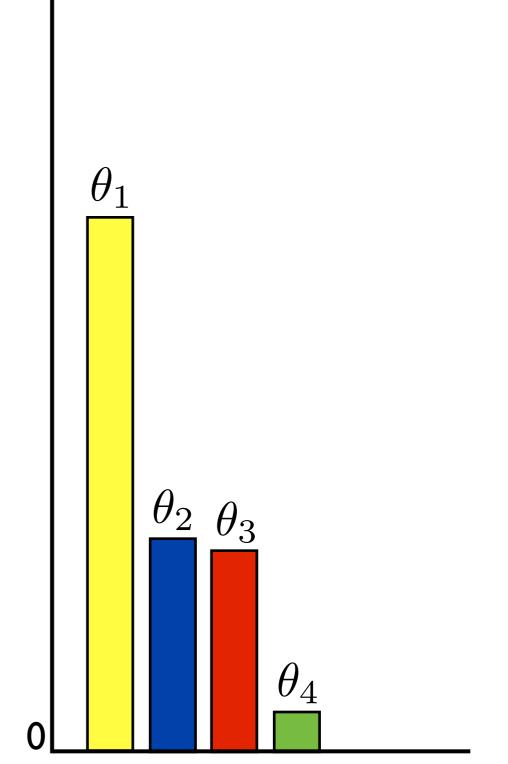
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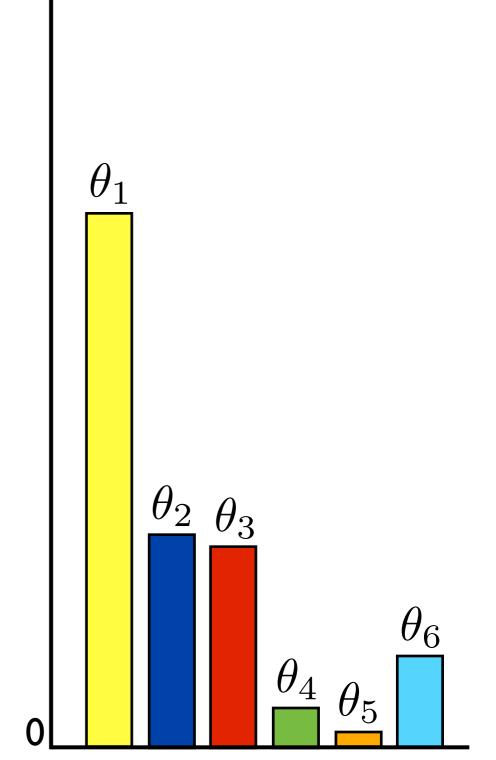
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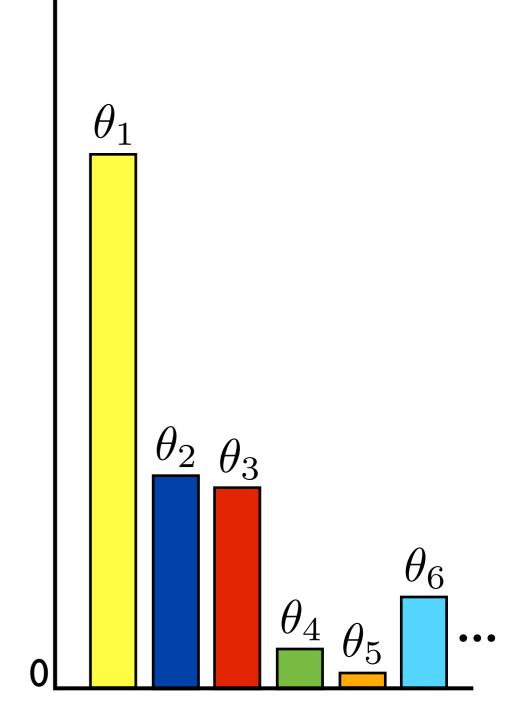
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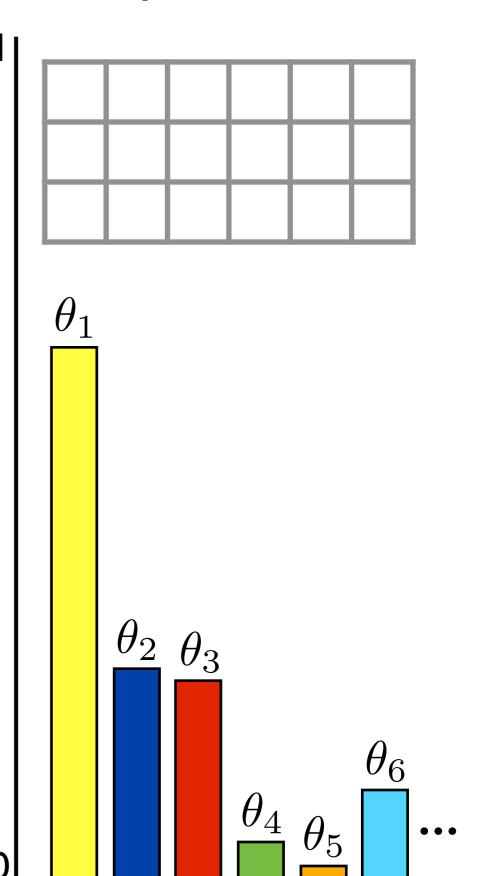
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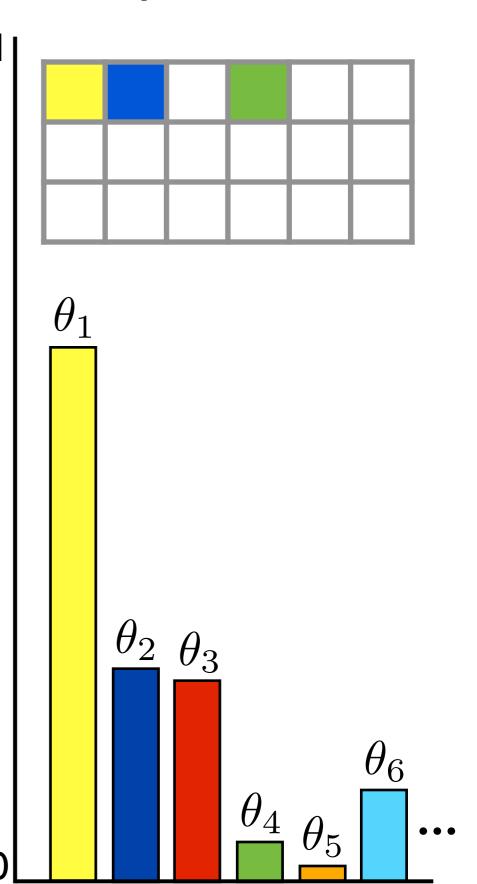
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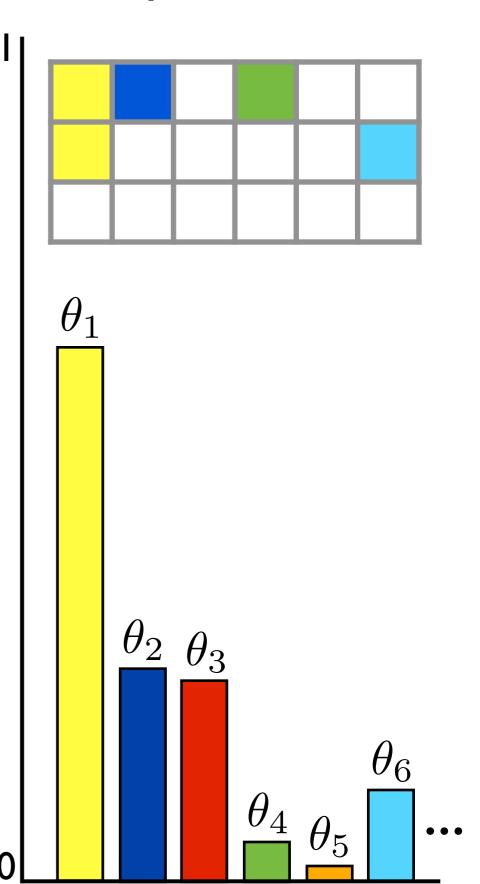
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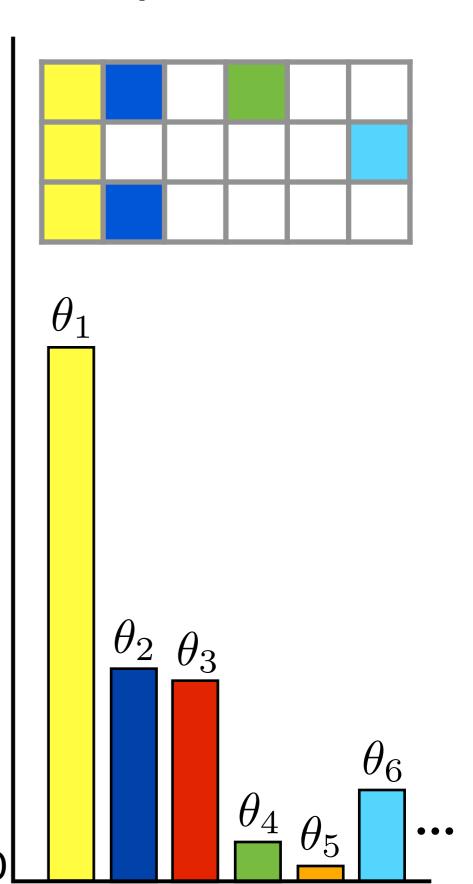
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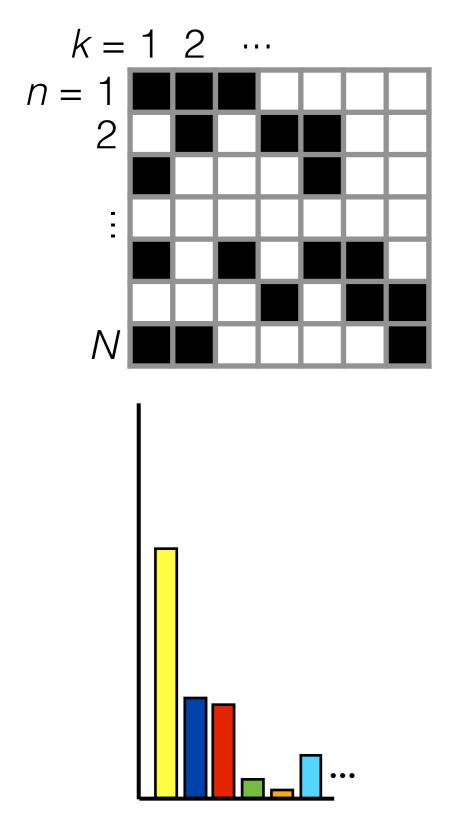


