

# Probabilistic Generative models

## Week 1 - Foundations

### Setup

A PGM, or simply a GM, is a methodology for generating data. In general, the GM is constructed and adjusted using observations with the aim to generate data with the same statistical properties of the observations. The 'probabilistic' nature of the PGMs considered follow from the fact that the data generated will be considered to be realisations of a random variable; e.g.  $X$ .

In this sense, the probability distribution of  $X$ , denoted  $P_X(x)$ , as well as its density  $p_X(x)$  will be central to the study of PGM. In particular, targeting the pdf is one way of constructing PGMs, in which case the whole PGM paradigm becomes equivalent to the classical statistical model approach. However, as we will see in the course, enforcing the sought-after PGM to have an explicit parametric pdf can be rather restrictive.

Throughout the course, we will consider a probability space  $(\Omega, \mathcal{F}, P)$  with 3 RVs given by the measurable maps

$$\begin{array}{ccc} X: \Omega \rightarrow \mathcal{X} & ; & Y: \Omega \rightarrow \mathcal{Y} & ; & Z: \Omega \rightarrow \mathcal{Z} \\ \text{(observed} & & \text{observed} & & \text{(latent} \\ \text{input)} & & \text{output)} & & \text{variable)} \end{array}$$

Remark: Not all three variables will be present in all our models. For instance, in classification there's no justification for  $Z$ , while in clustering there's no need for  $X$ . However, we build the general set up here for formality.

We will also consider the  $\sigma$ -algebra  $\mathcal{F}$  to be the product Borel  $\sigma$ -algebra.  $\mathcal{F} = \mathcal{B}(X) \otimes \mathcal{B}(Y) \otimes \mathcal{B}(Z)$ .

Furthermore, we will assume the joint probability of  $X$ ,  $Y$  and  $Z$  has a density, that is,  $\forall A \in \mathcal{B}(X), B \in \mathcal{B}(Y), C \in \mathcal{B}(Z)$ , we have

$$P(X \in A, Y \in B, Z \in C) = \int_{A \times B \times C} p(x, y, z) dx dy dz$$

We will also assume that all marginals and conditionals, have a density. This includes  $p(x, y)$ ,  $p(y|x)$ ,  $p(z|x, y)$  etc.

→ Discriminative v/s Generative. The generative approach aims to characterise the complete generative distribution  $p(x, y, z)$ , whereas in some application specific cases only the discriminative model  $p(y|x)$  is needed. Let's see the following example.

Example: generative v/s <sup>binary</sup> discriminative classification.

Consider the <sup>bi</sup> classification problem, where given an observation  $X=x$ , one needs to estimate its label  $Y$ . A discriminative model would directly construct  $p(Y|X=x)$ . Since this is a binary case, without loss of generality we can assume  $Y \in \{0, 1\}$ , and model  $P(Y=1|X=x)$ , since  $P(Y=0|X=x) = 1 - P(Y=1|X=x)$ . A model for this will only need to map  $x \in \mathbb{R}^n \rightarrow P(Y=1|X=x) \in [0, 1]$ . For instance

$$P(Y=1|X=x) = \frac{1}{1 + e^{-\theta^T x}}$$

which is known as the logistic regression.

Conversely, in a generative approach, we aim to model the joint probability  $p(Y=1, X=x)$ .

Modelling this distribution directly is not easy, however, we can observe that it can be factorised as

$$p(Y=1, X=x) = p(X=x|Y=1) p(Y=1)$$

which yields two much more intuitive distributions:

- the class prob  $p(Y=1) \rightarrow \pi, (1-\pi)$
- the class-conditional prob  $p(X=x|Y=1) \rightarrow$  choose...  
 $f_{0_1}, f_{0_2}$

