Bayesian Theory and Computation

Lecture 18: Dirichlet Processes

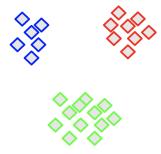


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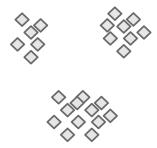
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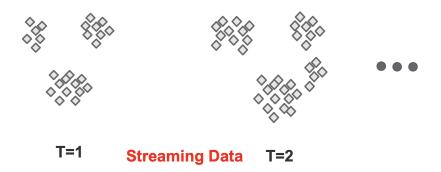
How to choose the number of clusters?



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How to choose the number of clusters?



- ► A generative approach to clustering
 - pick one of K clusters from a distribution $\pi = (\pi_1, \dots, \pi_K)$
 - ▶ generate a data point from a cluster-specific probability distribution
- ► This yields a finite mixture model:

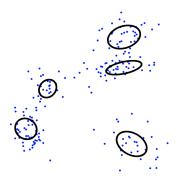
$$p(x|\phi,\pi) = \sum_{k=1}^{K} \pi_k p(x|\phi_k)$$

where π and $\phi = (\phi_1, \dots, \phi_K)$ are the parameters, and here we assume the sae parameterized family for each cluster for simplicity.

▶ Data $\{x_i\}_{i=1}^N$ are assumed to be generated conditionally iid from this mixture model.



▶ For Gaussian mixtures, $\phi_k = (\mu_k, \Sigma_K)$ and $p(x|\phi_k)$ is a Gaussian density with mean μ_k and covariance matrix Σ_k



- ▶ Mixture models make the assumption that each data point arises from a single mixture component, i.e., the *k*th cluster is by definition the set of data points arising from the *k*th mixture component.
- ► Can capture this explicitly via a latent multinomial variable Z:

$$p(x|\phi, \pi) = \sum_{k=1}^{K} p(Z = k|\pi)p(x|Z = k, \phi)$$
$$= \sum_{k=1}^{K} \pi_k p(x|\phi_k)$$

► Another way to express this: define an underlying measure

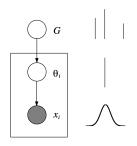
$$G = \sum_{k=1}^{K} \pi_k \delta_{\phi_k}$$

where δ_{ϕ_k} is an *atom* (Dirac delta function) at ϕ_k .

Now we can redefine the process of obtaining a sampling from a finite mixture model as follows. For i = 1, ..., n:

$$\theta_i \sim G$$
$$x_i \sim p(\cdot | \theta_i)$$

Note that each θ_i is equal to one of the underlying ϕ_k . Indeed, the subset of $\{\theta_i\}$ that maps to ϕ_k is exactly the kth cluster



$$G = \sum_{k=1}^{K} \pi_k \, \delta_{\phi_k}$$
 $\theta_i \sim G$
 $x_i \sim p(\cdot | \theta_i)$

- ▶ Bayesian approaches allow us to integrate out model parameters
- ▶ Need to place priors on the parameters ϕ and π
- ▶ The choice of prior for ϕ is model-specific; e.g., we may use conjugate normal/inverse-gamma priors for a Gaussian mixture model. Let us denote this prior as G_0 .
- ▶ What to choose for the mixture weights π ? A common choice is a symmetric Dirichlet prior, $Dir(\alpha_0/K,...,\alpha_0/K)$
 - ▶ the symmetry accords with the common assumption of the order-free of the labels of the mixture components
 - the concentration parameter α_0 controls concentration level of the labels



$$\phi_k \sim G_0
\pi_k \sim \text{Dir}(\alpha_0/K, \dots, \alpha_0/K)
G = \sum_{k=1}^K \pi_k \, \delta_{\phi_k}
\theta_i \sim G
x_i \sim p(\cdot | \theta_i)$$

 \blacktriangleright Note that G is now a random measure



- ► Posterior distributions can't be found analytically; nor can predictive distributions (for future observations)
- ► However, a variety of MCMC sampling algorithms are available
- \blacktriangleright Use the indicators Z within a Gibbs sampler. Give Z, we know which data points belong to which cluster, so:
 - $ightharpoonup p(\pi|Z,\phi)$: standard multinomial-Dirichlet conjugacy
 - ▶ $p(\phi|Z,\pi)$: separate updates for each cluster; i.e., for each ϕ_k (and conjugacy of G_0 and $p(\cdot|\phi)$ can make this easy)
 - $ightharpoonup p(Z|\pi,\phi)$: multinomial classification
- ▶ We can also use variational inference.