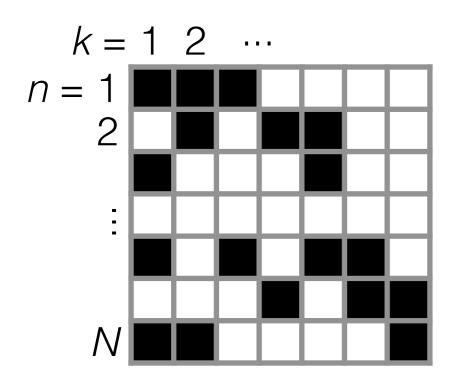
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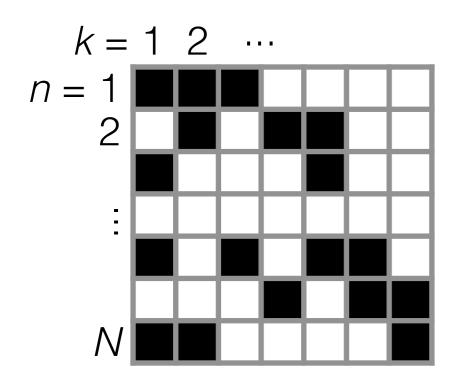
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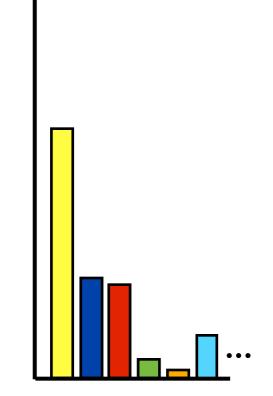


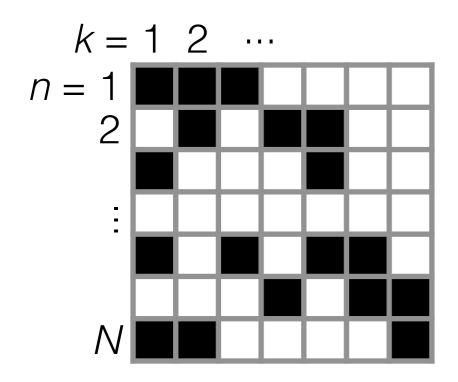




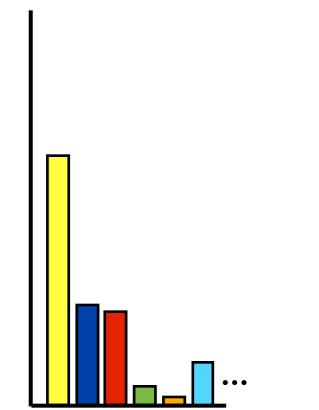


- Exchangeable (e.g.
 Gibbs sampling)
- Finite but unbounded

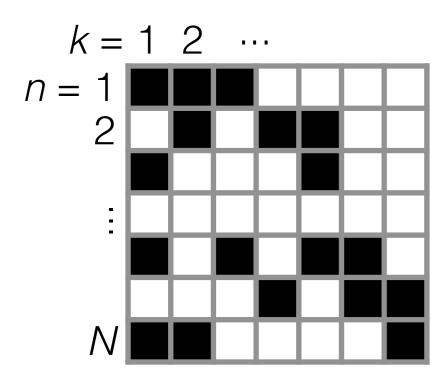


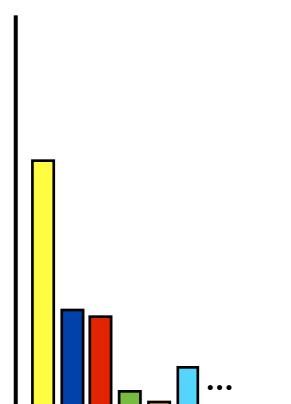


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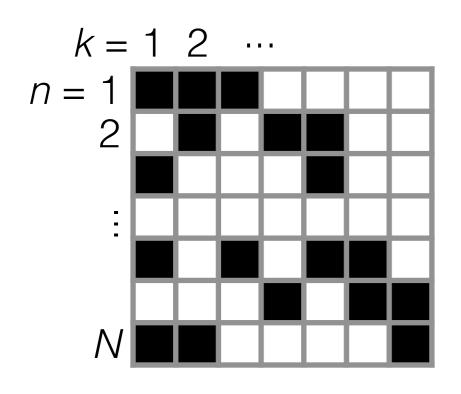
Hierarchical models





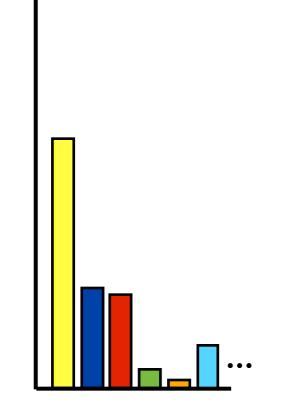
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- Hierarchical models
- (Countable) sequence of finite-dimensional distributions



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How do we come up with these models?

