IFT 6269: Probabilistic Graphical Models

Lecture 26 — December 4

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**Disclaimer:** These notes have only been lightly proofread.

# **Bayesian Non-Parametrics**

The aim of this lecture is to briefly introduce Stochastic processes and dive a little deeper into Gaussian processes and Dirichlet processes. However, note that this lecture will not cover the implementational aspects and hyperparameter selection. For a more in-depth/complete understanding of these topics, the readers are referred to the works of David MacKay, Yee Whye Teh, Kilian Weinberger, Tamara Broderick, and Michael Jordan.

# 26.1 Stochastic Process

A stochastic process is defined as a collection of random variables defined on a common probability space  $(\Omega, \mathcal{F}, \mathbb{P})$ , where  $\Omega$  is a sample space,  $\mathcal{F}$  is a sigma-algebra, and  $\mathbb{P}$  is a probability measure; and the random variables, indexed by some set T, all take values in the same space S, measurable with respect to some  $\sigma$ -algebra.

Mathematically, it can be understood as: For every element  $t \in T$ , there is a random variable X (from a common probability space) indexed by t. Therefore, we have a simple definition as:

$$\{X(t) \mid t \in T\}$$

# 26.1.1 Examples of Stochastic Process

The following are some examples of stochastic processes:

- Single random variable: Consider any singleton set T, say  $\{1\}$  for instance. Then, the collection of random variables indexed by T is given by  $\{X_1\}$ .
- IID random variables: Consider the finite set  $\{1, 2, ..., n\}$  to be the indexing set T. The corresponding collection of random variables is given by  $\{X_1, X_2, ..., X_n\}$ , where  $X_i$  and  $X_j$  are independent and follow a common distribution  $X^*$ .
- Deterministic function: Let f be a function  $f: T \to S$ , given the index set T, the collection is defined as  $\{X(t) := \delta_{f(t)} \mid t \in T\}$ , where  $\delta$  is the Dirac distribution.
- Wiener process, Poisson process, etc.
- Gaussian process
- Dirichlet process

# 26.2 Gaussian Processes

The goal is to do inference based on the inputs X and targets t and use that for prediction when presented with an unseen test input  $x^*$ . Inference is essentially calculating the posterior distribution over the models  $y(\cdot)$  using Bayes' theorem:

$$\mathbb{P}(y(\cdot) \mid \boldsymbol{t}, \boldsymbol{X}) = \frac{\mathbb{P}(\boldsymbol{t} \mid y(\cdot), \boldsymbol{X}) \mathbb{P}(y(\cdot))}{\mathbb{P}(\boldsymbol{t} \mid \boldsymbol{X})}$$
(26.1)

where,  $\mathbb{P}(t \mid y(\cdot), X)$  is the likelihood,  $\mathbb{P}(y(\cdot))$  is the prior over functions, and  $\mathbb{P}(t \mid X)$  is the evidence. It is important to note that the prior  $\mathbb{P}(y(\cdot))$  cannot be easily defined because it has to cover an infinite-dimensional space.

The posterior distribution is further used for prediction as it allows us to obtain the predictive distribution  $\mathbb{P}(t^* \mid \boldsymbol{x}^*, \boldsymbol{t}, \boldsymbol{X})$  defined as:

$$\mathbb{P}(t^* \mid \boldsymbol{x}^*, \boldsymbol{t}, \boldsymbol{X}) = \int \mathbb{P}(t^* \mid y(\cdot), x^*) \mathbb{P}(y(\cdot) \mid \boldsymbol{t}, \boldsymbol{X}) dy$$
(26.2)

where,  $\mathbb{P}(y(\cdot) \mid \boldsymbol{t}, \boldsymbol{X})$  is computed from equation (26.1).