- \blacktriangleright How to choose K, the number of mixture components?
- ▶ Various generic model selection methods can be considered: e.g., cross-validation, bootstrap, AIC, BIC, DIC, Laplace, bridge sampling, etc
- ightharpoonup Or we can place a parametric prior on K (e.g., Poisson) and use Bayesian methods
- ► The Dirichlet process provides a nonparametric Bayesian alternative.



- ▶ Make sure we always have more clusters than we need.
- ► How about infinite clusters a priori?

$$p(x|\phi,\pi) = \sum_{k=1}^{\infty} \pi_k p(x|\phi_k)$$

- ► A finite data set will always use a finite, but random, number of clusters.
- ► How to choose the prior?
- ▶ We need something like a Dirichlet prior, but with an infinte number of components.

▶ Relation to gamma distribution: If $\eta_k \sim \text{Gamma}(\alpha_k, \beta)$ independently, then

$$S = \sum_{k} \eta_k \sim \text{Gamma}\left(\sum_{k} \alpha_k, \beta\right)$$

and

$$V = (v_1, \dots, v_k) = (\eta_1/S, \dots, \eta_k/S) \sim \text{Dir}(\alpha_1, \dots, \alpha_K)$$

▶ Therefore, if $(\pi_1, \ldots, \pi_K) \sim \text{Dir}(\alpha_1, \ldots, \alpha_K)$ then

$$(\pi_1 + \pi_2, \pi_3, \dots, \pi_K) \sim \operatorname{Dir}(\alpha_1 + \alpha_2, \alpha_3, \dots, \alpha_K)$$

This is known as the collapsing property.



- ► The beta distribution is a Dirichlet distribution on the 1-simplex
- Let $(\pi_1, \ldots, \pi_K) \sim \text{Dir}(\alpha_1, \ldots, \alpha_K)$ and $\theta \sim \text{Beta}(\alpha_1 b, \alpha_1 (1 b)), \ 0 < b < 1.$
- ► Then

$$(\pi_1\theta, \pi_1(1-\theta), \pi_2, \dots, \pi_K) \sim \operatorname{Dir}(\alpha_1b_1, \alpha_1(1-b_1), \alpha_2, \dots, \alpha_K)$$

▶ More generally, if $\theta \sim \text{Dir}(\alpha_1 b_1, \alpha_1 b_2, \dots, \alpha_1 b_N)$, $\sum_i b_i = 1$, then

$$(\pi_1\theta_1,\ldots,\pi_1\theta_N,\pi_2,\ldots,\pi_K) \sim \text{Dir}(\alpha_1b_1,\ldots,\alpha_1b_N,\alpha_2,\ldots,\alpha_K)$$

This is known as the splitting property.



▶ Renormalization. If $(\pi_1, \ldots, \pi_K) \sim \text{Dir}(\alpha_1, \ldots, \alpha_K)$, and

$$V = (V_2, V_3, \dots, V_K), \quad V_k = \frac{\pi_k}{\sum_{k>2} \pi_k}$$

▶ What is the distribution of V?

$$V \sim \text{Dir}(\alpha_2, \dots, \alpha_K)$$

▶ All these properties can be easily verified using the aforementioned gamma distribution representation.

- ▶ Let G_0 be a distribution on some space Ω , e.g. a Gaussian distribution on the real line.
- \blacktriangleright Assume that π, ϕ have the following distributions

$$\phi_k \sim G_0$$

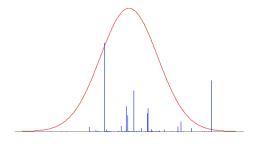
$$\pi \sim \lim_{K \to \infty} \operatorname{Dir}\left(\frac{\alpha_0}{K}, \dots, \frac{\alpha_0}{K}\right)$$

- ► Then $G := \sum_{k=1}^{\infty} \pi_k \delta_{\phi_k}$ defines an infinite distribution over G_0 .
- \blacktriangleright We say (informally) that G follows a Dirichlet Process

$$G \sim \mathrm{DP}(\alpha_0 G_0)$$



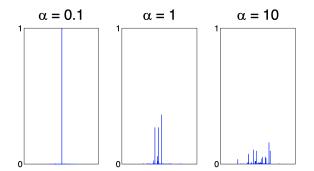
- ► Samples from the Dirichlet process are *discrete*.
- We call the point masses in the resulting distribution, atoms.



