### data-microscopes: Bayesian non-parametric inference made simple in Python

Stephen Tu tu.stephenl@gmail.com

SF Python - August 20, 2014

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- Countless more (sorry if I missed yours)

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#### Goal

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Doing it well means being correct and fast!

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- This will lead us to the Dirichlet process mixture model.

• Why not just *k*-means?

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$$\underset{C_1,...,C_k}{\operatorname{argmin}} \sum_{i=1}^k \sum_{x_j \in C_i} \|x_j - \mu_i\|^2 \qquad \mu_i = \frac{1}{|C_i|} \sum_{x_j \in C_i} x_j$$

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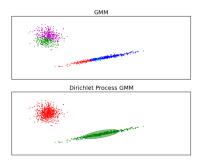


Figure: http://scikit-learn.org/stable/\_images/plot\_gmm\_0011.png

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• What if you have a *model* of the data? E.g. you know the data is from a mixture of gaussian distributions.

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- What if there is no metric on the data type? E.g. the data is categorical?

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- Describes the generative process of n i.i.d. observations  $Y_1, ..., Y_n$  as:

$$G \sim \mathsf{DirichletProcess}(lpha, H)$$
  
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where  $F(\cdot)$  is a likelihood model (e.g. Gaussian),  $H(\cdot)$  is the *prior* distribution (e.g. Normal-Inverse-Wishart) over the parameters of  $F(\cdot)$ , and  $\alpha \in \mathbb{R}^+$  is chosen a-priori.

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### Dirichlet process

#### Dirichlet process

#### Definition

Let  $H(\cdot)$  be a measure over S and  $\alpha > 0$ . We say G is drawn from a Dirichlet Process, written as  $G \sim DP(\alpha, H)$  if for any (measurable) partition of  $S = (P_1, ..., P_n)$  we have

$$(G(P_1),...,G(P_n)) \sim Dirichlet(\alpha H(P_1),...,\alpha H(P_n))$$

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Wait... what?!

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Don't worry! Alternative view known as the **Chinese Restaurant Process** which is *way* more intuitive.

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- Each table  $T_i$  has a vector  $\Theta^{(i)} = (\theta_1^{(i)}, ..., \theta_D^{(i)})$ , with each  $\theta_i^{(i)} \sim \text{Beta}(\gamma, \beta), j=1, ..., D$ .

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- To draw a value  $Y_i$ , first pick a table  $T_j$ , and then draw  $Y_i \sim (\mathsf{Bernoulli}(\theta_1^{(j)}), ..., \mathsf{Bernoulli}(\theta_D^{(j)}))$ .

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- To reference our previous notation,  $H = \text{Beta}(\gamma, \beta)^D$  and  $F = Bernoulli(\cdot)^D$ .

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- Note: can prove that expected number of tables filled is  $O(\alpha \log n)$ .



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data-microscopes only implements MCMC (for now!).

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#### Problem statement

Given n data points  $\mathcal{Y} = (Y_1, ..., Y_n)$ , our goal is to learn the distribution  $p(\mathcal{C}|\mathcal{Y})$ , where  $\mathcal{C}$  is the clustering (assignment vector) of  $\mathcal{Y}$ .

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**Note:** when we say "learn the distribution" we mean draw (independent) samples from.

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- Sampling  $c_i^{(t)} \leftarrow p(c_i|\mathcal{C}_{\neg i}^{(t-1)}, \mathcal{Y}), i = 1, ..., n$  over and over (and over) will eventually get us  $p(\mathcal{C}|\mathcal{Y})$ .

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- The above strategy is called Gibbs sampling.



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$$p(c_{i}=k|\mathcal{C}_{\neg i},\mathcal{Y}) \propto p(c_{i}=k,\mathcal{C}_{\neg i},\mathcal{Y})$$

$$= p(c_{i}=k|\mathcal{C}_{\neg i})p(Y_{i}|\mathcal{Y}^{(k)})$$

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If we pick H and F nicely, the integral on the RHS has an analytical solution!

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$$\frac{|\mathcal{Y}_{\neg i}^{(k)}|}{n-1+\alpha} \prod_{d=1}^{D} \frac{\left(\beta + \sum_{y_k \in \mathcal{Y}_{\neg i}^{(k)}} y_k^{(d)}\right)^{y_i^{(d)}} \left(\gamma + |\mathcal{Y}_{\neg i}^{(k)}| - \sum_{y_k \in \mathcal{Y}_{\neg i}^{(k)}} y_k^{(d)}\right)^{(1-y_i^{(d)})}}{\beta + \gamma + |\mathcal{Y}_{\neg i}^{(k)}|}$$

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$$\frac{\alpha}{n-1+\alpha} \prod_{d=1}^{D} \left( \frac{\beta}{\beta+\gamma} \right)^{y_i^{(d)}} \left( \frac{\gamma}{\beta+\gamma} \right)^{1-y_i^{(d)}}$$

when k is a new cluster.

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# Using data-microscopes

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Now let's see the library in action!