### 26.2.1 Gaussian Properties

We do a quick review of Gaussian properties as they are useful in the treatment of Gaussian processes. Consider the following setting where there are two random variables X, Y and they are Gaussian distributed with mean  $\mu$  and  $\Sigma$  as:

$$\begin{bmatrix} X \\ Y \end{bmatrix} \sim \mathcal{N} \left( \begin{bmatrix} \boldsymbol{\mu}_X \\ \boldsymbol{\mu}_Y \end{bmatrix}, \begin{bmatrix} \boldsymbol{\Sigma}_X & \boldsymbol{\Sigma}_{XY} \\ \boldsymbol{\Sigma}_{XY}^\top & \boldsymbol{\Sigma}_Y \end{bmatrix} \right)$$

The following are some of the important properties of Gaussian distributions:

**Property 26.2.1** (Tractable) Normalization -  $\int \tilde{p}(x) dx = (2\pi)^{d/2} |\Sigma|^{1/2}$ 

• Allows us to have a closed-form solution and is also easier for sampling.

Property 26.2.2 Marginalization -  $X \sim \mathcal{N}(\boldsymbol{\mu}_X, \boldsymbol{\Sigma}_X)$ 

• X is distributed according to a Gaussian with mean  $\mu_X$  and covariance  $\Sigma_X$ . Can be proved using the *characteristic function* of a Gaussian.

Property 26.2.3 Conditioning -  $X \mid Y = \boldsymbol{y} \sim \mathcal{N}(\boldsymbol{\mu}_X + \boldsymbol{\Sigma}_{XY}\boldsymbol{\Sigma}_Y^{-1}(\boldsymbol{y} - \boldsymbol{\mu}_Y), \boldsymbol{\Lambda}_X^{-1})$ 

• Where  $\Lambda := \Sigma^{-1}$ . Note that the conditional distribution is also a Gaussian.

Property 26.2.4 Addition -  $X + Y = \mathcal{N}(\boldsymbol{\mu}_X + \boldsymbol{\mu}_X, \boldsymbol{\Sigma}_X + \boldsymbol{\Sigma}_Y + 2\boldsymbol{\Sigma}_{XY})$ 

• Addition of two Gaussian random variables is also a Gaussian.

**Property 26.2.5** Product of densities -  $\mathcal{N}(x \mid \mu_1, \Sigma_1) \cdot \mathcal{N}(x \mid \mu_2, \Sigma_2) \propto \mathcal{N}(x \mid \mu_3, \Sigma_3)$ 

• Note: This is **not** a product of two Gaussian random variables, rather, it's a product of two Gaussian densities that results in an un-normalized Gaussian density.

### 26.2.2 Linear Regression

The goal is to perform inference on a linear model. We start with the following assumptions:

- Locations  $\{\boldsymbol{x}_i \mid i \in [1, N]\}$
- Basis functions  $\{\phi_h(\boldsymbol{x})\}_{h=1}^H$  some transformations of the input  $\boldsymbol{x}$  using H basis functions.
- Linear model  $y(\boldsymbol{\phi}(\boldsymbol{x}), \boldsymbol{w}) = \boldsymbol{w}^{\top} \boldsymbol{\phi}(\boldsymbol{x})$
- Prior  $\boldsymbol{w} \sim \mathcal{N}(\boldsymbol{0}, \sigma_w^2 \boldsymbol{I})$  a Gaussian defined over the weights  $\boldsymbol{w}$  with isotropic covariance.
- Observation noise  $t \mid y \sim \mathcal{N}(y, \sigma_{\nu}^2)$  typically defined as  $t = y + \sigma_{\nu} \epsilon$ , where y is the "true value" that is not observed directly.

Given these assumptions, we have that the joint distribution of the true values of the model  $\boldsymbol{y}$  is a Gaussian given by:  $\boldsymbol{y}_N \sim \mathcal{N}(\boldsymbol{0}, \sigma_w^2 \Phi_N \Phi_N^{\top})$  where  $\Phi_N$  is the  $n \times d$  design matrix where each  $\boldsymbol{\phi}(\boldsymbol{x}_i)$  is put as a row. Since we don't observe the true values directly but observe a noisy version of them, we arrive at a similar conclusion for the targets  $\boldsymbol{t}$  which is:  $\boldsymbol{t}_N \sim \mathcal{N}(\boldsymbol{0}, \sigma_w^2 \Phi_N \Phi_N^{\top} + \sigma_v^2 \boldsymbol{I}).$ 

**Remark 26.2.1** The term  $\Phi_N \Phi_N^{\top}$  is interesting because considering a value  $(\Phi_N \Phi_N^{\top})_{ij}$ , even before we observe the values of the noisy random variables, our prior encodes our belief that if features of datapoint i and features of datapoint j are similar, then the corresponding values of  $y_i$  and  $y_i$  will be close to each other.

