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## STATISTICAL MODELS WITH VARIATIONAL METHODS

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## ABSTRACT

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Some introduction about how important Variational methods are nowadays and what this project is about.

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## Part I

### BASIC CONCEPTS

In this chapter we will introduce the underlying concepts of probability and graph theory that we will need.

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## PROBABILITY

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All our theory will be made under the assumption that there is a *referential set*  $\Omega$ , set of all possible outcomes of an experiment. Any subset of  $\Omega$  will be called an *event*.

**Definition 1.** Let  $\mathcal{P}(\Omega)$  be the power set of  $\Omega$ . Then,  $\mathcal{F} \subset \mathcal{P}(\Omega)$  is called a  $\sigma$ -algebra if it satisfies:

- $\Omega \in \mathcal{F}$ .
- $\mathcal{F}$  is closed under complementation.
- $\mathcal{F}$  is closed under countable unions.

From these properties it follows that  $\emptyset \in \mathcal{F}$  and that  $\mathcal{F}$  is closed under countable intersections.

The tuple  $(\Omega, \mathcal{F})$  is called a *measurable space*.

**Definition 2.** A *probability*  $P$  over  $(\Omega, \mathcal{F})$  is a mapping  $P : \mathcal{F} \rightarrow [0, 1]$  which satisfies

- $P(\alpha) \geq 0 \quad \forall \alpha \in \mathcal{F}$ .
- $P(\Omega) = 1$ .
- $P$  is countably additive, that is, if  $\{\alpha_i\}_{i \in \mathbb{N}} \subset \mathcal{F}$ , is a countable collection of pairwise disjoint sets, then

$$P\left(\bigcup_{i \in \mathbb{N}} \alpha_i\right) = \sum_{i \in \mathbb{N}} P(\alpha_i).$$

The first condition guarantees non negativity. The second one states that the *trivial event* has the maximal possible probability of 1. The third condition implies that given a set of pairwise disjoint events, the probability of either one of them occurring is equal to the sum of the probabilities of each one.

From these conditions it follows that

- $P(\emptyset) = 0$
- $P(\alpha \cup \beta) = P(\alpha) + P(\beta) - P(\alpha \cap \beta)$

The triple  $(\Omega, \mathcal{F}, P)$  is called a *probability space*.

**Definition 3.** Given two events  $\alpha, \beta \in \mathcal{F}$ , with  $P(\beta) \neq 0$ , the conditional probability of  $\alpha$  given  $\beta$  is defined as the quotient of the probability of the joint events and the probability of  $\beta$ :

$$P(\alpha | \beta) = \frac{P(\alpha \cap \beta)}{P(\beta)}$$

**Theorem 1. (Bayes' theorem).** Let  $\alpha, \beta$  be two events of an experiment, given that  $P(\beta) \neq 0$ . Then

$$P(\alpha | \beta) = \frac{P(\beta | \alpha)P(\alpha)}{P(\beta)}$$

*Example 1.* Consider a study where the relation of a disease  $d$  and an habit  $h$  is being investigated. Suppose that  $P(d) = 10^{-5}$ ,  $P(h) = 0.5$  and  $P(h | d) = 0.9$ . What is the probability that a person with habit  $h$  will have the disease  $d$ ?

$$P(d | h) = \frac{P(d \cap h)}{P(h)} = \frac{P(h | d)P(d)}{P(h)} = \frac{0.9 \times 10^{-5}}{0.5} = 1.8 \times 10^{-5}$$

If we set the probability of having habit  $h$  to a much lower value as  $P(h) = 0.001$ , then the above calculation gives approximately  $1/100$ . Intuitively, a smaller number of people have the habit and most of them have the disease. This means that the relation between having the disease and the habit is stronger now compared with the case where more people had the habit.

**Definition 4.** We say that two events  $\alpha, \beta \in \mathcal{F}$  are *independent* if knowing one of them does not give any extra information about the other. Mathematically,

$$P(\alpha \cap \beta) = P(\alpha)P(\beta) \qquad P(\alpha | \beta) = P(\alpha)$$

Let  $\gamma \in \mathcal{F}$ , we say that  $\alpha$  and  $\beta$  are *conditionally independent* on  $\gamma$ ,  $\alpha \perp\!\!\!\perp \beta | \gamma$  if and only if

$$P(\alpha \cup \beta | \gamma) = P(\alpha | \gamma)P(\beta | \gamma)$$

Otherwise, they are said to be *conditionally dependent* on  $\gamma$ ,  $\alpha \not\perp\!\!\!\perp \beta | \gamma$ .

Now we are going to introduce the concept of *random variable* and some properties as we have done with events.

**Definition 5.** A function  $f : \Omega_1 \rightarrow \Omega_2$  between two measurable spaces  $(\Omega_1, \mathcal{F}_1)$  and  $(\Omega_2, \mathcal{F}_2)$  is said to be *measurable* if  $f^{-1}(\alpha) \in \mathcal{F}_1$  for every  $\alpha \in \mathcal{F}_2$ .

**Definition 6.** A *random variable* is a measurable function  $X : \Omega \rightarrow E$  from a probability space  $(\Omega, \mathcal{F}, P)$  to a measurable space  $(E, \mathcal{F}')$  verifying  $X(\omega) \in \mathcal{F}' \forall \omega \in \Omega$ .

The probability of  $X$  taking a value on a measurable set  $S \in \mathcal{E}$  is written as

$$P_X(S) = P(X \in S) = P(\{a \in \Omega \mid X(a) \in S\}).$$

We could make a question like “How likely is that the value of  $X$  equals  $a$ ?”. This is the same as asking for the probability of the set  $\{\omega \in \Omega \mid X(\omega) = a\}$ .

We will set the following notation that is going to be used, that is: random variables will be denoted with an upper case letter like  $X$  and a set of variables with a bold symbol like  $\mathbf{X}$ . The meaning of  $P(\text{state})$  will be clear without a reference to the variable. Otherwise  $P(X = \text{state})$  will be used. Using a lower case letter like  $P(x)$  will denote the probability of the corresponding upper case variable  $X$  taking a specific value.

**Definition 7.** The *cumulative distribution function* of a real-valued random variable  $X$  is the function given by:

$$F_X(x) = P(X \leq x)$$

where the right-hand side represents the probability of the random variable taking value below or equal to  $x$ .

**Definition 8.** When the image of a random variable  $X$  is countable, the random variable it is called a *discrete random variable*, its *probability mass function*  $p$  gives the probability of it being equal to some value.

$$p(x) = P(X = x)$$

If the image is uncountable and real, then  $X$  is called a *continuous random variable* if there exists a non-negative Lebesgue-integrable  $f$ , called its *probability density function* such that

$$F_X(x) = P(X \leq x) = \int_{-\infty}^x f(u) du$$

A *mixed random variable* is a random variable who is neither discrete nor continuous, it can be realized as the sum of a discrete and continuous random variables. An example of a random variable of mixed type would be based on an experiment where a coin is flipped and a random positive number is chose only if the result of the coin toss is heads,  $-1$  otherwise.

From now on,  $P(x)$  will denote  $f_X(x)$  when  $X$  is a continuous random variable. We will define the *probability distribution*  $P_X$  of a random variable  $X$  over the probability space  $(\Omega, \mathcal{F}, P)$  as the pushforward measure of it, that is,  $P_X = PX^{-1}$ .