Therefore, the classifier is

$$p(Y=1|X=x) = \frac{p(X=x|Y=1) p(Y=1)}{p(X=x)}$$

$$= \frac{1}{1 + \frac{p(X=x|Y=0)p(Y=0)}{p(X=x|Y=1)p(Y=1)}}$$

$$= \frac{1}{1 + e^{\log\left(\frac{1-\pi}{\pi} \cdot \frac{f_{0_0}(x)}{f_{0_1}(x)}\right)}} \quad (*)$$

Exercise: evaluate  $*$  for  $f_{0_1} = N(\mu_1, \Sigma), f_{0_0} = N(\mu_2, \Sigma)$

→ Push forward measure

Despite the abundant collection of well studied statistical models, in some scenarios we can construct a more ad hoc model by applying an appropriate transformation.

Def: Consider a RV  $X \in \mathcal{X}$  with measure  $P_X$ , and a non linear map  $T: \mathcal{X} \rightarrow \mathcal{X}$ . The measure of the transformed RV  $Y = T(X)$  is known as the pushforward measure of  $P_X$  through  $T$  and it is denoted by  $P_{X \# T}$ .

Remark: The transformations considered in the course will be s.t. the pushforward measure has a density, denoted  $P_{X \# T}$

Example 1: discrete push fwd

Example 2: continuous push fwd

Theorem: Change of variable. Consider  $X, Y \in \mathbb{R}^d$  R.V.s such that  $Y = f(X)$  with  $f: \mathbb{R}^d \rightarrow \mathbb{R}^d$  diff+invertible. If  $X, Y$  have densities  $P_X$  and  $P_Y$ , then

$$P_Y(y) = P_X(f^{-1}(y)) |\det f^{-1}(y)|$$

Remark: Though the above result provides a closed-form expression for the push-forward measure only in the invertible case, a new law  $P_Y$  can be induced for any measurable map. Furthermore, if  $f$  is 'locally invertible' and 'not too wild'  $P_Y$  has a density.

This is because

$$P(Y \in A) = \sum_{f(B_i) \in A} P(X \in B_i)$$

## - Maximise likelihood.

ML is going to be the canonical methodology for training our PGMs, and as we will see, it will recover other forms of training criteria in particular cases.

Consider a PGM for the RV  $Y$ , with density  $p_\theta(y)$ , where  $\theta \in \Theta$  is the model parameter. Also, observing realisations  $y_1, \dots, y_n$ .

Def: likelihood function. is the probability density of the data, given the model:

$$L(\theta) = p_\theta(y_1, \dots, y_n)$$

NB. Here, we abused notation stating the joint likelihood

Def: the ML estimator is

$$\theta_{ML} = \operatorname{argmax}_{\theta} L(\theta)$$

Remark. In general, we will be dealing with i.i.d samples from  $Y$ , in such cases note that

$$L(\theta) = p(y_1, \dots, y_n) = \prod_i p(y_i).$$

Furthermore, we consider the training objective

$$l(\theta) = \log L(\theta) = \sum_i \log p(y_i)$$

Example: Gaussian linear regression

Let us consider the PGM given by

$$y|x \sim N(ax, \sigma^2),$$

This is equivalent to  $y = ax + \epsilon$ ,  $\epsilon \sim N(0, \sigma^2)$ . The params here are  $a$  and  $\sigma^2$ . Consider obs  $(y_1 \dots y_n)$  at inputs  $(x_1 \dots x_n)$ .

$$L(\theta) = p(y_1 \dots y_n | x_1 \dots x_n) = \prod_i p(y_i | x_i)$$

where  $p(y_i | x_i) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2} (y_i - ax_i)^2\right)$ . Thus

$$l(\theta) = \sum_i -\frac{1}{2} \log(2\pi\sigma^2) + \frac{1}{2\sigma^2} (y_i - ax_i)^2 \quad (*)$$