**Remark 26.2.2** The covariance matrix  $(\sigma_w^2 \Phi_N \Phi_N^\top + \sigma_\nu^2 I)$  can be generalized to any positive semi-definite (PSD) similarity matrix. This has deep connections to the theory of kernel regression.

Note: From here on, the similarity matrix is represented using C wherein  $C(x_i, x_j)$  refers to the similarity between  $X_i$  and  $X_j$ .

### 26.2.3 Definition of a Gaussian Process

The probability distribution of a function  $y(\boldsymbol{x})$  is a Gaussian process if for any finite selection of points  $(\boldsymbol{x}_1, \boldsymbol{x}_2, \ldots, \boldsymbol{x}_N)$ , the vector  $[y(\boldsymbol{x}_1), y(\boldsymbol{x}_2), \ldots, y(\boldsymbol{x}_N)]$  follows a Gaussian distribution. Recalling what we defined earlier for stochastic processes, we have that:

- Index set T  $\mathcal{X}$  the set of possible inputs  $\boldsymbol{x}$
- Value space S  $\mathbb{R}^l$  (can also be multi-dimensional)
- Random variables Gaussians (it's essentially the function values acting as random variables for checking the distribution they follow)

Now, given all this information, we finally observe the targets  $t_N = \{t_i \mid i \in [1, N]\}$ and would like to infer a new target  $t_{N+1}$  for a new test point  $x_{N+1}$ . The similarity matrix for a new datapoint is given by:

$$\boldsymbol{C}_{N+1} = \begin{bmatrix} \boldsymbol{C}_N & \boldsymbol{k} \\ \boldsymbol{k}^\top & \boldsymbol{\kappa} \end{bmatrix}$$
(26.3)

where,  $C_N$  is the collection of all the similarity values for the N datapoints in the space, an element in  $\mathbf{k}$ , say  $k_j$ , is the similarity value between *j*-th and the new datapoint  $X_{N+1}$ , i.e.  $k_j = C(X_{N+1}, X_j)$ , and  $\kappa$  is the similarity value between the new datapoint with itself, i.e.  $C(X_{N+1}, X_{N+1})$ .

The predictive distribution is now given by:

$$\mathbb{P}(t_{N+1} \mid \boldsymbol{t}_N) = \frac{\mathbb{P}(t_{N+1}, \boldsymbol{t}_N)}{\mathbb{P}(\boldsymbol{t}_N)} \propto \exp\left[\frac{-1}{2} \begin{bmatrix} \boldsymbol{t}_N & t_{N+1} \end{bmatrix} \boldsymbol{C}_{N+1}^{-1} \begin{bmatrix} \boldsymbol{t}_N \\ t_{N+1} \end{bmatrix}\right]$$
(26.4)

For notational simplicity, the dependence on  $\Phi_N$  is omitted here. Note that this is only a distribution over  $t_{N+1}$ . Upon further simplification we have the following result:

$$t^* \mid t_1, \dots, t_N, x_1, \dots, x_n, x^* \sim \mathcal{N}(k^\top C_N^{-1} t_N, \kappa - k^\top C_N^{-1} k)$$
(26.5)

**Remark 26.2.3** The dependence is on  $C_N$  (and not  $C_{N+1}$ ). So, for prediction over multiple test points, we only need to calculate the inverse of the similarity matrix over the data space  $C_N$  once, augment the corresponding  $\mathbf{k}$  for all the new test points and then compute the Gaussian with mean and variance (using simple linear algebra).