**Definition 9.** As we did for events, we can define the *conditional probability* over random variables, let  $X, Y$  be random variables,

$$P(x \mid y) = \frac{P(x, y)}{P(y)}$$

It is required that  $P(y) \neq 0$  for the conditional probability to be defined.

We can also enunciate the *Bayes' theorem*.

$$P(x, y) = \frac{P(y | x)P(x)}{P(y)}$$

**Definition 10.** The *marginal distribution* of a subset of random variables is the probability distribution of the variables contained in that subset.

Let  $X, Y$  be two random variables, it follows that

$$P(x) = \sum_y P(x, y) \quad f_X(x) = \int_y f_{X,Y}(x, y)$$

**Definition 11.** Let  $\mathbf{X} = \{X_1, X_2, \dots, X_n\}$  be a set of random variables, the *joint probability distribution* for  $\mathbf{X}$  is function that gives the probability of each random variable  $X_i$  falling in a particular range or discrete set of values for that variable. It is called a *multivariate distribution*.

When using only two random variables, then is called a *bivariate distribution*.

This distribution can be expressed in terms of a joint cumulative distribution function

$$F_{\mathbf{X}}(\mathbf{x}) = F_{X_1, \dots, X_n}(x_1, \dots, x_n) = P(X_1 \leq x_1, \dots, X_n \leq x_n)^1$$

or using a probability density function (all variables must be continuous) or a probability mass function (all variables must be discrete).

**Definition 12.** We say that two random variables  $X$  and  $Y$  are *independent* if knowing one of them doesn't give any extra information about the other. Mathematically,

$$P(x, y) = P(x)P(y)$$

From this it follows that if  $X$  and  $Y$  are independent, then  $P(x | y) = P(x)$ .

**Definition 13.** Let  $X, Y$  and  $Z$  be three random variables, then  $X$  and  $Y$  are *conditionally independent* given  $Z$  if and only if

$$P(x, y | z) = P(x | z)P(y | z)$$

in that case we will denote  $X \perp\!\!\!\perp Y | Z$ . If  $X$  and  $Y$  are not conditionally independent, they are *conditionally dependent*  $X \not\perp\!\!\!\perp Y | Z$

Both independence definitions can be made over sets of variables  $\mathbf{X}, \mathbf{Y}$  and  $\mathbf{Z}$  in a straight forward way.

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<sup>1</sup> Where  $\mathbf{x} = (x_1, \dots, x_n)$



**Definition 14.** We say that a set of  $n$  random variables  $\{X_1, \dots, X_n\}$  defined to assume values in  $I \subset \mathbb{R}$  are *independent and identically distributed (i.i.d)* if and only if they are independent

$$F_{X_1, \dots, X_n}(x_1, \dots, x_n) = F_{X_1}(x_1) \dots F_{X_n}(x_n) \quad \forall x_1, \dots, x_n \in I$$

and are identically distributed

$$F_{X_1}(x_1) = F_{X_k}(x_k) \quad \forall k \in \{2, \dots, n\} \text{ and } \forall x \in I$$

**Definition 15.** A *multivariate random variable* or *random vector* is a column vector  $\mathbf{X} = (X_1, \dots, X_n)^T$  whose components are random variables that can be defined over different probability spaces.

Note that we use the same symbol  $\mathbf{X}$  for random vectors and sets of variables, but the meaning will be clear within the context.

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DISTRIBUTIONS

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In this section we will summarize some concepts concerning probability distributions among with some of the most used ones.

From now on, let  $X$  be a random variable and  $P$  its probability distribution.

**Definition 16.** The *mode*  $X_*$  of the probability distribution  $P$  is the state of  $X$  where the distribution takes it's highest value

$$X_* = \arg \max_x P(x)$$

A distribution could have more than one mode, in this case we say it is *multi-modal*.

**Definition 17.** The notation  $\mathbb{E}[X]$  is used to denote the *average* or *expectation* of the values a real-valued variable takes respect to its distribution. If  $X$  is non-negative, it is defined as

$$\mathbb{E}[X] = \int_{\Omega} X(\omega) dP(\omega)^1$$

For a general variable  $X$  it is defined as  $\mathbb{E}[X] = \mathbb{E}[X^+] - \mathbb{E}[X^-]$ . Where

$$X^+(\omega) = \max(X(\omega), 0) \quad X^-(\omega) = \min(X(\omega), 0)$$

Suppose now that  $X$  is a real-valued random variable, in case it is also continuous

$$\mathbb{E}[X] = \int_{-\infty}^{+\infty} x f(x) dx$$

and if it is discrete, let  $x_i$  be the values  $X$  can take

$$\mathbb{E}[X] = \sum_{i=1}^{+\infty} x_i p(x_i)$$

Let  $g : \mathbb{R} \rightarrow \mathbb{R}$  be a measurable function, then  $g \circ X$  is another random variable and we can talk about  $\mathbb{E}[g(X)]$ , so in case  $X$  is continuous, we have that

$$\mathbb{E}[g(X)] = \int_{-\infty}^{+\infty} g(x) f(x) dx$$

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<sup>1</sup>  $P$  is a measure over  $\Omega$

**Definition 18.** We define the  $k^{th}$  moment of a distribution as the average of  $X^k$  over the distribution

$$\mu_k = \mathbb{E}[X^k]$$

For  $k = 1$  it is typically denoted as  $\mu$ .

**Definition 19.** The *variance* of a distribution is defined as

$$\text{Var}(X) = \sigma^2 = \mathbb{E}[(X - \mathbb{E}[X])^2] = \mathbb{E}[X]^2 - \mathbb{E}[X^2]$$

The square root of the variance  $\sigma$  is called the *standard deviation*.

When using a multivariate distribution  $\mathbf{X} = (X_1, \dots, X_n)^T$  we can talk about the *covariance matrix*  $\Sigma$  whose elements are

$$\begin{aligned} \Sigma_{ij} &= \mathbb{E}[(X_i - \mathbb{E}[X_i])(X_j - \mathbb{E}[X_j])] = \mathbb{E}[(X_i - \mu_i)(X_j - \mu_j)] \\ &= \mathbb{E}[X_i X_j] - \mathbb{E}[X_i]\mathbb{E}[X_j] \end{aligned}$$

The following result will be helpful later on

**Proposition 1.** Let  $\{X_1, \dots, X_n\}$  be a set of random variables and  $P$  their joint probability distribution. It follows that the expectation of a function  $f$  on a subset of the variables  $\mathcal{X}_0 \subset \mathcal{X} = \{X_1, \dots, X_n\}$ , verifies

$$\mathbb{E}[f(\mathcal{X}_0)]_{P(\mathcal{X})} = \mathbb{E}[f(\mathcal{X}_0)]_{P(\mathcal{X}_0)}$$

that is, we only need to know the marginal distribution of the subset in order to carry out the average.

We are going to discuss now some examples of probability distributions that are going to be used from now on.

## 2.1 DISCRETE DISTRIBUTIONS

### 2.1.1 Bernoulli distribution

The Bernoulli distribution describes a discrete binary variable  $X$  that takes the value 1 with probability  $p$  and the value 0 with probability  $1 - p$ .

$$p(x) = \begin{cases} p & \text{if } x = 1 \\ 1 - p & \text{if } x = 0 \end{cases}$$

### 2.1.2 Binomial distribution

The binomial distribution describes the number of successes in a sequence of independent Bernoulli Trials. A discrete binary random variable  $X$  follows a *binomial distribution* of parameters  $n \in \mathbb{N}$  and  $p \in [0, 1]$ , denoted as  $X \sim B(n, p)$  if and only if

$$P(x = k) = \binom{n}{k} p^k (1 - p)^{n-k}$$

## 2.2 CONTINUOUS DISTRIBUTIONS

### 2.2.1 Univariate Normal distribution

The *normal* or *Gaussian distribution* is a type of continuous probability distribution for real-valued random variables.