→ optimal  $a$  and  $\sigma^2$

Remark: Recall the (\*): ML recovers least sq

Example (binary classification). Consider observations  $x_i \in \mathbb{R}^d$  and labels  $y_i \in \{0, 1\}$ . The classifier will be modelled by

$$p_\theta(y=1|x) = \sigma(s(x))$$

where:  $\sigma(s(x)) = \frac{1}{1+e^{-s(x)}}$ ; and  $s(\cdot)$  is a feature extractor (it could be linear)

Assuming i.i.d. (Bernoulli)

$$L(\theta) = \prod_i p(y_i | x_i) = \prod_i \sigma(s(x_i))^{y_i} (1 - \sigma(s(x_i)))^{1-y_i}$$

and

$$l(\theta) = \sum_i y_i \log \sigma(s(x_i)) + (1-y_i) \log (1 - \sigma(s(x_i)))$$

\* Cross entropy



Example: clustering. Let us consider a set of observations  $D = \{x_i\}_{i=1}^n$  and implement a clustering operation. We will assume that there are  $K$  clusters, each specified by a density  $p_k$  and that the prob of  $x$  coming from  $p_k$  is  $\pi_k$ ; where  $0 \leq \pi_k \leq 1$  and  $\sum_k \pi_k = 1$ . the prob of  $x$  is

$$p(x) = \sum_k \pi_k p_k(x) \quad (*)$$

and  $l(\theta) = \sum_i \log \sum_k \pi_k p_k(x_i)$ . [Say what parameters are]

Note that this has two probs.

i) this doesn't give the cluster assignments (though that can be solved by hand)

ii) optimising  $(*)$  is ill posed: if  $p_k = N(\mu_k, \Sigma_k)$  we can set  $\mu_k = x_i, \Sigma_k = 0$  for  $k = \infty$ .

We can sort this out introducing a latent variable that represent the cluster assignment, namely

$$z_{nk} = 1 \text{ iff } x_n \in C_k; \quad r_{nk} = P(z_{nk} = 1)$$

With this  $p(x_n | z_n) = \prod_k p_k^{r_{nk}}$

we have the "complete-data likelihood"

$$l(\theta) = \log \prod_n \prod_k p_k^{z_{nk}} = \sum_n \sum_k z_{nk} \log p_k(x_n)$$

Which is now optimisable but impractical, since we don't have the  $z$ 's

Workaround: estimate  $z_{nk}$  by  $r_{nk} = P(z_{nk} = 1 | x_n, \theta)$  which is equal to  $E(r_{nk} | x_n, \theta) \rightarrow$  can't do ML here.

# Some properties of ML

- Consistency
- Equivariance
- Asymptotic normality
- Asymptotic efficiency

## Intro to information theory

Motivation. Let us consider a RV  $X$  with  $P_X$ . Observe that  $-\log p_X(x)$  is represent a measure of information gained from obtain the value  $x$  as a sample of  $X$ . or consider a communication channel  $\mathcal{X}^A \rightarrow \mathcal{X}^B$  where  $A$  is transmitting samples from  $X$  to  $B$ .

$B$  sees samples from  $X$ , so it's average 'surprise' is

$$H(X) = - \int_{\mathcal{X}} p(x) \log(p(x)) dx \quad (\text{later, this})$$

$$H(x) = - \sum_x p(x) \log(p(x)) \geq 0$$

• Properties: (Jensen:  $\mathbb{E}(\log(p(x))) \leq \log \mathbb{E}(p(x))$ )

$$\textcircled{1}. H(X) = \sum p(x) \log\left(\frac{1}{p(x)}\right) \leq \log \sum 1 = \log K$$

$$\text{and } H(X_U) = \sum \frac{1}{K} (\log(K)) = \log K$$

$$\textcircled{2} \text{ if } \exists i \text{ s.t. } p(x_i) = 1 \quad p(x_{\neq i}) = 0 \rightarrow H(x) = 1 \cdot \log(1) = 0$$



Bernoulli example

$$X \sim p(x) = \theta^x (1-\theta)^{1-x}$$

$$H(X) = -\theta \log \theta + (1-\theta) \log (1-\theta)$$

→ show plot.