**Remark 26.2.4** The posterior mean is linear w.r.t  $\mathbf{t}_N$  weighted by the term  $\mathbf{k}^\top C_N^{-1}$ . For instance, if we have positive correlation between elements, then a higher target value would correspond to a new test point distribution with higher mean.

Note: A demo showing the working of a Gaussian process can be found at the following link: http://chifeng.scripts.mit.edu/stuff/gp-demo/

#### 26.2.4 GP Summary

Problem: y is a function i.e. "an infinite-dimensional vector". But the multivariate Gaussian distribution is only defined for **finite** dimensional vectors.

Definition: A GP is a (potentially infinite) collection of random variables such that the joint distribution of **every finite subset** of them is a multivariate Gaussian.

If we start with the assumption that the function values are distributed according to a Gaussian prior i.e.  $y(X_1), \ldots, y(X_N) \sim \mathcal{N}(\mathbf{0}, \mathbf{C}_N)$ , then the posterior distribution is also a Gaussian of the form:

$$t^* \mid t_1, \ldots, t_N, x_1, \ldots, x_n, x^* \sim \mathcal{N}(k^\top C_N^{-1} t_N, \kappa - k^\top C_N^{-1} k)$$

# 26.3 Dirichlet Processes

Considering the generative modelling perspective, we define our model (say, a latent variable model) as follows:

- Assignments  $z_n \mid \boldsymbol{\rho} \sim \text{Categorical}(\rho_1, \dots, \rho_K)$  the class assignments are obtained from a Categorical distribution with  $\rho_i \geq 0$  and  $\sum_i \rho_i = 1$ .
- Class conditionals  $\boldsymbol{x}_n \mid z_n \sim \mathcal{N}(\boldsymbol{\mu}_{z_n}, \boldsymbol{\Sigma}) = F_{z_n}$  the class conditionals depend on a specific value  $F_{z_n}$  of the latent variable.

We also make some prior assumptions as follows:

- Number of classes K is known.
- Class proportions  $\rho \sim \text{Dir}(\alpha_1, \ldots, \alpha_K)$  following a Dirichlet distribution.
- Class parameters  $\boldsymbol{\mu}_k \sim \mathcal{N}(\boldsymbol{\mu}_0, \boldsymbol{\Sigma}_0) = H$  the parameters defining the behaviour of  $F_{z_n}$  are themselves sampled from a Gaussian with fixed mean and covariance, called H.

### 26.3.1 Dirichlet Distribution

The Dirichlet distribution is formally defined as:

$$\mathbb{P}(\rho_1, \dots, \rho_K \mid \alpha_1, \dots, \alpha_K) = \frac{\Gamma(\Sigma_k \alpha_k)}{\prod_k \Gamma(\alpha_k)} \prod_{k=1}^K \rho_k^{\alpha_k - 1}$$
(26.6)

**Remark 26.3.1** The Dirichlet distribution can be thought of as a distribution over distributions. This is because whenever we sample a random variable from this distribution, we get a collection of  $\rho_1, \ldots \rho_K$  which in turn specifies a Categorical distribution. Note that  $\rho_1, \ldots \rho_K$ belong to a simplex of K - 1 dimensions.

**Remark 26.3.2** The Dirichlet distribution is a direct extension of the Beta distribution defined only over  $\rho_1, \rho_2$  and  $\alpha_1, \alpha_2$ .

An important property of the Gamma distribution is given below. It behaves similar to the factorial function.

#### Property 26.3.1

$$\Gamma(z+1) = z\Gamma(z) \qquad where, \quad z \in \mathbb{C}$$
(26.7)

**Property 26.3.2** Collapsibility property aka Aggregation

If  $\boldsymbol{\rho} = (\rho_1, \dots, \rho_K) \sim Dir(\alpha_1, \dots, \alpha_K)$ , then if the random variables at the *i*-th and *j*-th index are dropped and replaced with their sum, the new distribution is also Dirichlet with:

$$\boldsymbol{\rho}' = (\rho_1, \dots, \rho_i + \rho_j, \dots, \rho_K) \sim Dir(\alpha_1, \dots, \alpha_i + \alpha_j, \dots, \alpha_K)$$
(26.8)

It can be useful in deriving the marginal distribution of  $\rho_i$  from the joint above.