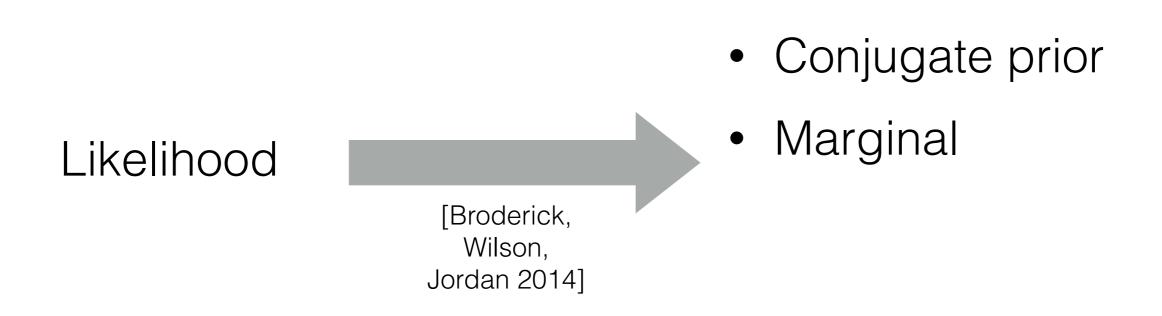
Likelihood

[Broderick,
Wilson,
Jordan 2014]

• Conjugate prior

Likelihood

[Broderick,
Wilson,
Jordan 2014]



Conjugate prior
Likelihood
Marginal
Size-biased atom sequence

Likelihood (e.g. Bernoulli)

- Conjugate prior
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Likelihood (e.g. Bernoulli)

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Likelihood (e.g. Bernoulli)

- Conjugate prior (e.g. BP)
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- Size-biased atom sequence (e.g. BP stickbreaking)

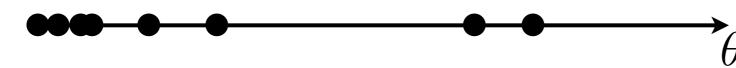
$$p(x|\theta) = \theta^x (1+\theta)^{-1}$$
 $x \in \{0,1\}$ $\theta > 0$

$$\theta > 0$$

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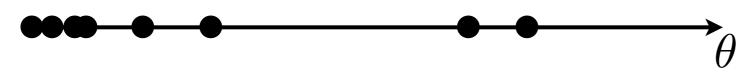
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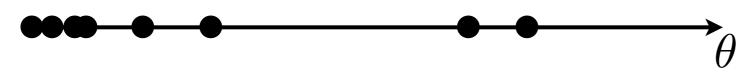
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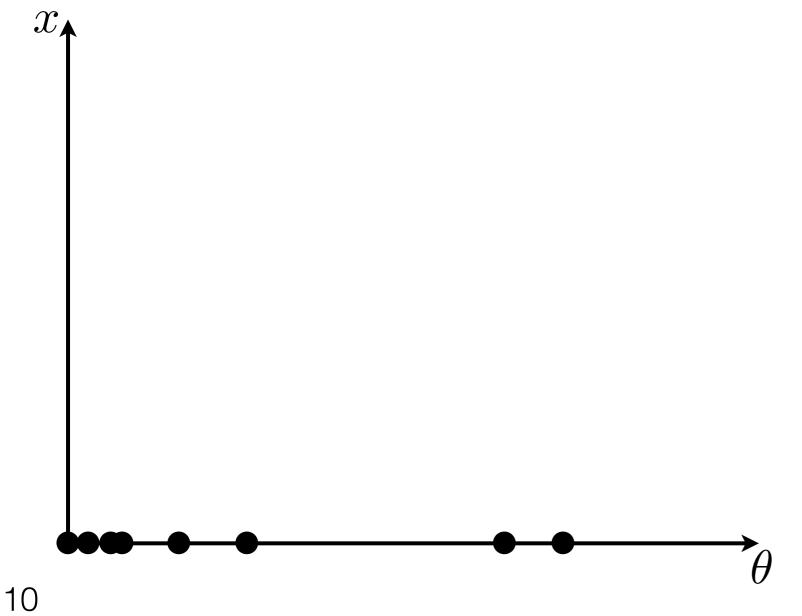
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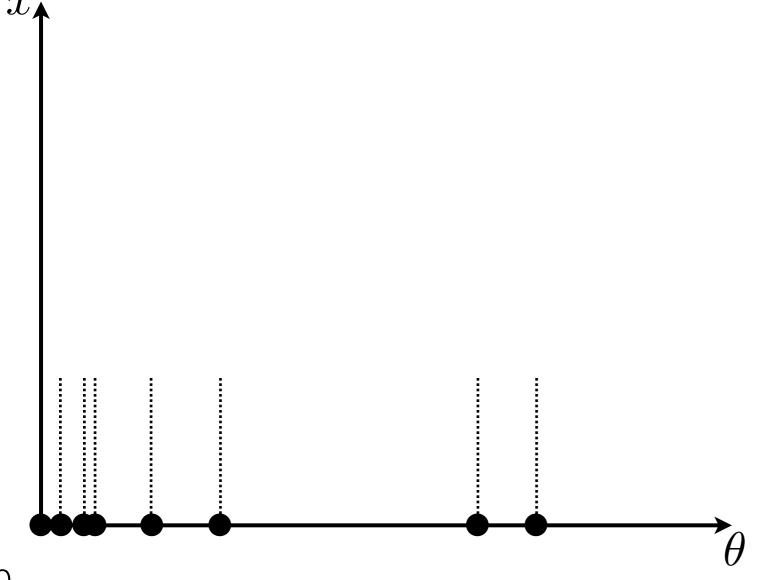
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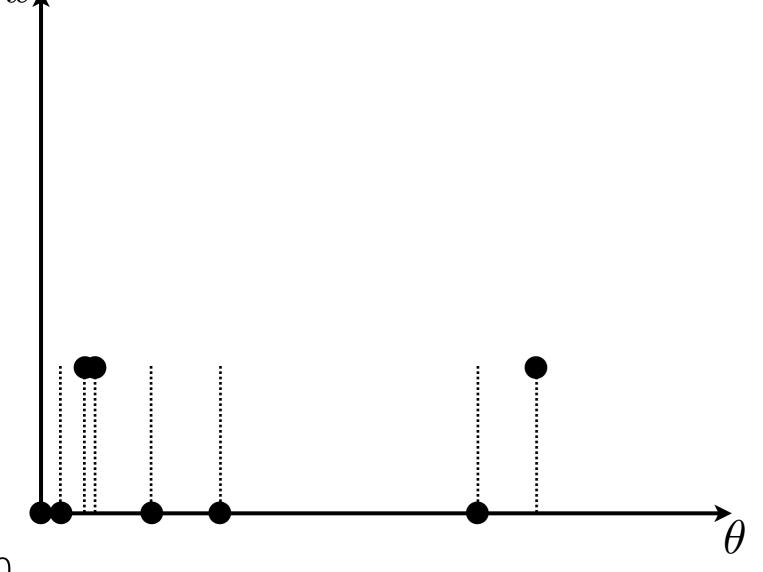
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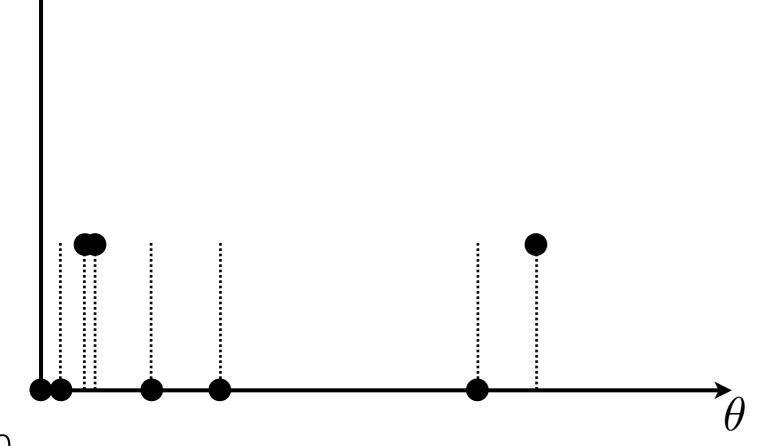