 \blacktriangleright The base measure G_0 determines the locations of the atoms.



- ► The concentration parameter α_0 determines the distribution over atom sizes.
- ▶ Small values of α_0 gives *sparse* distributions.



- ▶ Let (Ω, \mathcal{B}) be a measurable space, with G_0 a probability measure on the space, and let α_0 be a positive real number.
- ▶ A Dirichlet process is the distribution of a random probability measure G over (Ω, \mathcal{B}) such that, for any finite partition (A_1, \ldots, A_r) of Ω , the random vector $(G(A_1), \ldots, G(A_r))$ follows a finite-dimensional Dirichlet distribution:

$$(G(A_1),\ldots,G(A_r)) \sim \operatorname{Dir}(\alpha_0 G_0(A_1),\ldots,\alpha_0 G(A_r))$$

▶ We write $G \sim \mathrm{DP}(\alpha_0 G_0)$, and call G_0 the base measure, α_0 the concentration parameter.



▶ Let $A_1, ..., A_K$ be a partition of Ω . Let $G(A_k)$ be the mass assigned by $G \sim \mathrm{DP}(\alpha_0 G_0)$ to partition A_k . Then

$$(G(A_1),\ldots,G(A_K)) \sim \operatorname{Dir}(\alpha_0 G_0(A_1),\ldots,\alpha_0 G_0(A_K))$$

 \blacktriangleright If we see an observation in the j-th segment, then

$$(G(A_1), \dots, G(A_K) | \theta_1 \in A_j)$$

 $\sim \text{Dir}(\alpha_0 G_0(A_1), \dots, \alpha_0 G(A_j) + 1, \dots, \alpha_0 G_0(A_K)).$

- ▶ This is true for all possible partitions of Ω .
- ▶ Therefore, the posterior distribution of G, given an observation ϕ , is given by

$$G|\theta_1 = \phi \sim \mathrm{DP}(\alpha_0 G_0 + \delta_\phi)$$



- ► The Dirichlet process clusters observations.
- ► A new data point can either join an existing cluster, or start a new cluster.
- ▶ Question: What is the predictive distribution for a new data point?
- Assume G_0 is a continuous distribution on Ω. This means for every point ϕ in Ω, $G_0(\phi) = 0$.
- ► First data point:
 - ► Start a new cluster
 - ▶ Sample a parameter $\phi_1 \sim G_0$ for that cluster.



- ▶ We have now split our parameter space in two: the singleton ϕ_1 , and everything else.
- ▶ Let π_1 be the size of atom at ϕ_1 .
- ▶ The combined mass of all the other atoms is $\pi_* = 1 \pi_1$.
- ► According to the DP,

$$(\pi_1, \pi_*) \sim \text{Dir}(0, \alpha_0)$$

▶ Given $\theta_1 = \phi_1$, the posterior is

$$(\pi_1, \pi_*)|\theta_1 = \phi_1 \sim \operatorname{Dir}(1, \alpha_0)$$

▶ If we integrate out π_1 , we get

$$p(\theta_2 = \phi_k | \theta_1 = \phi_1) = \int p(\theta_2 = \phi_k | (\pi_1, \pi_*)) p((\pi_1, \pi_*) | \theta_1 = \phi_1) d\pi_1$$

$$= \int \pi_k \operatorname{Dir}((\pi_1, 1 - \pi_1) | 1, \alpha_0) d\pi_1$$

$$= \mathbb{E}_{\operatorname{Dir}(1, \alpha_0)} \pi_k$$

$$= \begin{cases} \frac{1}{1 + \alpha_0} & \text{if } k = 1 \\ \frac{\alpha_0}{1 + \alpha_0} & \text{for new } k. \end{cases}$$