#### 26.3.2 Understanding the influence of Dirichlet parameters

Let us consider the following extension of the collapsibility property where we fix a  $\rho_i$  and sum over all others:

$$\left(\rho_i, \sum_{j \neq i} \rho_j\right) \sim \operatorname{Dir}\left(\alpha_i, \sum_{j \neq i} \alpha_j\right)$$

Recall that if we have a collection of random variables following a Dirichlet distribution, then each random variable follows a Beta distribution i.e.  $\rho_i \sim \text{Beta}(\alpha_i, \alpha_0 - \alpha_i)$ , where,  $\alpha_0 = \sum_i \alpha_i$ . Likewise, we can also compute the mean, variance and the covariance as:

$$\mathbb{E}[\rho_i] = \frac{\alpha_i}{\alpha_i + \alpha_0 - \alpha_i} = \frac{\alpha_i}{\alpha_0}$$
(26.9)

$$\operatorname{Var}(\rho_i) = \frac{\alpha_i(\alpha_0 - \alpha_i)}{\alpha_0^2(\alpha_0 + 1)}$$
(26.10)

$$\operatorname{Cov}(\rho_i, \rho_j) = \frac{-\alpha_i \alpha_j}{\alpha_0^2(\alpha_0 + 1)}$$
(26.11)

**Remark 26.3.3** Note that in equation 26.10 if  $\alpha_i$  is too small or if  $\alpha_i \approx \alpha_0$  the variance will be small (i.e.  $\approx 0$ ). Similarly, considering the scale of the parameters, if all the parameters are large, the denominator becomes much larger and hence the variance will be small.

**Remark 26.3.4** The covariance in equation 26.11 is negative to account for the fact that the random variables lie in a probability simplex (summing up to 1), so, intuitively, if one of the variables gets large then the others have to be small for this to be true.

Figure 26.1 shows an example of how different values of Dirichlet parameters determine the spread of the probability density function (pdf) on the 2-simplex. In the top-left image,  $\mathbb{E}[\rho_i] = 0.33$  and  $\alpha_0 = 4.5$  showing concentric shapes for each random variable. In the topright image,  $\alpha_0 = 15$  which shows that the variance is small hence the concentric shapes are concentrated in the middle of the simplex. For the bottom-right image,  $\mathbb{E}[\rho_3] = \frac{8}{14} = 0.57$ which explains the skewness towards the random variable  $\rho_3$ .

### 26.3.3 Dirichlet Simulation

The goal here is to outline a mechanism for sampling from a Dirichlet distribution. We have:

$$\rho_1 \sim \text{Beta}(\alpha_1, \sum_{k=1}^K \alpha_k - \alpha_1) \quad \perp \quad \frac{(\rho_2, \dots, \rho_K)}{1 - \rho_1} \sim \text{Dir}(\alpha_2, \dots, \alpha_K)$$
(26.12)

$$\nu_2 \sim \text{Beta}(\alpha_2, \sum_{k=2}^{K} \alpha_k - \alpha_2) \qquad \rho_2 = (1 - \nu_1)\nu_2 \qquad (26.13)$$



Figure 26.1: Probability density functions for the Dirichlet distribution on the 2-simplex. Source: Dirichlet distribution wikipedia.

:  

$$\nu_l \sim \text{Beta}(\alpha_l, \sum_{k=l}^K \alpha_k - \alpha_l) \qquad \rho_l = \left(\prod_{k=1}^{l-1} (1 - \nu_k)\right) \nu_l \qquad (26.14)$$

$$\rho_K = 1 - \sum_{k=1}^{K-1} \rho_k \tag{26.15}$$

**Remark 26.3.5** Consider equation 26.12 where  $\rho_1$  is sampled from a Beta distribution. If the distribution is re-normalized excluding the most recent sample ( $\rho_1$  in this case), we again obtain a Dirichlet distribution without the corresponding parameter  $\alpha_1$  for the random variable  $\rho_1$ . In other words, the re-normalized distribution, once  $\rho_1$  is observed, only depends on the values of the remaining random variables.

