

# CS 2750 Machine Learning

## Lecture 12b

# Bayesian belief networks

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# Density estimation

**Data:**  $D = \{D_1, D_2, \dots, D_n\}$   
 $D_i = \mathbf{x}_i$  a vector of attribute values

## Attributes:

- modeled by random variables  $\mathbf{X} = \{X_1, X_2, \dots, X_d\}$  with:
  - **Continuous values**
  - **Discrete values**

E.g. *blood pressure* with numerical values  
or *chest pain* with discrete values

[no-pain, mild, moderate, strong]

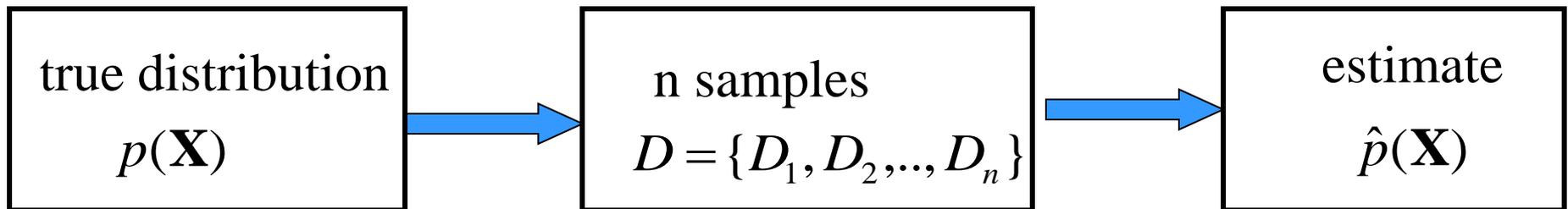
## Underlying true probability distribution:

$$p(\mathbf{X})$$

# Density estimation

**Data:**  $D = \{D_1, D_2, \dots, D_n\}$   
 $D_i = \mathbf{x}_i$  a vector of attribute values

**Objective:** try to estimate the underlying true probability distribution over variables  $\mathbf{X}$ ,  $p(\mathbf{X})$ , using examples in  $D$



**Standard (iid) assumptions: Samples**

- are **independent** of each other
- come from the same **(identical) distribution** (fixed  $p(\mathbf{X})$ )

# Learning via parameter estimation

In this lecture we consider **parametric density estimation**

## Basic settings:

- A set of random variables  $\mathbf{X} = \{X_1, X_2, \dots, X_d\}$
- **A model of the distribution** over variables in  $\mathbf{X}$

with parameters  $\Theta$  :

$$\hat{p}(\mathbf{X} | \Theta)$$

- **Data**  $D = \{D_1, D_2, \dots, D_n\}$

**Objective:** find the parameters  $\Theta$  that explain best the observed data

# Parameter estimation

- **Maximum likelihood (ML)**

maximize  $p(D | \Theta, \xi)$

- yields: one set of parameters  $\Theta_{ML}$
- the target distribution is approximated as:

$$\hat{p}(\mathbf{X}) = p(\mathbf{X} | \Theta_{ML})$$

- **Bayesian parameter estimation**

- uses the posterior distribution over possible parameters

$$p(\Theta | D, \xi) = \frac{p(D | \Theta, \xi) p(\Theta | \xi)}{p(D | \xi)}$$

- Yields: all possible settings of  $\Theta$  (and their “weights”)
- The target distribution is approximated as:

$$\hat{p}(\mathbf{X}) = p(\mathbf{X} | D) = \int_{\Theta} p(\mathbf{X} | \Theta) p(\Theta | D, \xi) d\Theta$$

# Parameter estimation

## Other possible criteria:

- **Maximum a posteriori probability (MAP)**

maximize  $p(\Theta | D, \xi)$  (mode of the posterior)

– Yields: one set of parameters  $\Theta_{MAP}$

– Approximation:

$$\hat{p}(\mathbf{X}) = p(\mathbf{X} | \Theta_{MAP})$$

- **Expected value of the parameter**

$\hat{\Theta} = E(\Theta)$  (mean of the posterior)

– Expectation taken with regard to posterior  $p(\Theta | D, \xi)$

– Yields: one set of parameters

– Approximation:

$$\hat{p}(\mathbf{X}) = p(\mathbf{X} | \hat{\Theta})$$

# Density estimation

- So far we have covered density estimation for “simple” distribution models:
  - Bernoulli
  - Binomial
  - Multinomial
  - Gaussian
  - Poisson

## But what if:

- The dimension of  $\mathbf{X} = \{X_1, X_2, \dots, X_d\}$  is large
  - Example: patient data
- Compact parametric distributions do not seem to fit the data
  - E.g.: multivariate Gaussian may not fit
- We have only a “small” number of examples to do accurate parameter estimates

# How to learn complex distributions

How to learn complex multivariate distributions  $\hat{p}(\mathbf{X})$  with large number of variables?

**One solution:**

- **Decompose the distribution using conditional independence relations**
- **Decompose the parameter estimation problem to a set of smaller parameter estimation tasks**

Decomposition of distributions under conditional independence assumption is the main idea behind **Bayesian belief networks**

# Example

## Problem description:

- **Disease:** pneumonia
- **Patient symptoms (findings, lab tests):**
  - Fever, Cough, Paleness, WBC (white blood cells) count, Chest pain, etc.