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- Poisson process rate measure $\nu(d\theta)$
- Marked Poisson process rate measure $\nu(d\theta)p(x|\theta)$

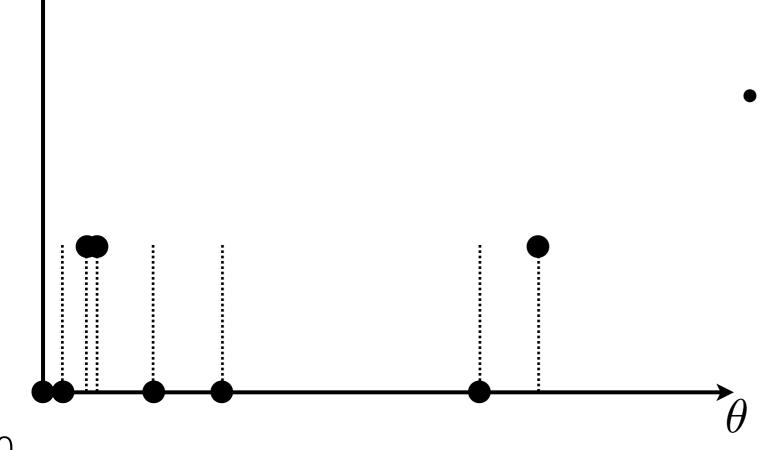


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- Conjugate prior:



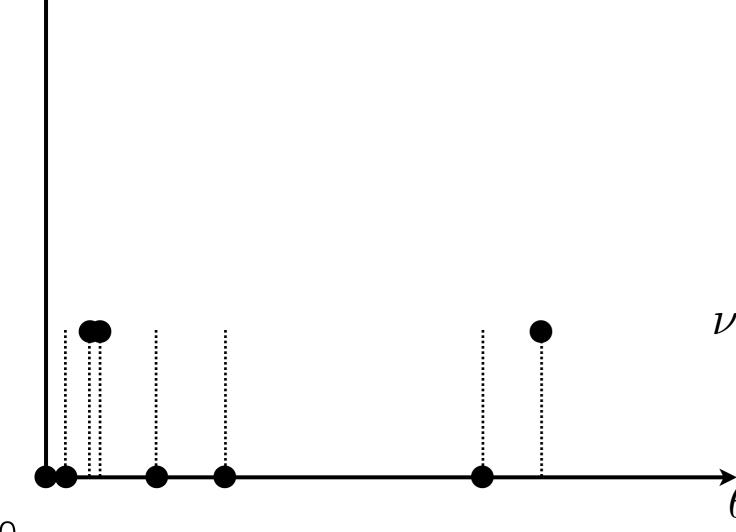
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- Conjugate prior:
 - Rate measure

$$\nu(d\theta) = \gamma \theta^{\alpha - 1} (1 - \theta)^{-\alpha - \beta} d\theta$$
$$\alpha \in (-1, 0], \beta > 0, \gamma > 0$$



$$p(x|\theta) = \theta^x (1+\theta)^{-1}$$

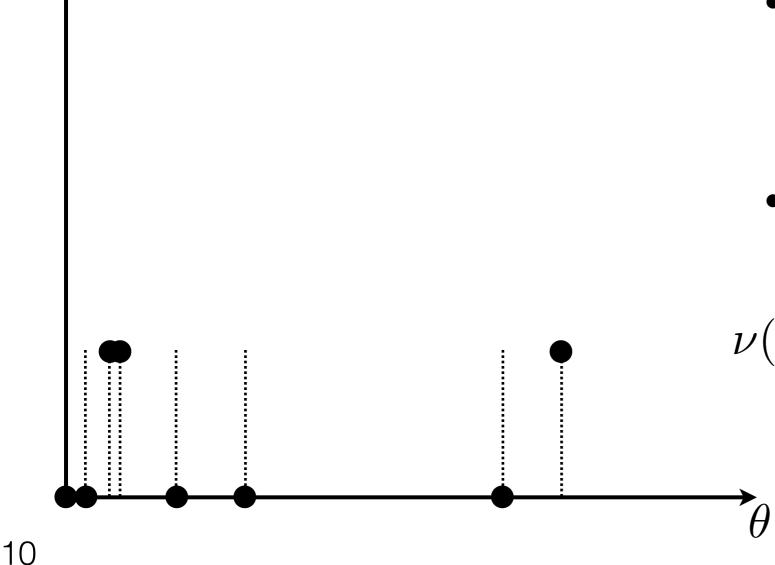
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 Beta prime fixed atoms