**Definition 20.** We say the real valued random variable  $X$  follows a *normal distribution* of parameters  $\mu, \sigma \in \mathbb{R}$ , denoted as  $X \sim N(\mu, \sigma)$  if and only if, its probability density function exists and is

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{1}{2}\left(\frac{x-\mu}{\sigma}\right)^2}$$

The parameter  $\mu$  is the mean or expectation of the distribution and  $\sigma$  is its standard deviation.

The simplest case of a normal distribution is known as *standard normal distribution*, denoted as  $Z$ . It is a special case where  $\mu = 0$  and  $\sigma = 1$ , then its density function is

$$f(x) = \frac{1}{\sqrt{2\pi}} e^{-\frac{1}{2}x^2}$$

One of the properties of the normal distribution is that if  $X \sim N(\mu, \sigma)$ ,  $a, b \in \mathbb{R}$  and  $f : \mathbb{R} \rightarrow \mathbb{R}$  be defined as  $f(x) = ax + b$ , then  $f(X) \sim N(\mu + b, a^2\sigma)$

### 2.2.2 Multivariate Normal Distribution

This distribution plays a fundamental role in this project so we will discuss its properties in more detail.

This distribution is an extension of the uni-variate one when having a multivariate random variable.

**Definition 21.** We say that a random vector  $\mathbf{X} = (X_1, \dots, X_p)$  follows a *multivariate normal distribution* of parameters  $\boldsymbol{\mu} \in \mathbb{R}^p$  and  $\boldsymbol{\Sigma} \in \mathbb{M}_n(\mathbb{R})$ , denoted as  $\mathbf{X} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma})$  if and only if its probability density function is

$$f(\mathbf{x}) = \frac{1}{\sqrt{\det(2\pi\boldsymbol{\Sigma})}} e^{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu})}$$

Where  $\boldsymbol{\mu}$  is the mean vector of the distribution, and  $\boldsymbol{\Sigma}$  the covariance matrix. The inverse matrix  $\boldsymbol{\sigma}^{-1}$  is called *precision*. It also satisfies that

$$\boldsymbol{\mu} = \mathbb{E}[\mathbf{X}] \quad \boldsymbol{\Sigma} = \mathbb{E}[(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^T]$$

As  $\boldsymbol{\Sigma}$  is a real symmetric matrix, it can be eigendecomposed

$$\boldsymbol{\Sigma} = \mathbf{E} \boldsymbol{\Delta} \mathbf{E}^T$$

where  $\mathbf{E}^T \mathbf{E} = \mathbf{I}$  and  $\boldsymbol{\Delta} = \text{diag}(\lambda_1, \dots, \lambda_n)$ .

Using the transformation

$$\mathbf{y} = \boldsymbol{\Delta}^{\frac{1}{2}} \mathbf{E}^T (\mathbf{x} - \boldsymbol{\mu})$$

we get that

$$(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma} (\mathbf{x} - \boldsymbol{\mu}) = \mathbf{y}^T \mathbf{y}$$

Using this, the multivariate Normal Distribution reduces to a product of  $n$  univariate standard normal distributions.

TODO: Add more properties as they are needed

### 2.2.3 Beta Distribution

Another continuous distribution that we are going to use is the *Beta distribution*.

**Definition 22.** We say that a continuous random variable  $X$  defined on the interval  $[0, 1]$  follows a *Beta distribution* of parameters  $\alpha, \beta > 0$ , denoted as  $X \sim \text{Beta}(\alpha, \beta)$  if and only if its density function is

$$f(x) = \frac{1}{B(\alpha, \beta)} x^{\alpha-1} (1-x)^{\beta-1}$$

where  $B(\alpha, \beta)$  is the *beta function* defined as

$$B(\alpha, \beta) = \int_0^1 x^{\alpha-1} (1-x)^{\beta-1} dx$$

The mean is given by  $\mathbb{E}[X] = \frac{\alpha}{\alpha+\beta}$

### 2.2.4 Dirichlet Distribution

The Dirichlet distribution is a family of continuous multivariate probability distributions parameterized by a vector  $\alpha$  of positive reals. It is a multivariate generalization of the Beta Distribution.

**Definition 23.** We say that a continuous random multivariate variable  $\mathbf{X}$  with order  $K \geq 2$ , follows a *Dirichlet Distribution* with parameters  $\alpha = (\alpha_1, \dots, \alpha_K)$ , if and only if its density function is defined as

$$f(\mathbf{x}) = \frac{1}{B(\alpha)} \prod_{k=1}^K x_k^{\alpha_k-1}$$

and it satisfies that

$$\sum_{k=1}^K x_k = 1 \text{ and } x_k > 0 \forall k = 1, \dots, K$$

Where the normalization constant is the beta function

$$B(\alpha) = \frac{\prod_k \Gamma(\alpha_k)}{\Gamma(\sum_k \alpha_k)}$$

## 2.3 KULLBACK-LEIBLER DIVERGENCE

**Definition 24.** Let  $P$  and  $Q$  be two probability distributions over the same set of random variables  $\mathbf{X}$ , the *Kullback-Leibler divergence*  $KL(Q|P)$  measures the ‘difference’ between both distributions as

$$KL(Q|P) = \mathbb{E}[\log Q(x) - \log P(x)]_Q$$

**Proposition 2.** *The Kullback-Leibler divergence is always non-negative.*

*Proof.* As the logarithm is bounded by  $x - 1$ , we can bound  $\log \frac{P(x)}{Q(x)}$

$$\log x \leq x - 1 \implies \frac{P(x)}{Q(x)} - 1 \geq \log \frac{P(x)}{Q(x)}$$

Since probabilities are non-negative, we can multiply by  $Q(x)$

$$P(x) - Q(x) \geq Q(x) \log P(x) - Q(x) \log Q(x)$$

Now we integrate (sum in case of discrete variables) both sides

$$0 \geq \mathbb{E}[\log P(x) - \log Q(x)]_Q \implies \mathbb{E}[\log Q(x) - \log P(x)]_Q \geq 0$$

□

As a result, the Kullback-Leibler divergence is 0 if and only if the two distributions are equal almost everywhere.

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GRAPH THEORY

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**Definition 25.** A graph  $G = (V, E)$  is a set of vertices or nodes  $V$  and edges  $E \subset V \times V$  between them. If  $V$  is a set of ordered pairs then the graph is called a *directed graph*, otherwise if  $V$  is a set of unordered pairs it is called an *undirected graph*.

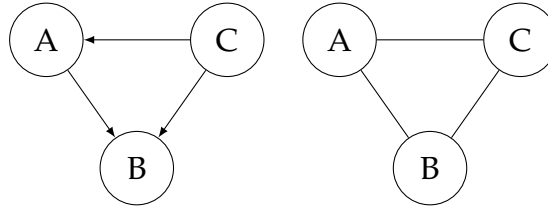


Figure 1: Example of directed and undirected graph, respectively.

**Definition 26.** In a directed graph  $G = (V, E)$ , a *directed path*  $A \rightarrow B$  is a sequence of vertices  $A = A_0, A_1, \dots, A_{n-1}, A_n = B$  where  $(A_i, A_{i+1}) \in E \forall i \in 0, \dots, n-1$ .