- Lets say we choose to start a new cluster, and sample a new parameter $\phi_2 \sim G_0$. Let π_2 be the size of the atom at ϕ_2 .
- ► Similarly, the posterior is

$$(\pi_1, \pi_2, \pi_*)|\theta_1 = \phi_1, \theta_2 = \phi_2 \sim \text{Dir}(1, 1, \alpha_0)$$

▶ If we integrate out $\pi = (\pi_1, \pi_2, \pi_*)$, we get

$$p(\theta_3 = \phi_k | \theta_1 = \phi_1, \theta_2 = \phi_2)$$

$$= \int p(\theta_3 = \phi_k | \pi) p(\pi | \theta_1 = \phi_1, \theta_2 = \phi_2) d\pi$$

$$= \mathbb{E}_{\text{Dir}(1,1,\alpha_0)} \pi_k$$

$$= \begin{cases} \frac{1}{2+\alpha_0} & \text{if } k = 1\\ \frac{1}{2+\alpha_0} & \text{if } k = 2\\ \frac{\alpha_0}{2+\alpha_0} & \text{for new } k. \end{cases}$$



▶ In general, if m_k is the number of times we have seen $X_i = k$, and K is the total number of observed values,

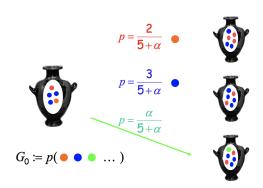
$$p(\theta_{n+1} = \phi_k | \theta_1, \dots, \theta_n) = \int p(\theta_{n+1} = \phi_k | \pi) p(\pi | \theta_1, \dots, \theta_n) d\pi$$

$$= \mathbb{E}_{\text{Dir}(m_1, \dots, m_K, \alpha_0)} \pi_k$$

$$= \begin{cases} \frac{m_k}{n + \alpha_0} & \text{if } k \leq K \\ \frac{\alpha_0}{n + \alpha_0} & \text{for new cluster.} \end{cases}$$

- ► We tend to see observations that we have seen before, i.e., rich-get-richer property
- ▶ We can always add new features, a typical nonparametric behavior.





Adapted from Eric Xing

▶ Joint: $G(\overset{\bullet}{\Psi}) \sim \mathrm{DP}(\alpha_0 G_0)$

► Marginal: $\theta_{n+1}|\theta_{\leq n}, \alpha_0, G_0 \sim \sum_{k=1}^K \frac{m_k}{n+\alpha_0} \delta_{\phi_k} + \frac{\alpha_0}{n+\alpha_0} G_0$.

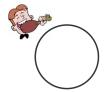


- ► The resulting distribution over data points can be thought of using the following urn scheme (Blackwell and MacQueen, 1973).
- ▶ An urn initially contains a black ball of mass α_0 .
- For n = 1, 2, ..., sample a ball from the urn with probability proportional to its mass.
- ▶ If the ball is black, choose a previously unseen color, record that color, and return the black ball plus a unit-mass ball of the new color to the urn.
- ► If the ball is not black, record it's color and return it, plus another unit-mass ball of the same color, to the urn.



► The distribution over partitions can also be described in terms of the following restaurant metaphor:

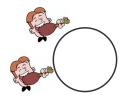
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- ▶ The first customer enters a restaurant, and picks a table.







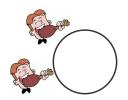
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- ▶ The first customer enters a restaurant, and picks a table.
- ► The *n*-th customer enters the restaurant. He sits at an existing table with probability $\frac{m_k}{n-1+\alpha_0}$, where m_k is the number of people sat the table k. He starts a new table with probability $\frac{\alpha_0}{n-1+\alpha_0}$.