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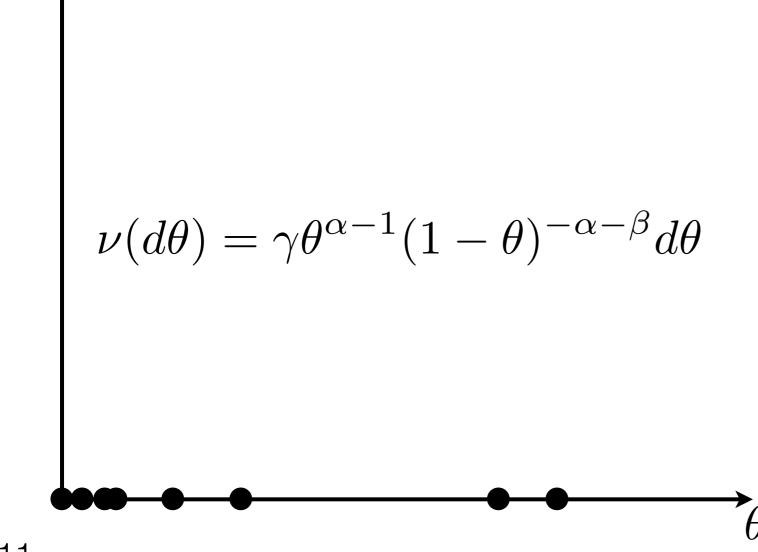
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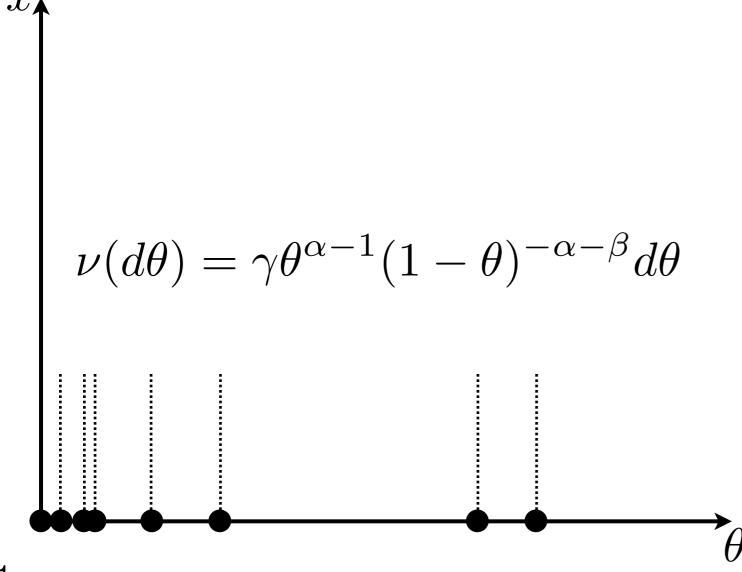
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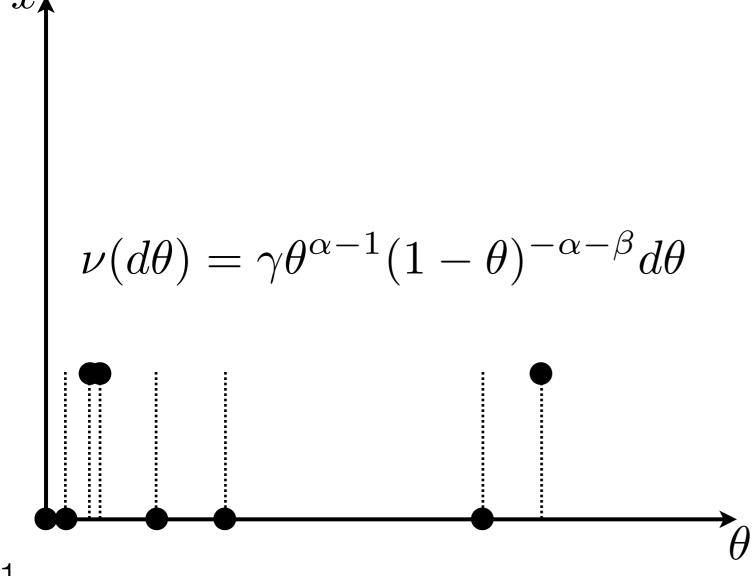
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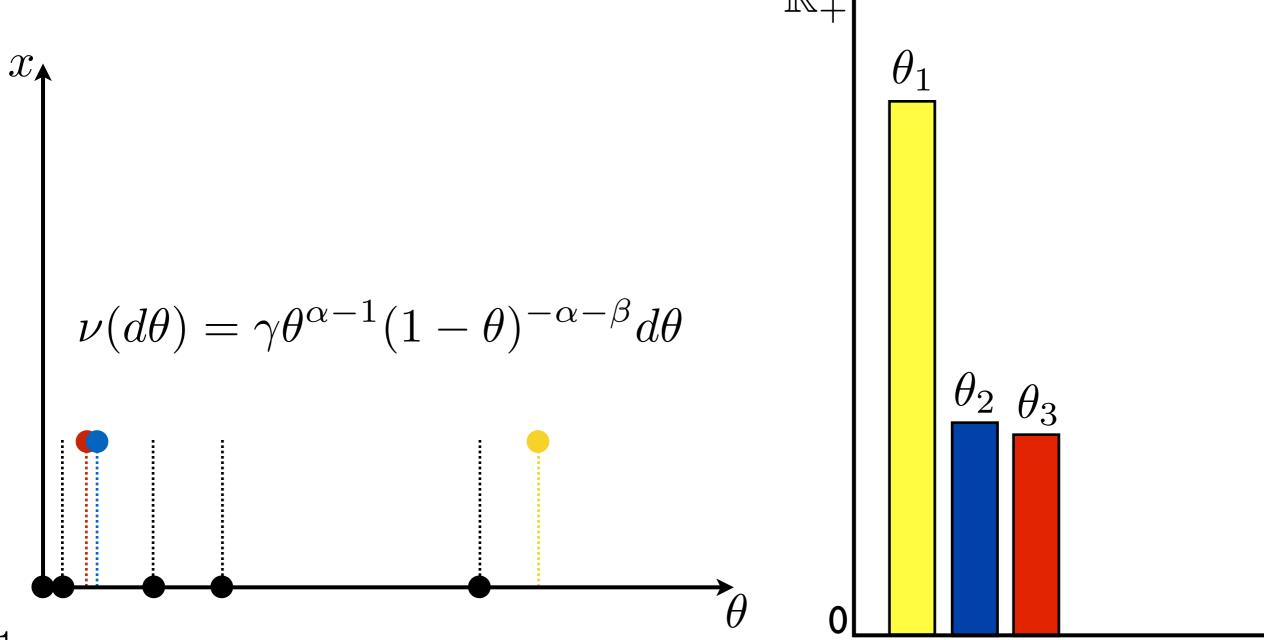
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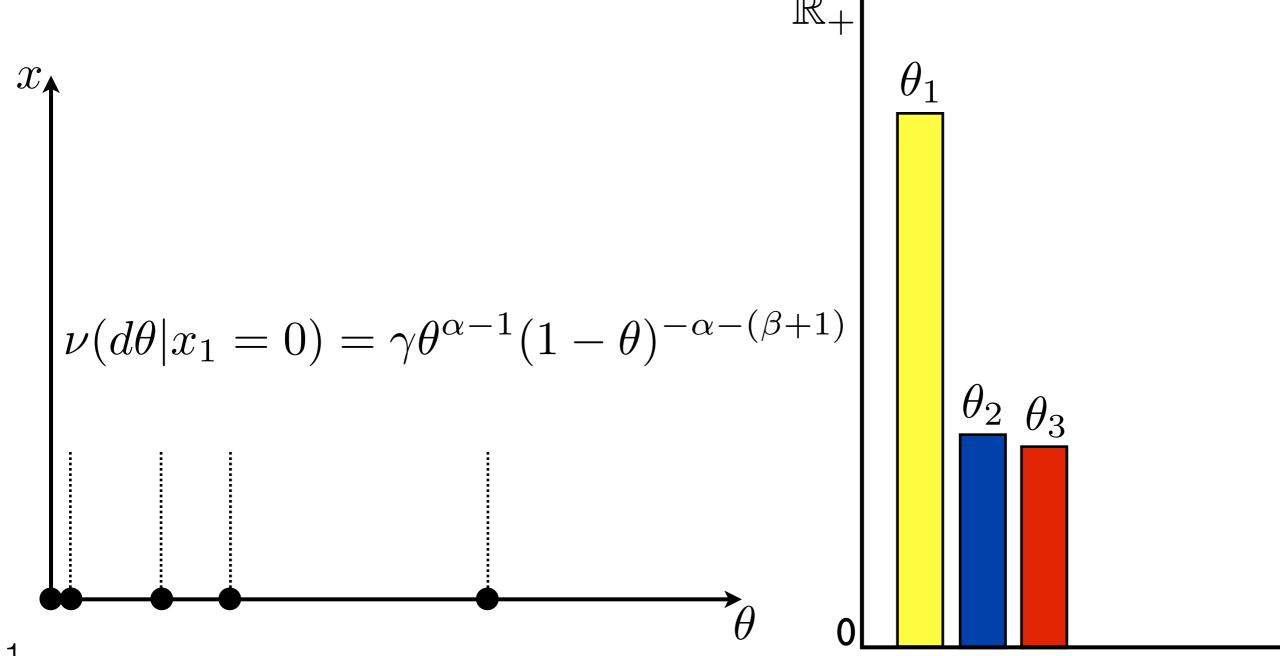