- The entropy, in addition to the measure of disorder, can be understood as the cost of the optimal form of compression. Think of a compression strategy using symbols  $s_1, \dots$ , where with increasing size. E.g.  $s_1 = 0$   $s_2 = 1$   $s_3 = 10$   $s_4 = 11$   $\vdots$   
the strategy would be to assign  $s_i$  to the  $i$ th most frequent.  
therefore  $H(X) = \sum_i p(x) \log \left( \frac{1}{p(x)} \right)$  is the average message size of the code.

Now assume a communication channel  $A \rightarrow B$  where  $B$  receives samples  $X \sim p$  but mistakenly thinks that  $X \sim q$ . His assessment of info would be

$$H_{CE}(p, q) = - \sum \log(q(x_i)) p(x_i). \rightarrow \text{this is also the cost of compressing } X \sim p \text{ via } q$$

Discussion. Would  $H(p, q)$  greater or smaller than  $H(p, p) = H(p)$ ?

$$\text{Let's see: } H(p) - H(p, q) = \sum p \log\left(\frac{q}{p}\right) \\ \leq \log \sum p \cdot \frac{q}{p} = \log 1 = 0$$

$\Rightarrow H(p) \leq H(p, q)$  with e.g. attainable only for  $p=q$

therefore minimising crossentropy wrt. to one arg. is precisely an attempt to match  $p=q$ .

The notion of entropy/crossentropy can be extended to continuous RVs with densities via

$$\cdot H(x) = \int p(x) \log(p(x)) dx$$

$$\cdot H(p, q) = \int p(x) \log(q(x)) dx$$

Remark:  $H(x)$  can be positive, negative or zero.

$$H(U_{(0,1)}) = \int_0^1 \frac{1}{x} \log\left(\frac{1}{x}\right) dx = \log(e) = 1. \text{ However, } H(p, q) \neq H(p)$$

Quantifying this discrepancy gives

$$KL(p||q) = H(p, q) - H(p) = \sum p \log\left(\frac{p}{q}\right) \geq 0$$

KL is a divergence measure, that is, a function that quantifies how far is  $p$  from  $q$ , that is always positive and  $D(p, q) = 0$  iff  $p = q$ . (a.e.) - identity of the indiscernibles.

• However, note that

- KL is not symmetric

- Does not have  $\Delta$  inequality.

- it's only defined when  $p \gg q$

KL as a metric to compare  $p$  and  $q$

In the continuous case, it is interesting to understand what type of convergence KL gives

• Let us consider other two divergences

$$L_1: D_{L_1}(p, q) = \int |p - q| dx$$

$$X^2: D_{X^2}(p, q) = \int \frac{|p - q|^2}{q} dx$$

$$\text{Scenario 1 } p = U_{[0,1]}, \quad q = \begin{cases} e^{-n} & x \in [0, 1/n] \\ c_n & x \in (1/n, 1) \end{cases}$$

$$D_{L_1}(p, q) = \int_0^{1/n} |e^{-n} - 1| dx + \int_{1/n}^1 |c_n - 1| dx = \frac{1 - e^{-n}}{n} + \frac{|c_n - 1|}{\frac{n-1}{n}} \xrightarrow[n \rightarrow \infty]{} 0$$

$$KL(p, q) = \int p(x) \log \frac{p(x)}{q(x)} = \int_0^{1/n} -\log e^{-n} + \int_{1/n}^n -\log C_n = n \cdot \underbrace{\frac{1}{n}}_{\rightarrow 0} + \underbrace{\frac{-n}{n-1} \log C_n}_{\rightarrow 0} = KL \text{ not conv.}$$