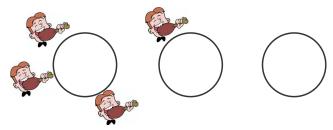
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- ▶ An interesting fact: the distribution over the clustering of the first N customers does not depend on the order in which they arrived.
- ► However, the customers are not independent. They tend to sit at popular tables.
- ▶ We say that distributions like this are *exchangeable*.

$$p(\theta_1, \dots, \theta_N) = p(\theta_{\sigma(1)}, \dots, \theta_{\sigma(n)})$$

▶ By **de Finetti**'s theorem, there exists a random distribution G and a prior P(G) such that

$$p(\theta_1, \dots, \theta_N) = \int \prod_{i=1}^N G(\theta_i) dP(G)$$

▶ In our setting, the prior p(G) is just $DP(\alpha_0G_0)$, thus establishing existence.



▶ Define an infinite sequence of Beta random variables:

$$\beta_k \sim \text{Beta}(1, \alpha_0), \quad k = 1, 2, \dots$$

▶ Now define an infinite sequence of mixing proportions as:

$$\pi_1 = \beta_1$$

$$\pi_k = \beta_k \prod_{\ell=1}^{k-1} (1 - \beta_\ell), \quad k = 2, 3, \dots$$

▶ This can be viewed as breaking off portions of a stick:

$$\beta_1$$
 β_2 $(1-\beta_1)$...

▶ We now have an explicit formula for each π_k :

$$\pi_k = \beta_k \prod_{\ell=1}^{k-1} (1 - \beta_\ell)$$

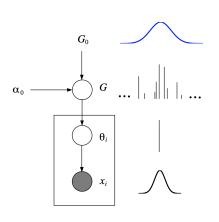
▶ We can easily see that $\sum_{k=1}^{\infty} \pi_k = 1$:

$$1 - \sum_{k=1}^{\infty} \pi_k = 1 - \beta_1 - \beta_2 (1 - \beta_1) - \beta_3 (1 - \beta_1) (1 - \beta_2) - \cdots$$
$$= (1 - \beta_1) (1 - \beta_2 - \beta_3 (1 - \beta_2) - \cdots)$$
$$= \prod_{k=1}^{\infty} (1 - \beta_k) = 0$$

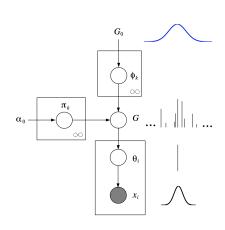
▶ Let $\phi_k \sim G_0, \forall k, G = \sum_{k=1}^{\infty} \pi_k \delta_{\phi_k}$ has a clean definition as a random measure. In fact,

$$G \sim \mathrm{DP}(\alpha_0 G_0).$$





Polya urn construction



Stick breaking construction



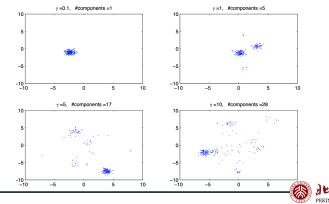
- ▶ Now we can use a Dirichlet process as the prior for an unknown mixture distribution (with potentially infinite mixture components).
- ▶ Suppose we have $x_1, ..., x_n$ observations from some unknown distribution.
- ▶ We can model the unknown distribution of x as a mixture of simple distributions of the form $f(\cdot|\theta)$.
- We denote the mixing distribution over θ as G and let the prior over G be a Dirichlet process

$$x_i | \theta_i \sim f(\theta_i)$$

 $\theta_i | G \sim G$
 $G \sim DP(\alpha_0 G_0)$



- Multiple subjects can be mapped to the same ϕ . This creates a clustering of subjects.
- ▶ The following graphs shows 4 different data sets (n = 200) randomly generated from distributions sampled from Dirichlet process mixture priors with α_0 .



 \blacktriangleright We can integrate out G to get the CRP. Note that the CRP is exchangeable, which induces the conditional priors

$$p(\theta_i|\theta_{-i},\alpha_0,G_0) = \frac{\alpha_0}{n-1+\alpha_0}G_0(\theta_i) + \sum_{k=1}^{K^{(-i)}} \frac{m_k^{(-i)}}{n-1+\alpha_0}\delta_{\phi_k^{(-i)}}$$

- ▶ Let z_i be the cluster allocation of the *i*-th data point. The collapsed Gibbs sampler alternates between
 - ightharpoonup update z_i

$$p(z_i = k | x_i, z_{-i}, \phi_{1:K}) \propto \begin{cases} m_k^{(-i)} f(x_i | \phi_k^{(-i)}) & k \le K^{(-i)} \\ \alpha_0 \int f(x_i | \theta) dG_0(\theta) & k = K^{(-i)} + 1 \end{cases}$$

ightharpoonup update ϕ_k

$$p(\phi_k|z_{1:n},x_{1:n}) \propto G_0(\phi_k) \prod_i f(x_i|\phi_k)$$

▶ If G_0 is conjugate to f, the above steps can be evaluated accurately.