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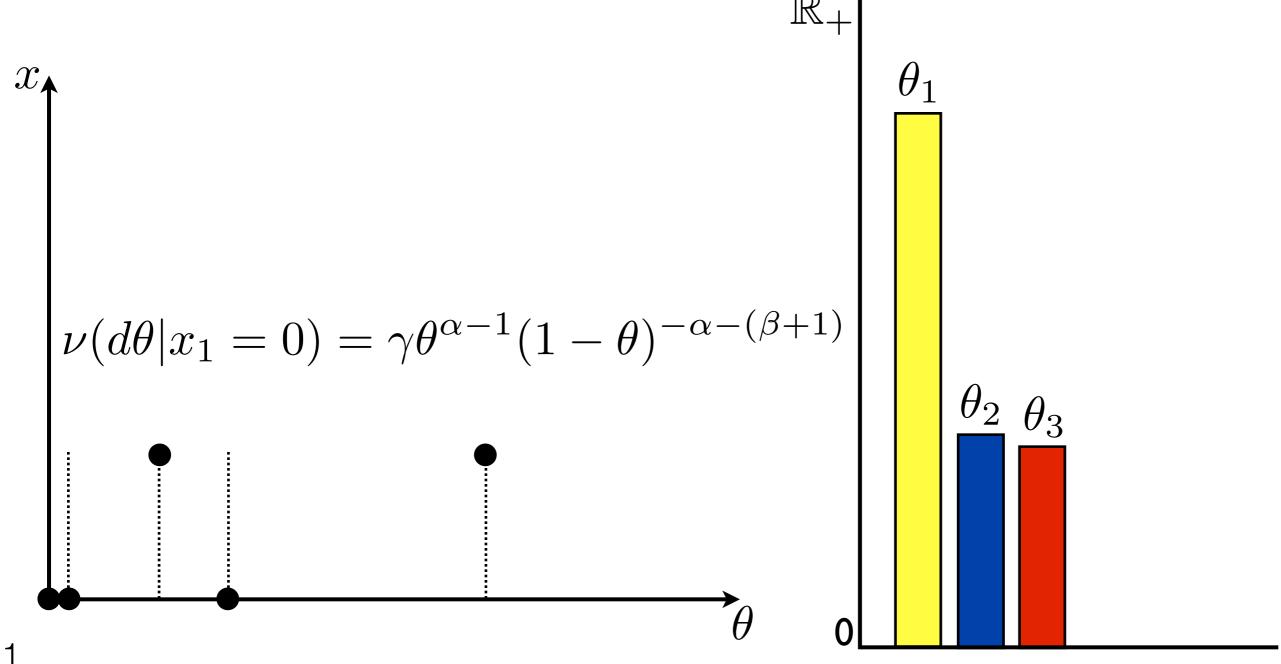
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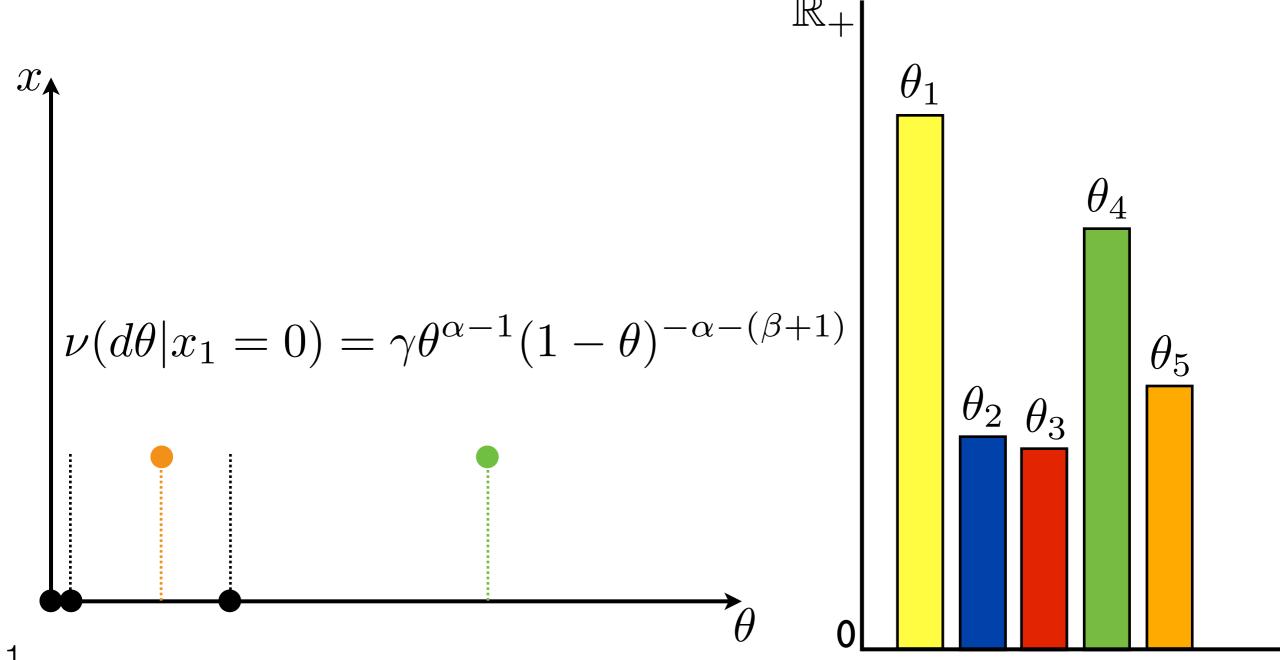
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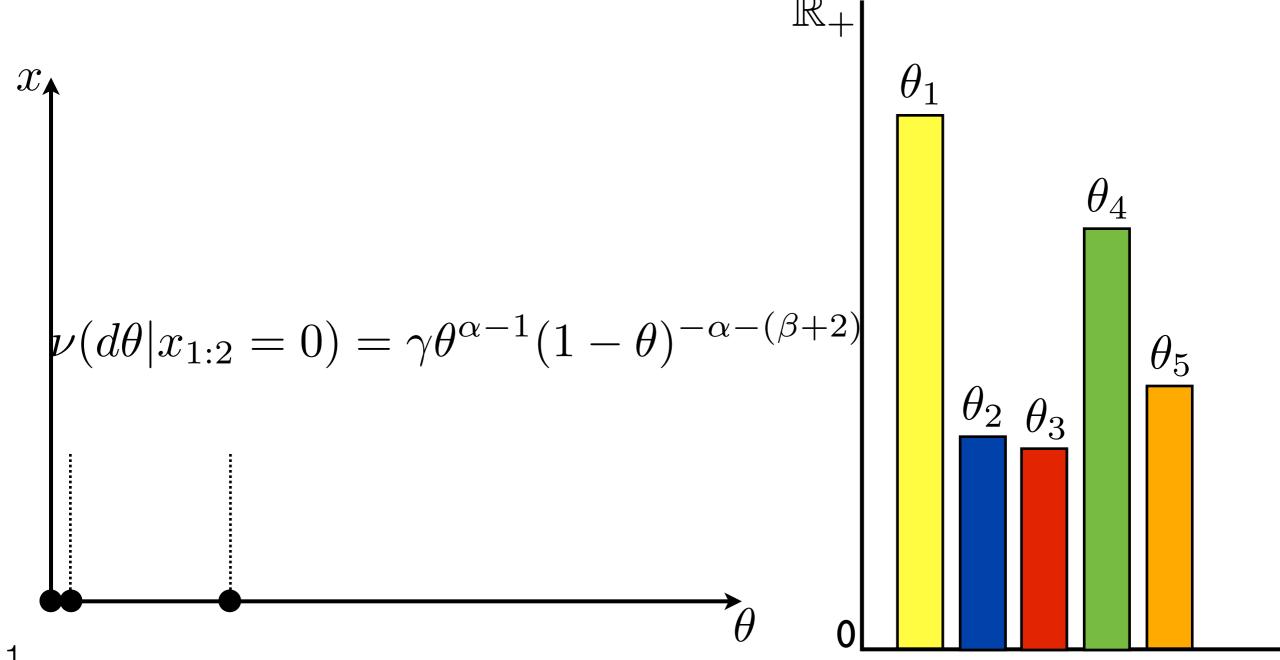
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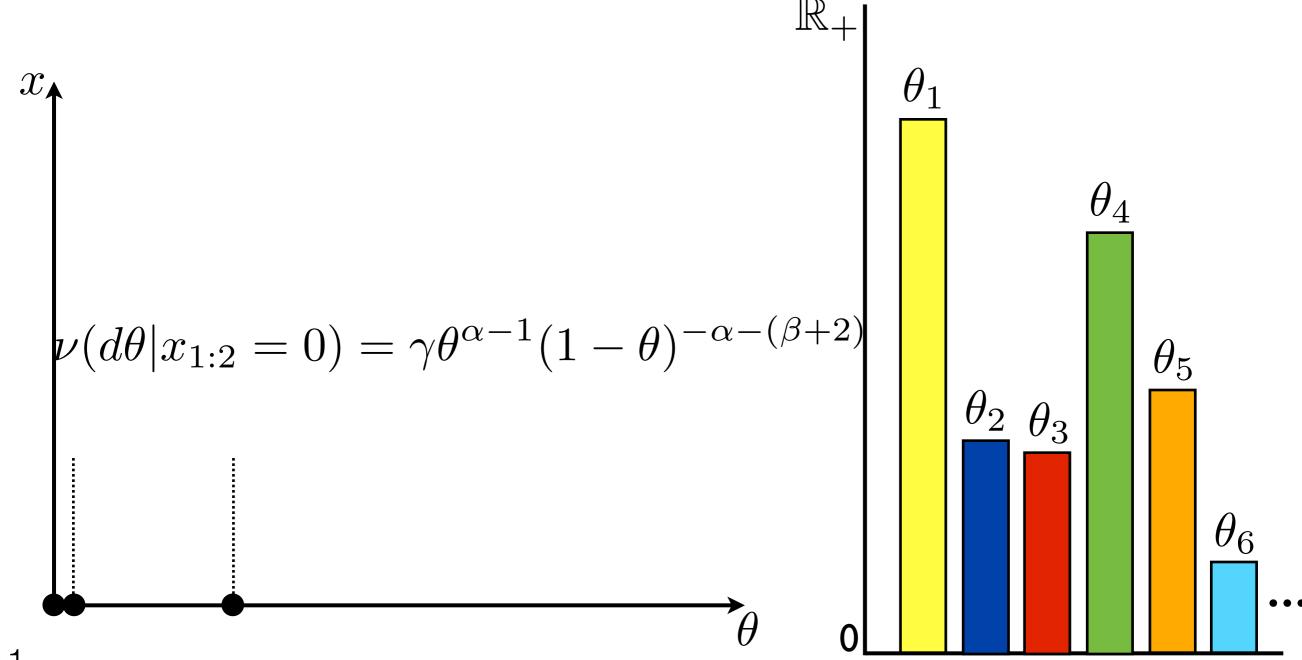
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Size-biased atoms, beta prime process