**Remark 26.3.6** Considering equation 26.13, note that  $\nu_2$  denotes the sample from the renormalized distribution (without  $\rho_1$ ). Therefore,  $\rho_2$  is obtained by multiplying  $\nu_2$  with the remaining portion denoted by  $(1 - \nu_1)$ .

Therefore, we have the following outline where we sample from a Beta distribution, renormalize it, sample another variable from the updated distribution, re-normalize it, and so on until we reach K - 1 random variables. The K-th can then be sampled by subtracting all the previously sampled variables from 1 (equation 26.15).

Note: This is also analogous to the stick-breaking mechanism where, with each sampled variable we chop off a certain portion of the stick and continue sampling from the remaining portion.

#### **26.3.4** How do we choose K?

We don't! Instead, we directly consider an infinite-dimensional mixture model. The goal here is to extend our knowledge of sampling from a finite Dirichlet to an infinite setting when we don't know K to begin with.

The solution is to keep continuing the "stick-breaking" process till infinity i.e.

$$\nu_{1} \sim \text{Beta}(\alpha_{1}, \beta_{1}) \qquad \rho_{1} = \nu_{1}$$

$$\nu_{2} \sim \text{Beta}(\alpha_{2}, \beta_{2}) \qquad \rho_{2} = (1 - \nu_{1})\nu_{2}$$

$$\vdots \qquad \vdots$$

$$\nu_{l} \sim \text{Beta}(\alpha_{l}, \beta_{l}) \qquad \rho_{l} = \left(\prod_{k=1}^{l-1} (1 - \nu_{k})\right) \nu_{l}$$

However, the values for  $\alpha_l$  and  $\beta_l$  must be chosen carefully for proper normalization as they dictate whether the portion of the stick chopped off is too small or too large.

Towards that end, a method proposed by Griffiths, Engen and McCloskey (GEM) is to use the following strategy:

$$\nu_l = \text{Beta}(1, \alpha) \qquad \rho_1, \rho_2, \ldots \sim \text{GEM}(\alpha)$$

**Remark 26.3.7** Using the GEM strategy, we can observe that  $\mathbb{E}[\nu_l] = \frac{1}{1+\alpha}$ , where  $\mathbb{E}[\nu_l] \approx 1$  if  $\alpha \ll 1$  and  $\mathbb{E}[\nu_l] \approx 0$  if  $\alpha \gg 1$ .

#### 26.3.5 Random Measures

We have an infinite collection of proportions  $\rho_1, \rho_2, \ldots \sim \text{GEM}(\alpha)$  and a distribution over the parameters of the distribution  $\phi_1, \phi_2, \ldots \overset{\text{iid}}{\sim} H$  (e.g.  $\mathcal{N}(\boldsymbol{\mu_0}, \boldsymbol{\Sigma_0})$ ), which comprise two sources of randomness. Recall the generative modelling setting in the beginning of section 26.3 where we had the parameters of the distribution specified by a fixed mean depending on the class and a shared covariance matrix. So, we have  $S \in \mathbb{R}^d \times \{\boldsymbol{\Sigma}\}$ , with each point in this space being a vector that is parametrized by the mean (for each class) and the shared covariance matrix. We then define a *random measure* over the space S:

$$G = \sum_{k=1}^{\infty} \rho_k \delta_{\phi_k} \tag{26.16}$$

Its interpretation is as follows: we sample infinitely many proportions  $\rho$ 's and the parameters of the distribution  $\phi$ 's according to the distribution H (which is also a distribution on the space S) and define a random measure such that for each location  $\phi_k$  in the space S, a Dirac distribution is assigned with a height (given by the corresponding proportion  $\rho_k$ ) that is associated with the location  $\phi_k$ .