If  $G$  is a undirected graph,  $A \rightarrow B$  is an *undirected path* if  $\{A_i, A_{i+1}\} \in E \forall i \in 0, \dots, n-1$

**Definition 27.** Let  $A, B$  be two vertices of a directed graph  $G$ . If  $A \rightarrow B$  is a directed path and  $B \not\rightarrow A$  (meaning there isn't a directed path from  $B$  to  $A$ ), then  $A$  is called an *ancestor* of  $B$  and  $B$  is called a *descendant* of  $A$ .

For example, in the figure 1,  $C$  is an ancestor of  $B$ .

**Definition 28.** A *directed acyclic graph (DAG)* is a directed graph such that no directed path between any two nodes revisits a vertex.

As we can see in the figure 2,  $A \rightarrow B \rightarrow C \rightarrow A \rightarrow B$  is a path from  $A$  to  $B$  that revisits  $A$ .

Now we are going to define some relations between nodes in a DAG.

**Definition 29.** The *parents* of a node  $A$  is the set of nodes  $B$  such that there is a directed edge from  $B$  to  $A$ . The same applies for the *children* of a node.



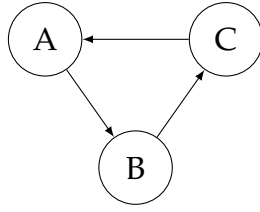


Figure 2: Example of graph which isn't a DAG.

The *Markov blanket* of a node is composed by the node itself, its children, its parents and the parents of its children.

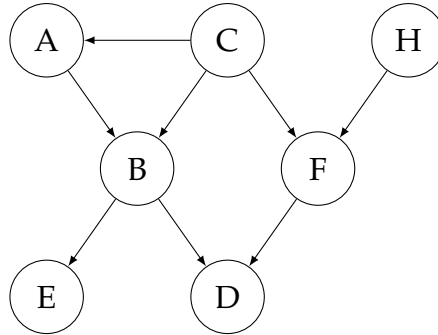


Figure 3: Directed acyclic graph

**Definition 30.** In a graph, the *neighbors* of a node are those directly connected to it.

We can use figure 3 to reflect on these definitions. The parents of  $B$  are  $pa(B) = \{A, C\}$  and its children are  $ch(B) = \{E, D\}$ . Taking this into account, its neighbors are  $ne(B) = \{A, C, E, D\}$  and its Markov blanket is  $\{A, B, C, D, E, F\}$ .

**Definition 31.** Let  $G$  be a DAG,  $U$  be a path between two vertex and  $A \in U$

- $A$  is called a *collider* if  $\forall B \in ne(A) \cap U, (B, A) \in E$ .
- $A$  is called a *fork* if  $\forall B \in ne(A) \cap U, (A, B) \in E$ .

Notice, a vertex can be a collider for a path but not for others.

For example in figure 3,  $D$  is a collider and  $C$  is a fork.

**Definition 32.** Let  $G$  be an undirected graph, a *clique* is a maximally connected subset of vertices. That is, all the members of the clique are connected to each others and there is no bigger clique that constains another.

Formally,  $S \subset V$  is a *clique* if and only if  $\forall A, B \in S, \{A, B\} \in E$  and  $\nexists C \in V \setminus S$  such that  $\forall A \in S, \{A, C\} \in E$ .

## Part II

### GRAPHICAL MODELS

A *graphical model* is a statistical model for which a graph expresses the conditional dependence structure between random variables.

Commonly, they provide a graph-based representation for encoding a multi-dimensional distribution representing a set of independences that hold in the specific distribution. The most commonly used are *Bayesian networks* and *Markov random fields*, which differ in the set of independences they can encode and the factorization of the distribution that they include.

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## BAYESIAN NETWORKS

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Consider we have  $N$  variables with the corresponding distribution  $P(x_1, \dots, x_N)$ . Let  $\mathcal{E}$  be a set of indexes such as evidence =  $\{X_e = x_e \mid e \in \mathcal{E}\}$ . Inference could be made by brute force:

$$P(X_i = x_i \mid \text{evidence}) = \frac{\int_{j \notin \mathcal{E}, j \neq i} P(\text{evidence}, x_j, X_i = x_i)}{\int_{j \notin \mathcal{E}} P(\text{evidence}, x_j)}$$

The notation when using discrete variables is analogous replacing integration with summations.

Lets suppose all these variables are binary, this calculation will require  $O(2^{N-\#\mathcal{E}})$  operations. Also, all entries of a table  $P(x_1, \dots, x_N)$  take  $O(2^N)$  space.

This is unpractical when taking into account millions of variables. The underlying idea of belief networks is to specify which variables are independent of others, factoring the joint probability distribution.

**Definition 33.** Let  $G = (V, E)$  be a graph where  $V = \{X_1, \dots, X_n\}$  is a set of random variables. We say that the joint probability  $P(x_1, \dots, x_n)$  *factorizes* according to  $G$  if and only if

$$P(x_1, \dots, x_N) = \prod_{i=1}^N P(x_i \mid pa(x_i))$$

**Definition 34.** A *belief network* or *Bayesian network* is a pair  $(G, P)$  where  $P$  factorizes over  $G$ . It is a probabilistic graphical model that represents conditional dependencies of a set of variables  $X_1, \dots, X_n$ .

Any probability distribution can be written as a Bayesian Network, even though it may end up been a fully-connected DAG. To set the specification of the Belief Network, we need to define all elements of the probability tables  $P(x_i \mid pa(x_i))$ . When the number of variables is large, this is still intractable so the tables are generally parameterized in a low dimensional manner.

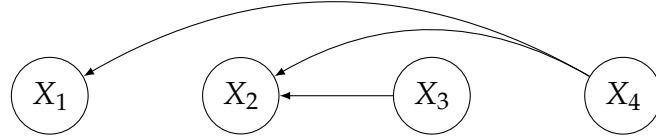


Figure 4: Bayesian Network factorizing  $P(x_1, x_2, x_3, x_4) = P(x_1|x_4)P(x_2|x_3, x_4)P(x_3)P(x_4)$

Bayesian Networks are good for encoding conditional independence over the variables, but aren't for encoding dependence. For example, with the following network  $P(x, y) = P(y|x)P(x)$  represented as  $x \rightarrow y$  in a DAG. It may appear to encode dependence between both variables but the conditional  $P(y|x)$  could happen to equal  $P(y)$ , giving  $P(x, y) = P(x)P(y)$ .

How could we check if two variables are conditionally independent given a Bayesian Network? For example in figure 3,  $X_1 \perp\!\!\!\perp X_2 \mid X_4$  as<sup>1</sup>:

$$\begin{aligned}
 P(x_2|x_4) &= \frac{1}{P(x_4)} \int_{x_1, x_3} P(x_1, x_2, x_3, x_4) = \frac{1}{P(x_4)} \int_{x_1, x_3} P(x_1|x_4)P(x_2|x_3, x_4)P(x_3)P(x_4) \\
 &= \int_{x_3} P(x_2|x_3, x_4)P(x_3) \\
 P(x_1, x_2|x_4) &= \frac{1}{P(x_4)} \int_{x_3} P(x_1, x_2, x_3, x_4) = \frac{1}{P(x_4)} \int_{x_3} P(x_1|x_4)P(x_2|x_3, x_4)P(x_3)P(x_4) \\
 &= P(x_1|x_4) \int_{x_3} P(x_2|x_3, x_4)P(x_3) = P(x_1|x_4)P(x_2|x_4)
 \end{aligned}$$

Now we are going to define two central concepts to determine conditional independence in any Bayesian Network, these are *d-connection* and *d-separation*.