$$\alpha = 0$$

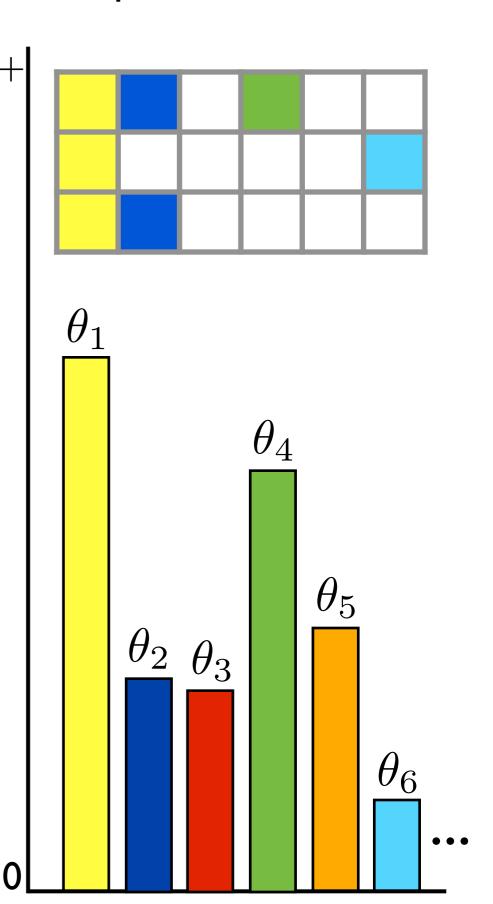
For m = 1, 2, ...

1. Draw

$$K_m^+ \sim \text{Poisson}\left(\gamma \frac{\beta}{\beta + m - 1}\right)$$

2. For $k = 1, ..., K_m^+$ Draw a rate of size

$$\theta_k \sim \text{BetaPrime}(1, \beta + m - 1)$$



Size-biased atoms, beta prime process

$$\alpha = 0$$

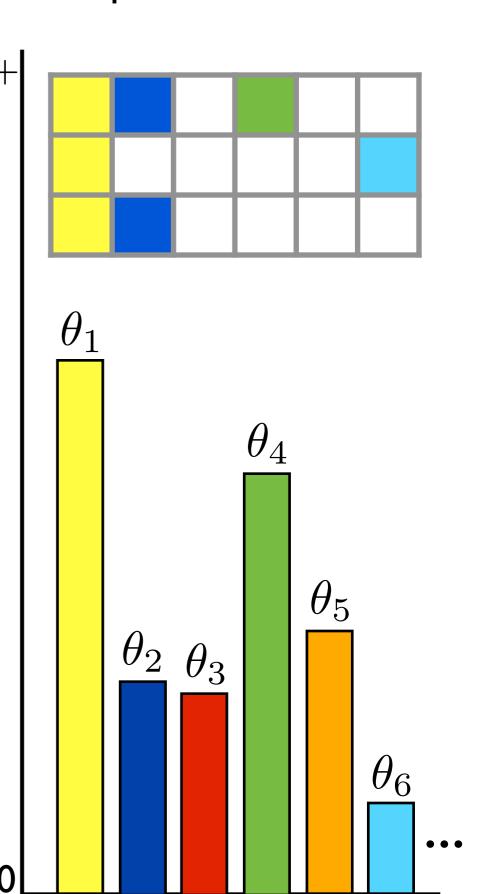
For m = 1, 2, ...