The sampling is then done as follows:

$$z_n \mid G \sim G \tag{26.17}$$

$$\boldsymbol{x}_n \mid \boldsymbol{z}_n \sim \boldsymbol{F}_{\boldsymbol{z}_n} \tag{26.18}$$

Equation 26.17 tells that once we have defined the random measure G, we sample the class assignments from G and note that realizations of  $z_n$  are nothing but  $\phi_k$ 's parameterized by  $\mathcal{N}(\boldsymbol{\mu}_k, \boldsymbol{\Sigma})$ . Given the class assignments  $z_n$ , we can then obtain the observations by sampling from F, which depends on  $z_n$  by  $\mathcal{N}(\boldsymbol{\mu}_{z_n}, \boldsymbol{\Sigma})$ .

#### Why is G a random measure?

First, note that G is also a distribution on S. Consider a subset of the space S, say A, and apply the random measure G on the subset A denoting it as  $G(A) \in [0, 1]$ . Now, we see that for any subset in S, the result of applying this measure on that subset will be random because G itself is a random measure with  $\rho$ 's and  $\phi$ 's being its sources of randomness. Since G(A) is random, we can analyze it further by calculating its expectation  $\mathbb{E}_G[G(A)]$ .

$$\mathbb{E}_{G}[G(A)] = \mathbb{E}_{G}\left[\sum_{k} \rho_{k} \delta_{\phi_{k}}(A)\right]$$
$$= \mathbb{E}_{\rho}\left[\sum_{k} \rho_{k} \mathbb{E}_{\phi_{k}}[\delta_{\phi_{k}}(A)]\right]$$
$$= \mathbb{E}_{\rho}\left[\sum_{k} \rho_{k} H(A)\right] \quad (\because \delta_{\phi_{k}}(A) = 1 \text{ if } \phi_{k} \text{ lies in } A)$$

This is because the  $\mathbb{E}_{\phi_k}[\delta_{\phi_k}(A)]$  is the probability that  $\phi_k$  belongs to A. But,  $\phi$ 's are defined according to H, therefore, the probability *measure* w.r.t which it is obtained is H.

= H(A) (:: H(A) is independent of  $\rho$ )

**Remark 26.3.8** This result tells us that whatever random measures G we obtain by creating subsets on the space S, their expectation always results in the probability measure H being applied to the subsets on the space S. In other words, the probability that a random measure assigns to a subset in expectation is same as the probability that H assigns to the subset.

### 26.3.6 Definition of a Dirichlet Process

Given a measurable space  $(S, \mathcal{M})$ , a base probability distribution H and  $\alpha > 0$ , the probability distribution of a (random) measure G is a Dirichlet process if for any finite partition  $\{A_1, \ldots, A_r\}$  of S, the vector  $[G(A_1), \ldots, G(A_r)] \sim \text{Dir}(\alpha H(A_1), \ldots, \alpha H(A_r))$ . As for every stochastic process, we have:

- Random variables Dirichlet (every component of random vector arising from a finite partition of the space S will follow a Dirichlet distribution)
- Index set T  $\mathcal{M}$  (the "measureable" subsets of the space)

Exercise to the Reader: Explore the behavior of a Dirichlet process  $DP(\alpha, H)$  when: (i)  $\alpha \to 0$ ? and (ii)  $\alpha \to \infty$ ? This should also lead you to understand why  $\alpha$  is known as the concentration parameter.

### 26.3.7 DP Summary

Problem:  $\rho_1, \rho_2, \ldots$  is an infinite-dimensional probability vector. But, the Dirichlet distribution is defined for **finite** dimensional spaces.

Definiton: A DP is a distribution over probability measures such that for **every finite partition**, the probabilities of the partition elements follow a joint Dirichlet distribution.

$$[G(A_1),\ldots,G(A_r)] \sim \operatorname{Dir}(\alpha H(A_1),\ldots,\alpha H(A_r))$$