Case 2  $p = (1-\epsilon, \epsilon)$ ,  $q = (1-\epsilon^2, \epsilon^2)$  (Bernoullis)

$$KL = \sum p_i \log \left( \frac{p_i}{q_i} \right) = \underbrace{(1-\epsilon) \log \left( \frac{1-\epsilon}{1-\epsilon^2} \right)}_{\rightarrow 0} + \underbrace{\epsilon \log \left( \frac{\epsilon}{\epsilon^2} \right)}_{\rightarrow -\infty} \rightarrow 0 \quad (L'Hôpital)$$

$$\chi^2: \frac{\|1-\epsilon - 1+\epsilon^2\|^2}{1-\epsilon^2} + \frac{\|\epsilon - \epsilon^2\|^2}{\epsilon^2} = \frac{\|\epsilon^2 - \epsilon\|^2}{1-\epsilon^2} + \|1-\epsilon\|^2 \rightarrow 1$$

This means that, even though KL provides a stronger sense of convergence than  $L_1$ , there are 'stronger' divergences.

• Direct  $\forall$ s reverse KL.

Since  $KL(p||q)$  is not symmetric, it makes sense to study  $KL(q||p)$ , some observation.

Since  $p \gg q$  is needed for direct  $KL(p||q)$ , we have that  $KL(p||q)$ , as a metric to penalise  $q$  as an approximation of  $p$ , will promote  $q$  covers all the support of  $p$ .

Example:  $KL(N(\mu_0, \sigma_0) || N(\mu_1, \sigma_1)) = \log \frac{\sigma_1}{\sigma_0} + \frac{\sigma_0^2}{2\sigma_1^2} (\mu_0 - \mu_1)^2 - 1/2$

When  $\mu_0 = \mu_1$ ;  $KL = \log \frac{\sigma_1}{\sigma_0} + \frac{\sigma_0^2}{2\sigma_1^2} - \frac{1}{2}$  Recall that " $\sigma_0^2$ " is the true param. and that  $\sigma_1 = \sigma_0$  achieves  $KL = 0$ , However let's see what's worse:  $\sigma_1 > \sigma_0$  or  $\sigma_1 < \sigma_0$ .

Plot  $KL = \log \frac{\sigma_1}{\sigma_0} + \frac{\sigma_0^2}{2\sigma_1^2} - \frac{1}{2}$  for  $\sigma_0^2 = 1$  as a fn of  $\sigma_1$   
(and vice versa)

Plot: Consider  $p = \text{GMM}_1$ ,  $q = \text{GMM}_2$ ; plot the optimal KL.

## KL and Maximum Likelihood.

Let's now turn to our setting of a learning problem. Consider a PM given by  $p_0(x)$  and iid observations  $x_1 \dots x_n \sim p(x)$ . We could use the KL as a metric to adjust  $p_0$  to  $p$ . That is

$$\Theta = \arg \min KL(p || p_\Theta)$$

This is unfeasible in practice, since  $p$  is not known. However, we

$$\Theta = \arg \min \int \log p(x) dx - \int \log q_\Theta(x) dx$$

$$= \arg \max \mathbb{E}_p \log q_\Theta$$

$$= \arg \max \sum_{x_i} \log q_\Theta(x_i)$$

$\Rightarrow$  minimizing KL is equivalent to ML (large data)

As we'll see in detail during the module when designing a PGM, we have the following options:

- Closed-form likelihood based models. These are classical statistical models that we can choose, or design for, first principles.
  - $N$ , cat, expo, even half-normal, log normal.
- likelihood based push forward models. This is a transformation of a simple base (or 'source') model via a learned map. The params of the map appear explicitly in the likelihood. For instance take a base measure  $p(z)$  and model  $x = T_\theta(z)$ , with  $z \sim \text{Gaussian}$ .

$$\text{this way } p_x = p_z(T^{-1}(x)) |\det \nabla T^{-1}(x)|$$

this only works for invertible maps - otherwise likelihood is not available.

- Implicit generative models.

Take a base measure, e.g.,  $z \sim N(0, I)$  and map with e.g. a neural net. This does not admit ML, and models need to be learnt in some other way.

[Plot these flow models]