1. Draw

$$K_m^+ \sim \text{Poisson}\left(\gamma \frac{\beta}{\beta + m - 1}\right)$$

2. For $k = 1, ..., K_m^+$ Draw a rate of size

$$\theta_k \sim \text{BetaPrime}(1, \beta + m - 1)$$



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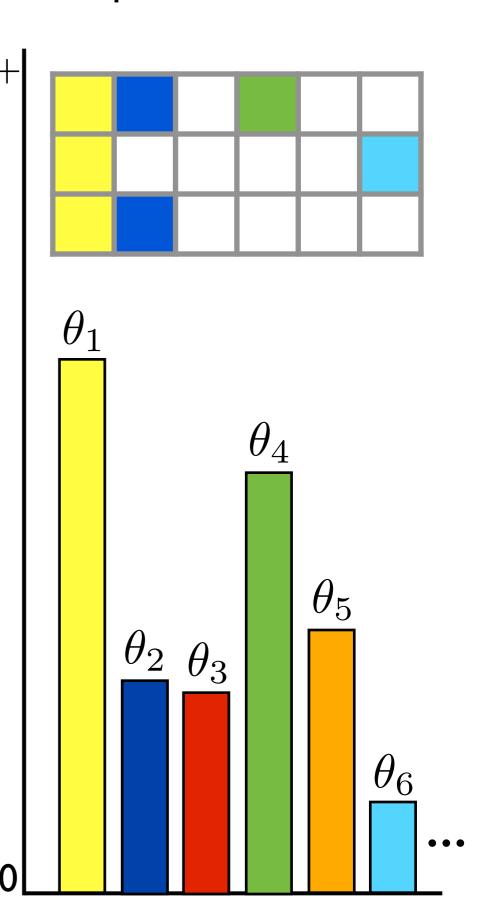
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Marginal process derivation is similar



Exponential family likelihood

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Conjugate prior

PPP rate measure $\nu(d\theta) = \gamma \exp\{\langle \xi, \eta(\theta) \rangle + \lambda [-A(\theta)]\}d\theta$ + fixed atoms $f(d\theta) = \exp\{\langle \xi_k, \eta(\theta) \rangle + \lambda_k [-A(\theta)] - B(\xi_k, \lambda_k)\}d\theta$

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$$K_m^+ \sim \text{Poisson}\left(\int_{x>0} \gamma \cdot \kappa(0)^{m-1} \cdot \kappa(x) \cdot \exp\left\{B(\xi + (m-1)\phi(0) + \phi(x), \lambda + m)\right\} dx\right)$$
$$f(d\theta) \propto \int_{x>0} \exp\left\{\langle \xi + (m-1)\phi(0) + \phi(x), \eta(\theta)\rangle + (\lambda + m)[-A(\theta)]\right\} dx$$

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Marginal process

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Marginal process

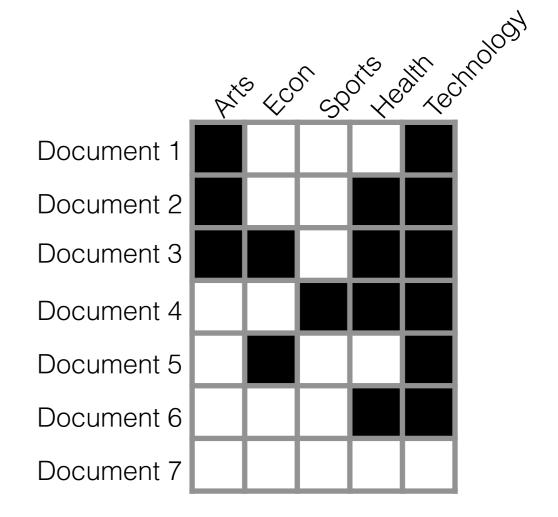
$$K_n^+ \text{ as above } \\ p(x_n|x_{1:(n-1)}) = \kappa(x_n) \exp\left\{-B(\xi + \sum_{m=1}^{n-1} x_m, \lambda + n - 1) + B(\xi + \sum_{m=1}^{n-1} x_m + x_n, \lambda + n)\right\}$$

 To satisfy BNP desiderata, likelihood must have a point mass at 0

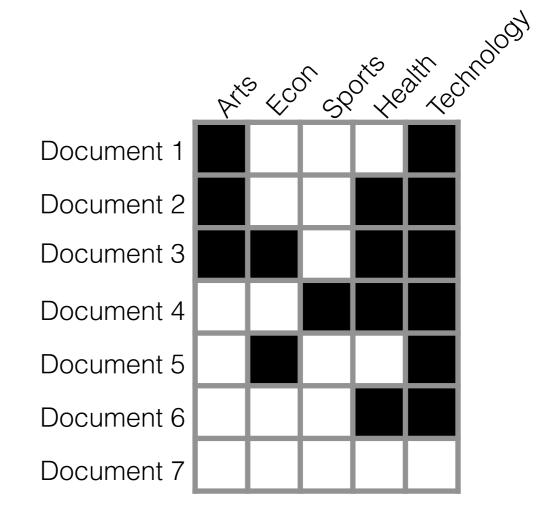
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- Poisson distribution direct result of Poisson process

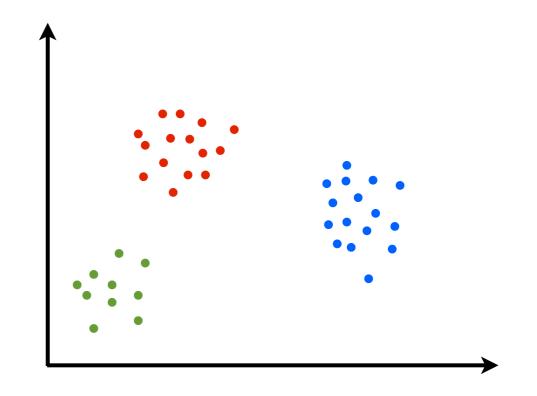
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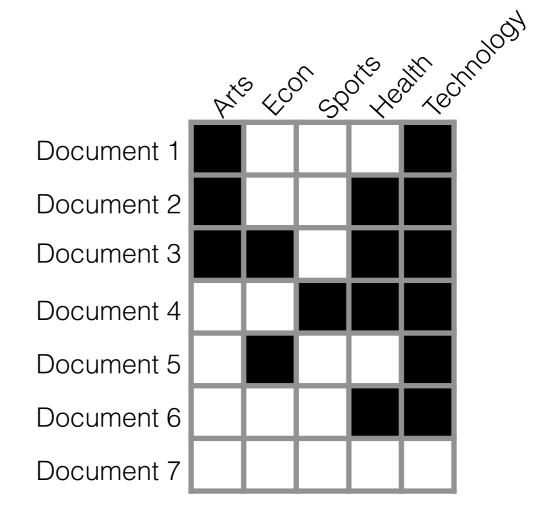
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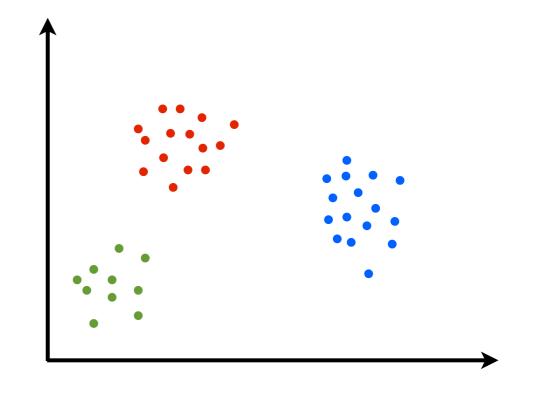




- To satisfy BNP desiderata, likelihood must have a point mass at 0
- Poisson distribution direct result of Poisson process
- Much previous work on conjugacy at a different level of a BNP hierarchy
- Can be used with arbitrary

 (i.e., discrete, continuous, or other) data likelihood





References

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