**Definition 35.** Let  $G$  be a DAG where  $X, Y$  and  $Z$  are disjoint sets of vertices. We say that  $X$  and  $Y$  are *d-connected* by  $Z$  if and only if there exists an undirected path  $U$  from any vertex in  $X$  to any vertex in  $Y$  such that:

- For any collider  $C$ , itself or any it's descendants is in  $Z$
- No non-collider on  $U$  is on  $Z$

**Definition 36.** Let  $G$  be a DAG where  $X, Y$  and  $Z$  are disjoint sets of vertices.  $X$  and  $Y$  are *d-separated* by  $Z$  if and only if they are not d-connected by  $Z$  in  $G$

For example, in figure 5  $d$  d-separates  $a$  and  $c$  ( $b$  is a collider in the path that isn't in  $\{d\}$ ), and  $\{d, e\}$  d-connect them.

**Theorem 2 (Pearl & Dechter (2013)).** Let  $G$  be a DAG where  $X, Y$  and  $Z$  are disjoint sets of vertices. If  $X$  and  $Y$  are d-separated by  $Z$ , then they are independent conditional on  $Z$  in all probability distributions that  $G$  can represent.

<sup>1</sup> Continuous variable notation is used

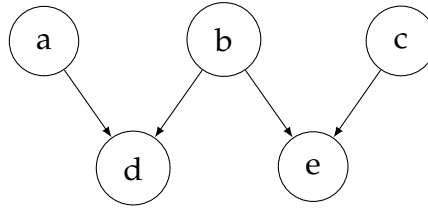


Figure 5: D-separation example

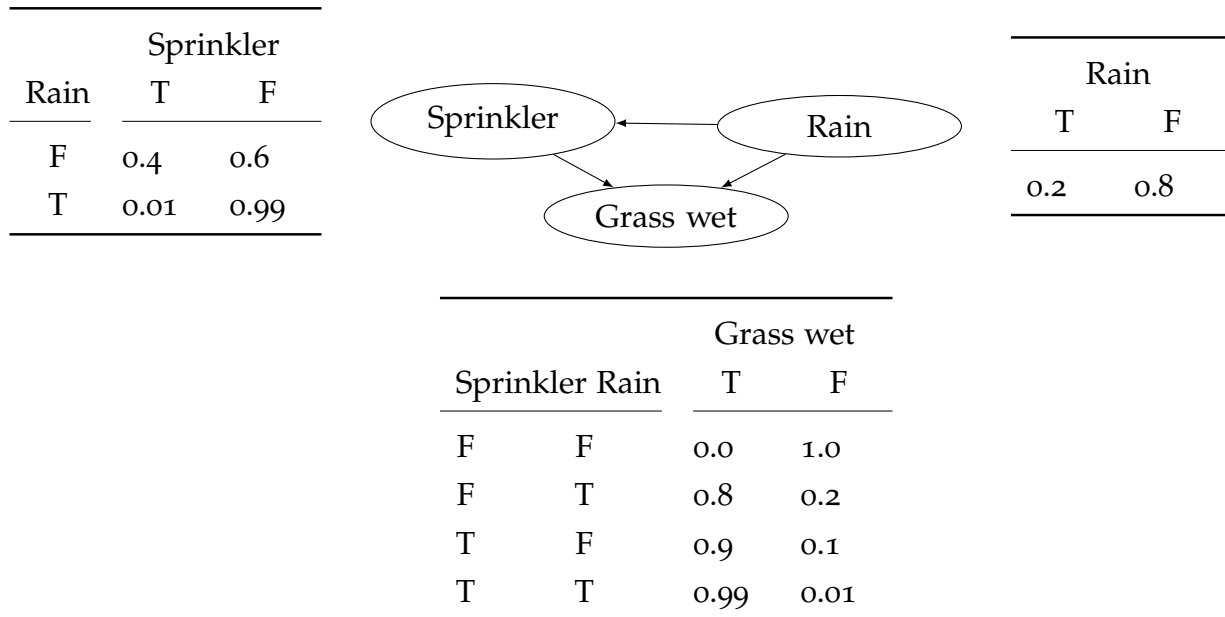
The Bayes Ball algorithm [Shachter \(2013\)](#) provides a linear time complexity algorithm that computes conditional independent using this theorem.

*Example 2.* In this example we are modeling three discrete random variables: Sprinkler ( $S$ ), Rain ( $R$ ) and Grass wet ( $G$ ).

The joint probability function is:

$$P(s, r, g) = P(s|r)P(g|s, r)P(r)$$

The following DAG illustrates the Bayesian Network among with the probability tables we are using.



This model can answer questions about the presence of a cause given the presence of an effect. For example, What is the probability that it has been raining given the grass is wet?

$$P(R = T|G = T) = \frac{P(G = T, R = T)}{P(G = T)} = \frac{\sum_s P(G = T, R = T, s)}{\sum_{r,s} P(G = T, r, s)}$$

Using the expression of the joint probability along with the tables we can compute every term. For example:

$$\begin{aligned} P(G = T, R = T, S = T) &= P(S = T | R = T) P(G = T | R = T, S = T) P(R = T) \\ &= 0.01 * 0.99 * 0.2 = 0.00198 \end{aligned}$$

In some situations our Belief Networks will contain a number of nodes that are essentially the same but repeated a number of times, for this, we are going to introduce the *plate notation*. Suppose we have the situation that figure 6 shows on the left. Then we can collapse all  $B_i$  variables in a box, indicating the number of variables inside it.

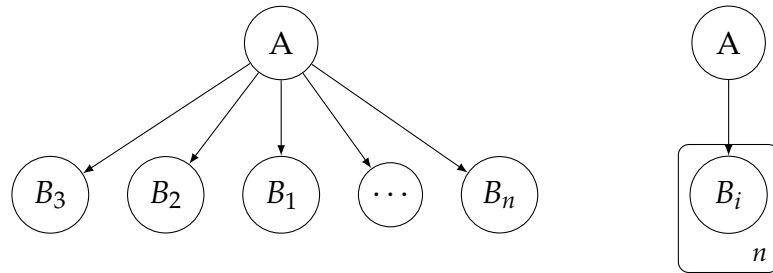


Figure 6: Plate notation example. Standard notation on the left and plate on the right

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MARKOV RANDOM FIELDS

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**Definition 37.** A *potential*  $\phi$  is a non-negative function. It is worth to mention that a probability distribution is a special case of a potential.

**Definition 38.** Let  $\mathbf{X}$  be a set of random variables,  $G$  an undirected graph,  $\mathbf{X}_c, c \in \{1, \dots, C\}$  be the maximal cliques of  $G$  and  $P$  a probability distribution over  $\mathbf{X}$ . The pair  $(G, P)$  is said to be a *Markov network* or *Markov random field* if, and only if

$$P(x_1, \dots, x_n) = \frac{1}{Z} \prod_{c=1}^C \phi_c(\mathbf{X}_c)$$

where  $Z$  is a constant that ensures normalization.