## Representation of a patient case:

- Symptoms and disease are represented as random variables

## Our objectives:

- **Describe a multivariate distribution representing the relations between symptoms and disease**
- **Design of inference and learning procedures for the multivariate model**

# Modeling uncertainty with probabilities

- **Full joint distribution:**

- Assume  $\mathbf{X} = \{X_1, X_2, \dots, X_d\}$  are all random variables that define the domain
- Full joint:  $P(\mathbf{X})$  or  $P(X_1, X_2, \dots, X_d)$

**Full joint it is sufficient** to do any type of probabilistic inference:

- Computation of joint probabilities for sets of variables

$$P(X_1, X_2, X_3) \quad P(X_1, X_{10})$$

- Computation of conditional probabilities

$$P(X_1 \mid X_2 = \text{True}, X_3 = \text{False})$$

# Marginalization

## Joint probability distribution (for a set variables)

- Defines probabilities for all possible assignments to values of variables in the set

$P(\text{pneumonia}, \text{WBCcount})$   $2 \times 3$  table

		<i>WBCcount</i>			$P(\text{Pneumonia})$
		<i>high</i>	<i>normal</i>	<i>low</i>	
<i>Pneumonia</i>	<i>True</i>	0.0008	0.0001	0.0001	0.001
	<i>False</i>	0.0042	0.9929	0.0019	0.999
		0.005	0.993	0.002	

$P(\text{WBCcount})$

**Marginalization** (summing of rows, or columns)

- summing out variables

# Variable independence

- The joint distribution over a subset of variables can be always computed from the joint distribution through marginalization
- Not the other way around !!!
  - **Only exception:** when variables are independent

$$P(A, B) = P(A)P(B)$$

$P(\text{pneumonia}, \text{WBCcount})$		$\text{WBCcount}$			$P(\text{Pneumonia})$
		<i>high</i>	<i>normal</i>	<i>low</i>	
<i>Pneumonia</i>	<i>True</i>	0.0008	0.0001	0.0001	0.001
	<i>False</i>	0.0042	0.9929	0.0019	0.999
		0.005	0.993	0.002	

$P(\text{WBCcount})$   $\longrightarrow$

# Conditional probability

## Conditional probability :

- Probability of A given B

$$P(A | B) = \frac{P(A, B)}{P(B)}$$

- Conditional probability is defined in terms of joint probabilities
- Joint probabilities can be expressed in terms of conditional probabilities

$$P(A, B) = P(A | B)P(B) \quad \text{(product rule)}$$

$$P(X_1, X_2, \dots, X_n) = \prod_{i=1}^n P(X_i | X_1, \dots, X_{i-1}) \quad \text{(chain rule)}$$

- Conditional probability – is useful for **various probabilistic inferences**

$$P(\text{Pneumonia} = \text{True} | \text{Fever} = \text{True}, \text{WBCcount} = \text{high}, \text{Cough} = \text{True})$$

# Inference

Any query can be computed from the full joint distribution !!!

- **Joint over a subset of variables** is obtained through marginalization

$$P(A = a, C = c) = \sum_i \sum_j P(A = a, B = b_i, C = c, D = d_j)$$

- **Conditional probability over a set of variables**, given other variables' values is obtained through marginalization and definition of conditionals

$$\begin{aligned} P(D = d \mid A = a, C = c) &= \frac{P(A = a, C = c, D = d)}{P(A = a, C = c)} \\ &= \frac{\sum_i P(A = a, B = b_i, C = c, D = d)}{\sum_i \sum_j P(A = a, B = b_i, C = c, D = d_j)} \end{aligned}$$

# Inference

- Any joint probability can be expressed as a product of conditionals via the **chain rule**.

$$\begin{aligned}P(X_1, X_2, \dots, X_n) &= P(X_n | X_1, \dots, X_{n-1})P(X_1, \dots, X_{n-1}) \\ &= P(X_n | X_1, \dots, X_{n-1})P(X_{n-1} | X_1, \dots, X_{n-2})P(X_1, \dots, X_{n-2}) \\ &= \prod_{i=1}^n P(X_i | X_1, \dots, X_{i-1})\end{aligned}$$

- It is often easier to define the distribution in terms of conditional probabilities:
  - E.g.  $\mathbf{P}(\textit{Fever} | \textit{Pneumonia} = T)$   
 $\mathbf{P}(\textit{Fever} | \textit{Pneumonia} = F)$

# Modeling uncertainty with probabilities

- **Full joint distribution:** joint distribution over all random variables defining the domain
  - it is sufficient to represent the complete domain and to do any type of probabilistic inferences

## Problems:

- **Space complexity.** To store full joint distribution requires to remember  $O(d^n)$  numbers.
  - $n$  – number of random variables,  $d$  – number of values
- **Inference complexity.** To compute some queries requires  $O(d^n)$  steps.
- **Acquisition problem.** Who is going to define all of the probability entries?

# Pneumonia example. Complexities.

- **Space complexity.**

- Pneumonia (2 values: T,F), Fever (2: T,F), Cough (2: T,F), WBCcount (3: high, normal, low), paleness (2: T,F)
- Number of assignments:  $2*2*2*3*2=48$
- We need to define at least 47 probabilities.

- **Time complexity.**

- Assume we need to compute the probability of Pneumonia=T from the full joint

$$\begin{aligned} P(\text{Pneumonia} = T) &= \\ &= \sum_{i \in T, F} \sum_{j \in T, F} \sum_{k=h, n, l} \sum_{u \in T, F} P(\text{Fever} = i, \text{Cough} = j, \text{WBCcount} = k, \text{Pale} = u) \end{aligned}$$

- Sum over  $2*2*3*2=24$  combinations

# Bayesian belief networks (BBNs)

## Bayesian belief networks.

- Represent the full joint distribution over the variables more compactly with a **smaller number of parameters**.
- Take advantage of **conditional and marginal independences** among random variables

- **A and B are independent**

$$P(A, B) = P(A)P(B)$$

- **A and B are conditionally independent given C**

$$P(A, B | C) = P(A | C)P(B | C)$$

$$P(A | C, B) = P(A | C)$$