 \blacktriangleright For the concentration parameter α_0 , we have

$$p(K|\alpha_0) \propto \alpha_0^K \frac{\Gamma(\alpha_0)}{\Gamma(\alpha_0 + n)}$$

where K is the number of unique ϕ 's (e.g., the number of clusters).

▶ Therefore, given K and the prior distribution $P(\alpha_0)$ we can sample from the posterior distribution of α_0 using the MH algorithm or the Gibbs sampling method of Escobar and West (1995).

- ▶ We are only updating one data point at a time.
- ► Imagine two "true" clusters are merged into a single cluster, a single data point is unlikely to "break away".
- ► Getting to the true distribution involves going through low probability states, i.e., mixing can be slow.
- ▶ If the likelihood is not conjugate, integrating out parameter values for new features can be difficult.
- ▶ Neal (2000) offers a variety of algorithms.



- ► The stick-breaking representation orders the mixture components so that the weights are stochastically decreasing. For a sufficiently large T, we will have $\sum_{k>T} \pi_k \approx 0$.
- ► Therefore, we can truncate the stick-breaking construction at a fixed value T and let $\beta_T = 1$.
- ▶ This implies $\pi_k = 0, \forall k > T$, and the distribution of

$$G_T = \sum_{k=1}^{T} \pi_k \delta_{\phi_k}$$

is known as a truncated Dirichlet process.

Variational distance between distributions of marginals from a DP and from its truncation at T is approximately $4n \exp(-(T-1)/\alpha_0)$. T doesn't have to be very large to get a good approximation.

- ► State of the Markov chain: $(\beta_{1:T-1}, \phi_{1:T}, z_{1:n})$.
- ▶ Update z_i by multinomial sampling with

$$p(z_i = k | \beta, \phi, x_i) \propto \pi_k f(x_i | \phi_k)$$

▶ Update β_k by sampling from the conditional posterior

$$\beta_k \sim \text{Beta}(1 + m_k, \alpha_0 + \sum_{j>k} m_j)$$

▶ Update ϕ_k by sampling from the conditional posterior

$$p(\phi_k|z_{1:n}, x_{1:n}) \propto G_0(\phi_k) \prod_{i:z_i=k} f(x_i|\phi_k)$$

▶ One can monitor $\max_i z_i$ to verify that truncation at T is good enough, and increase T if necessary.

▶ We can also use truncated steak-breaking representation to form a mean field approximation of DP mixtures

$$q(\beta, \phi, z) = \prod_{i=1}^{n} q(z_i | w_i) \prod_{k=1}^{T} q(\phi_k | \tau_k) \prod_{k=1}^{T-1} q(\beta_k | \gamma_k)$$

▶ For a conjugate DP mixture in the exponential family

$$\gamma_{k,1} = 1 + \sum_{i=1}^{n} w_{i,k}, \quad \gamma_{k,2} = \alpha_0 + \sum_{i=1}^{n} \sum_{j>k} w_{i,j}$$
$$\tau_{k,1} = \lambda_1 + \sum_{i=1}^{n} w_{i,k} t(x_i), \quad \tau_{k,2} = \lambda_2 + \sum_{i=1}^{n} w_{i,k}$$
$$w_{i,k} \propto \exp(S_k)$$

where

$$S_k = \mathbb{E}\log\beta_k + \sum_{j \le k} \mathbb{E}\log(1-\beta_j) + \mathbb{E}\phi_k^T t(x_i) - \mathbb{E}A(\phi_k)$$



► The approximate predictive distribution given by variational inference at different stages of the algorithm. The data are 100 points generated by a Gaussian DP mixture model with fixed diagonal covariance.



Initial state



1st iteration



5th (and last) iteration



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