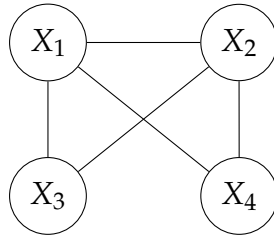


Figure 7: Markov Network  $P(x_1, x_2, x_3, x_4) = \phi(x_1, x_2, x_3)\phi(x_2, x_3, x_4)/Z$

In figure 7 we can see an example of the factorization, without giving any reference of the potentials.

Let  $(G, P)$  be a Markov network, then it satisfies the following properties known as Markov properties:

- Pairwise Markov property. Any two non-adjacent variables are conditionally independent given all other variables.
- Local Markov property. A variable is conditionally independent over all other variables given it's neighbors. That is,

$$P(x_i | x_{\setminus i}) = p(x_i | ne(x_i))^{1}$$

---

<sup>1</sup>  $X_{\setminus i} = \{X_j \mid j \neq i\} \subset \mathbf{X}$

- Global Markov property. Any two subsets of variables are conditionally independent given a separating subset (any path from one set to the other passes through this one).

*Remark 1.* A Markov network can also be defined as a pair  $(G, P)$  such as all Markov properties are satisfied. The clique factorization definition is a special case of these properties.

**Definition 39.** Let  $G$  be an undirected graph and  $P$  a probability distribution over a set of random variables  $\mathbf{X}$ . The pair  $(G, P)$  is called a Markov Random Field if and only if it follows the local Markov Property.



## Part III

NAME THIS PART

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## LEARNING AS INFERENCE

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In Machine Learning and related fields, the distributions are not fully specified and need to be learned from the data.

From now on,  $\mathcal{V}$  will denote the known data and  $\theta$  the set of parameters of the data distributions. The main task is to determine this set of parameters using the information given by the data.

**Definition 40.** *Priors* and *posteriors* typically refer to the parameter distribution before and after seeing the data, respectively. Using Bayes' rule

$$P(\theta \mid \mathcal{V}) = \frac{P(\mathcal{V} \mid \theta)P(\theta)}{P(\mathcal{V})}$$

The factor  $P(\mathcal{V} \mid \theta)$  is called the *likelihood*.

Let us see an example of our goal, in it we will try to learn the bias of a coin, given a set of tossing results.

*Example 3.* Let  $\{v_n\}_{n \in 0, \dots, N}$  be the results of tossing a coin  $N \in \mathbb{N}$  times, let 1 symbolize *heads* and 0 *tails*.

Our objective is to estimate the probability  $\theta$  that the coin will be head  $P(v_n = 1 \mid \theta)$ , for this we have the random variables  $v_1, \dots, v_n$  and  $\theta$ , and we require a model  $P(v_1, \dots, v_n, \theta)$ . We are considering the variables  $v_i$  to be independent to each others, we have a Belief Network depicted in figure 8

$$P(v_1, \dots, v_n, \theta) = P(\theta) \prod_{n=1}^N P(v_n \mid \theta)$$

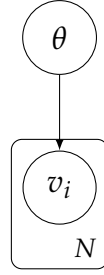


Figure 8: Belief network for coin tossing

We want to calculate

$$P(\theta \mid v_1, \dots, v_n) = \frac{P(v_1, \dots, v_n \mid \theta)P(\theta)}{P(v_1, \dots, v_n)}$$

to do so, we need to specify the prior  $P(\theta)$ , we are using a discrete model where

$$P(\theta = 0.2) = 0.1 \quad P(\theta = 0.5) = 0.7 \quad P(\theta = 0.8) = 0.2$$

This means that we have a 70% belief that the coin is fair, a 10% belief that is biased to tails and 20% that is biased to heads. Notice that  $P(v_n \mid \theta) = \theta$  if  $v_n = 1$  and  $P(v_n \mid \theta) = 1 - \theta$  if  $v_n = 0$ .

Let  $n_h$  be the number of heads in our observed data and  $n_t$  the number of tails

$$P(\theta \mid v_1, \dots, v_n) = \frac{P(\theta)}{P(\mathcal{V})} \theta^{n_h} (1 - \theta)^{n_t}$$

Suppose now that  $n_h = 2$  and  $n_t = 0.8$ , then

$$\begin{aligned} P(\theta = 0.2 \mid \mathcal{V}) &= \frac{1}{P(\mathcal{V})} \times 0.1 \times 0.2^2 \times 0.8^8 = \frac{1}{P(\mathcal{V})} \times 6.71 \times 10^{-4} \\ P(\theta = 0.5 \mid \mathcal{V}) &= \frac{1}{P(\mathcal{V})} \times 0.7 \times 0.5^2 \times 0.5^8 = \frac{1}{P(\mathcal{V})} \times 6.83 \times 10^{-4} \\ P(\theta = 0.8 \mid \mathcal{V}) &= \frac{1}{P(\mathcal{V})} \times 0.2 \times 0.2^2 \times 0.8^8 = \frac{1}{P(\mathcal{V})} \times 3.27 \times 10^{-7} \end{aligned}$$

Now, we can compute

$$\frac{1}{P(\mathcal{V})} = 6.71 \times 10^{-4} + 6.83 \times 10^{-4} + 3.27 \times 10^{-7} = 0.00135$$

So,

$$\begin{aligned} P(\theta = 0.2 \mid \mathcal{V}) &= 0.4979 \\ P(\theta = 0.5 \mid \mathcal{V}) &= 0.5059 \\ P(\theta = 0.8 \mid \mathcal{V}) &= 0.00024 \end{aligned}$$

These are the posterior parameter beliefs of our experiment. Given this, if we were to choose a single value for the posterior it would be  $\theta = 0.5$ . This result is intuitive since, we had a strong belief of the coin being fair and even though the number of tails was quite bigger than heads, it was not enough to make the difference. Obviously the posterior of the coin being biased to tails is now bigger than the prior.

Let us use an uniform prior distribution so that  $P(\theta) = k \implies \int_0^1 P(\theta) d\theta = k = 1$  due to normalization.

Using the previous calculations we have

$$P(\theta | \mathcal{V}) = \frac{1}{P(\mathcal{V})} \theta^{n_h} (1 - \theta)^{n_t}$$

where

$$P(\mathcal{V}) = \int_0^1 \theta^{n_h} (1 - \theta)^{n_t} d\theta$$

this implies that

$$P(\theta | \mathcal{V}) = \frac{\theta^{n_h} (1 - \theta)^{n_t}}{\int_0^1 u^{n_h} (1 - u)^{n_t} du} \equiv \text{Beta}(n_h + 1, n_t + 1)$$

**Definition 41.** If the posterior distribution is in the same probability distribution family as the prior distribution, they are then called *conjugate distributions*, and the prior is called a *conjugate prior* of the likelihood distribution.

Let's use a Beta distribution as the prior in the last example

$$P(\theta) = \frac{1}{B(\alpha, \beta)} \theta^{\alpha-1} (1 - \theta)^{\beta-1} \equiv \text{Beta}(\alpha, \beta)$$

then, repeating the same as before we get that

$$P(\theta, \mathcal{V}) = \frac{1}{B(\alpha + n_h, \beta + n_t)} \theta^{\alpha+n_h-1} (1 - \theta)^{\beta+n_t-1} \equiv \text{Beta}(\alpha + n_h, \beta + n_t)$$

So both the prior and posterior are Beta distributions, then the Beta distribution is called “conjugate” of the Binomial distribution.

## 6.1 UTILITY

The Bayesian posterior says nothing about how to summarize the beliefs it represents, in order to do this we need to specify the utility of each decision.

With this idea we define an utility function over the parameters

$$U(\theta, \theta_{true}) = \alpha \mathbb{I}[\theta = \theta_{true}] - \beta \mathbb{I}[\theta \neq \theta_{true}]$$

where  $\theta_{true}$  symbolizes the true value of the parameter, and  $\alpha, \beta \in \mathbb{R}$ .

Then the expected utility of a parameter  $\theta_0$  is calculated as

$$U(\theta = \theta_0) = \sum_{\theta_{true}} U(\theta = \theta_0, \theta_{true}) P(\theta = \theta_{true} \mid \mathcal{V})$$

Using the last example, we may define out utility function as

$$U(\theta, \theta_{true}) = 10 \mathbb{I}[\theta = \theta_{true}] - 20 \mathbb{I}[\theta \neq \theta_{true}]$$

so the expected utility of the decision that the parameter is  $\theta = 0.2$  is

$$\begin{aligned} U(\theta = 0.2) &= U(\theta = 0.2, \theta_{true} = 0.2) P(\theta_{true} = 0.2 \mid \mathcal{V}) \\ &\quad + U(\theta = 0.2, \theta_{true} = 0.5) P(\theta_{true} = 0.5 \mid \mathcal{V}) \\ &\quad + U(\theta = 0.2, \theta_{true} = 0.8) P(\theta_{true} = 0.8 \mid \mathcal{V}) \end{aligned}$$

## 6.2 MAXIMUM A POSTERIORI AND MAXIMUM LIKELIHOOD

The posterior reflects our beliefs about the full range of probabilities, but we may want to summarize all this information, even though, we may lose loads of it.

**Definition 42.** Maximum Likelihood is calculated as

$$\theta^{ML} = \arg \max_{\theta} p(\mathcal{V} \mid \theta)$$

it refers to the value of the parameter  $\theta$  for which the observed data better fits the model.

**Definition 43.** Maximum A Posteriori is calculated as

$$\theta^{MAP} = \arg \max_{\theta} p(\mathcal{V} \mid \theta) P(\theta)$$

The decision of taking the Maximum A Posteriori can be motivated using an utility that equals zero for all but the correct parameter

$$U(\theta, \theta_{true}) = \mathbb{I}[\theta = \theta_{true}]$$

using this, the expected utility of a parameter  $\theta = \theta_0$  is

$$U(\theta = \theta_0) = \sum_{\theta_{true}} \mathbb{I}[\theta_{true} = \theta_0] P(\theta = \theta_{true} \mid \mathcal{V}) = P(\theta_0 \mid \mathcal{V})$$

This means that the maximum utility decision is to take the value  $\theta_0$  with the highest posterior value.

It is worth mentioning that when using a flat prior  $\theta^{ML} = \theta^{MAP}$ .

Let  $\{x_1, \dots, x_m\}$  be a set of discrete variables, we can define the empirical distribution as a distribution of the variables whose mass probability function is

$$Q(x) = \frac{1}{m} \sum_{i=1}^m \mathbb{I}[x = x_i]$$

Now, we are going to show the relation between the Maximum Likelihood and the Kullback-Leibler divergence of the empirical distribution and our model. We may calculate this Kullback-Leibler divergence and study their functional independence.

$$KL(Q | P) = \mathbb{E}[\log(Q(x))]_Q - \mathbb{E}[\log(P(x))]_Q$$

Notice the term  $\mathbb{E}[\log(Q(x))]_Q$  is a constant and assume the data is i.i.d, so that

$$\mathbb{E}[\log(P(x))]_Q = \frac{1}{m} \sum_{i=1}^m \log P(x_i)$$

where the right side is the log likelihood under  $P(x)$ . As the logarithm is a strictly increasing function, maximizing the log likelihood equals to maximize the likelihood itself, and we can see here how it is equivalent to minimize the Kullback-Leibler divergence between the empirical distribution and our distribution.

In case  $P(x)$  is unconstrained, the optimal choice is  $P(x) = Q(x)$ , that is, the maximum likelihood distribution corresponds to the empirical distribution.

For a Belief Network  $P(x)$  presents the following constraint

$$P(x) = \prod_{i=1}^K P(x_i | pa(x_i))$$

We now want to minimize the Kullback-Leibler divergence between the empirical distribution  $Q(x)$  and  $P(x)$  in order to get the Maximum Likelihood.

$$\begin{aligned} KL(Q | P) &= -\mathbb{E}\left[\sum_{i=1}^K \log P(x_i | pa(x_i))\right]_Q + \mathbb{E}\left[\sum_{i=1}^K \log P(x_i | pa(x_i))\right]_P \\ &= -\sum_{i=1}^K \mathbb{E}\left[\log P(x_i | pa(x_i))\right]_Q + \sum_{i=1}^K \mathbb{E}\left[\log P(x_i | pa(x_i))\right]_P \end{aligned}$$

We can now use that  $\log P(x_i | pa(x_i))$  only depends on  $Q(x_i | pa(x_i))$  to rewrite it as

$$\begin{aligned}
KL(Q | P) &= \sum_{i=1}^K \mathbb{E} \left[ \log Q(x_i | pa(x_i)) \right]_{Q(x_i, pa(x_i))} - \mathbb{E} \left[ \log P(x_i | pa(x_i)) \right]_{Q(x_i, pa(x_i))} \\
&= \sum_{i=1}^K \mathbb{E} \left[ KL \left( Q(x_i | pa(x_i)) | P(x_i | pa(x_i)) \right) \right]_{Q(x_i, pa(x_i))}
\end{aligned}$$

The minimal setting is then

$$P(x_i | pa(x_i)) = Q(x_i | pa(x_i))$$

in terms of the initial data it is to set  $P(x_i | pa(x_i))$  to the number of times the state appears in it.

### 6.3 BAYESIAN BELIEF NETWORK TRAINING

A Bayesian approach where we set a distribution over the parameters is an alternative to Maximum Likelihood training of a Bayesian Network.

To illustrate this, we are using the following scenario, consider a disease  $D$  and two habits  $A$  and  $B$ , following the Bayesian Network

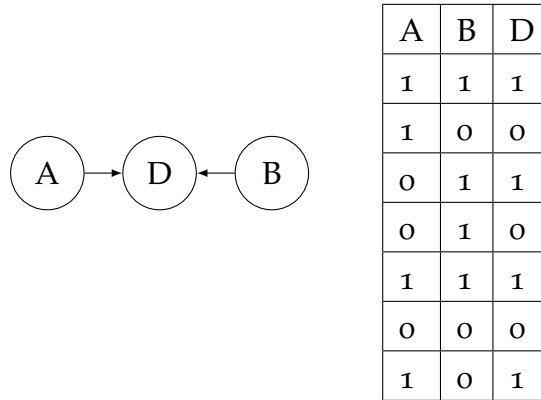


Figure 9: Model for the relationship between  $D$ ,  $A$  and  $B$ , and observations

$$P(a, b, d) = P(d|a, b)P(a)P(b)$$

Consider a set of observations  $\mathcal{V} = \{(a_n, b_n, d_n), n = 1, \dots, N\}$

We need a notation for the parameters, as all the variables are binary we are going to use

$$\begin{aligned}
P(A = 1 | \theta_a) &= \theta_a, & P(B = 1 | \theta_b) &= \theta_b, & P(D = 1 | A = 0, B = 1, \theta_d) &= \theta_1 \\
\theta_d &= (\theta_0, \theta_1, \theta_2, \theta_3)
\end{aligned}$$

Using the binary number  $A$  and  $B$  create (01) and its correspondent decimal form in the sub-index of  $\theta$ .

We need to specify a prior and since dealing with multi-dimensional continuous distributions is computationally problematic it is normal to use uni-variate distributions.

A convenient assumption is that the prior factorizes, this is usually called *global parameter independence*. We assume then

$$P(\theta_a, \theta_b, \theta_d) = P(\theta_a)P(\theta_b)P(\theta_d)$$

Assuming our data is i.i.d, we have

$$P(\theta_a, \theta_b, \theta_d, \mathcal{V}) = P(\theta_a)P(\theta_b)P(\theta_d) \prod_n P(a_n | \theta_a)P(b_n | \theta_b)P(d_n | a_n, b_n, \theta_d)$$

Learning then corresponds to inference

$$\begin{aligned} P(\theta_a, \theta_b, \theta_d | \mathcal{V}) &= \frac{P(\mathcal{V} | \theta_a, \theta_b, \theta_d)P(\theta_a, \theta_b, \theta_d)}{P(\mathcal{V})} = \frac{P(\mathcal{V} | \theta_a, \theta_b, \theta_d)P(\theta_a)P(\theta_b)P(\theta_d)}{P(\mathcal{V})} \\ &= \frac{1}{P(\mathcal{V})}P(\theta_a) \prod_n P(a_n | \theta_a)P(\theta_b) \prod_n P(b_n | \theta_b)P(\theta_d) \prod_n P(d_n | a_n, b_n, \theta_d) \\ &= P(\theta_a | \mathcal{V}_a)P(\theta_b | \mathcal{V}_b)P(\theta_d | \mathcal{V}) \end{aligned}$$

Where  $\mathcal{V}_i$  is the subset of the data restricted to the variable  $i$ . If we further assume that  $P(\theta_d)$  factorizes as  $P(\theta_d) = P(\theta_0)P(\theta_1)P(\theta_2)P(\theta_3)$ , this is called *local parameter independence*, then it follows that

$$P(\theta_d | \mathcal{V}) = P(\theta_0 | \mathcal{V})P(\theta_1 | \mathcal{V})P(\theta_2 | \mathcal{V})P(\theta_3 | \mathcal{V})$$

The simplest cases to continue are  $P(a | \theta_a)$  and  $P(b | \theta_b)$  since they require only a univariate prior distribution  $P(\theta_a)$  or  $P(\theta_b)$ . Lets illustrate it using  $P(\theta_a)$ :

The posterior is

$$P(\theta_a | \mathcal{V}_a) = \frac{1}{P(\mathcal{V}_a)}P(\theta_a)\theta_a^{\#(a=1)}(1 - \theta_a)^{\#(a=0)}$$

The most convenient choice for the prior is a Beta distribution as conjugacy will hold.

$$\begin{aligned} P(\theta_a) &= B(\theta_a | \alpha_a, \beta_a) = \frac{1}{B(\alpha_a, \beta_a)}\theta_a^{\alpha_a-1}(1 - \theta_a)^{\beta_a-1} \\ P(\theta_a | \mathcal{V}_a) &= B(\theta_a | \alpha_a + \#(a = 1), \beta_a + \#(a = 0)) \end{aligned}$$



The marginal is then

$$\begin{aligned}
 P(a = 1 \mid \mathcal{V}_a) &= \frac{P(a = 1, \mathcal{V}_a)}{P(\mathcal{V}_a)} = \int_{\theta_a} \frac{P(a = 1, \mathcal{V}_a, \theta_a)}{P(\mathcal{V}_a)} = \int_{\theta_a} \frac{P(a = 1 \mid \mathcal{V}_a, \theta_a) P(\mathcal{V}_a, \theta_a)}{P(\mathcal{V}_a)} \\
 &= \int_{\theta_a} \frac{P(a = 1 \mid \mathcal{V}_a, \theta_a) P(\theta_a \mid \mathcal{V}_a) P(\mathcal{V}_a)}{P(\mathcal{V}_a)} = \int_{\theta_a} P(\theta_a \mid \mathcal{V}_a) \theta_a = \mathbb{E}[\theta_a \mid \mathcal{V}_a] \\
 &= \frac{\alpha_a + \#(a = 1)}{\alpha_a + \#(a = 1) + \beta_a + \#(a = 0)}
 \end{aligned}$$

For  $P(d \mid a, b)$  the situation is more complex, the most convenient way is to specify a Beta prior for each one of the four components of  $\theta_d$ . Lets focus on  $P(d = 1 \mid a = 1, b = 0)$ , notice the parameters  $\alpha$  and  $\beta$  we used before now depend on  $a$  and  $b$ , for this reason we are using  $\alpha_d(a, b)$  and  $\beta_d(a, b)$  as prior parameters, these are called *hyperparameters*.

$$P(\theta_2) = B(\theta_2 \mid \alpha_d(1, 0) + \#(d = 1, a = 1, b = 0), \beta_d(1, 0) + \#(d = 0, a = 1, b = 0))$$

As before we got that

$$P(d = 1 \mid a = 1, b = 0, \mathcal{V}) = \frac{\alpha_d(1, 0) + \#(d = 1, a = 1, b = 0)}{\alpha_d(1, 0) + \beta_d(1, 0) + \#(a = 1, b = 0)}$$

In case we had no preference, we could set all hyperparameters to the same value, and, a complete ignorance prior would correspond to set them to 1.

Let now consider two limit possibilities, the one where we have no data at all, and the one where we have infinite data.

In case we have no data, the marginal probability corresponds to the prior which in the last case is

$$P(d = 1 \mid a = 1, b = 0, \mathcal{V}) = \frac{\alpha_d(1, 0)}{\alpha_d(1, 0) + \beta_d(1, 0)}$$

Note that equal hyperparameters would give a result of 0.5.

When infinite data is available, the marginal is generally dominated by it, this corresponds to the Maximum Likelihood solution.

$$P(d = 1 \mid a = 1, b = 0, \mathcal{V}) = \frac{\#(d = 1, a = 1, b = 0)}{\#(a = 1, b = 0)}$$

This happens unless the prior has a pathologically strong effect.

Consider the data given in the table in figure 9, and equal parameters and hyperparameters 1. Then we can compute the differences between this and using the Maximum Likelihood technique.

$$P(a = 1 \mid \mathcal{V}) = \frac{1 + \#(a = 1)}{2 + N} = \frac{5}{9} \approx 0.556$$

By comparison, the Maximum Likelihood result is  $4/7 = 0.571$ , the Bayesian result is more prudent than this one, which fits in with our belief that any setting is equally probable i.e 0.5.

The natural generalization to more than two states is using a Dirichlet distribution as prior, assuming i.i.d data and local and global prior independence.

Consider a variable  $X$  with  $Dom(X) = \mathcal{I}$ , then

$$P(x \mid \theta) = \prod_{i \in \mathcal{I}} \theta_i^{\mathbb{I}[x=i]} \text{ with } \sum_{i \in \mathcal{I}} \theta_i = 1$$

So that the posterior (considering  $N$  observations of the variable) is

$$P(\theta) \prod_{n=1}^N \prod_{i \in \mathcal{I}} \theta_i^{\mathbb{I}[x_n=i]} = P(\theta) \prod_{i \in \mathcal{I}} \theta_i^{\sum_n \mathbb{I}[x_n=i]}$$

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