# Inference and Representation

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### Lecture 1, September 8, 2015

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Inference and Representation

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One of the **most exciting advances** in machine learning (AI, signal processing, coding, control, ...) in the last decades

How can we gain global insight based on local observations?

- **Represent** the world as a collection of random variables  $X_1, \ldots, X_n$  with joint distribution  $p(X_1, \ldots, X_n)$
- Learn the distribution from data
- Perform "inference" (compute conditional distributions p(X<sub>i</sub> | X<sub>1</sub> = x<sub>1</sub>,..., X<sub>m</sub> = x<sub>m</sub>))

- As humans, we are continuously making predictions under uncertainty
- Classical AI and ML research ignored this phenomena
- Many of the most recent advances in technology are possible because of this new, *probabilistic*, approach



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## Applications: Speech recognition



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## input: two images

# output: disparity





- **Represent** the world as a collection of random variables  $X_1, \ldots, X_n$  with joint distribution  $p(X_1, \ldots, X_n)$ 
  - How does one compactly describe this joint distribution?
  - Directed graphical models (Bayesian networks)
  - Undirected graphical models (Markov random fields, factor graphs)
- Learn the distribution from data
  - Maximum likelihood estimation. Other estimation methods?
  - How much data do we need?
  - How much computation does it take?
- Perform "inference" (compute conditional distributions p(X<sub>i</sub> | X<sub>1</sub> = x<sub>1</sub>,..., X<sub>m</sub> = x<sub>m</sub>))

- We will study Representation, Inference & Learning
- First in the simplest case
  - Only discrete variables
  - Fully observed models
  - Exact inference & learning
- Then generalize
  - Continuous variables
  - Partially observed data during learning (hidden variables)
  - Approximate inference & learning
- Learn about algorithms, theory & applications

### • Class webpage:

- http://cs.nyu.edu/~dsontag/courses/inference15/
- Sign up for Piazza!
- **Book:** *Machine Learning: a Probabilistic Perspective* by Kevin Murphy, MIT Press (2012)
  - Required readings for each lecture posted to course website.
  - A good optional reference is *Probabilistic Graphical Models: Principles* and *Techniques* by Daphne Koller and Nir Friedman, MIT Press (2009)
- Office hours: Thurs 3:30-4:30pm. 715 Broadway, 12th floor, #1204 (except for 9/22, 11/3, 11/10, 11/24, 12/8: Tues 10:30-11:30am)
- Lab: Wednesdays, 7:10-8:00pm in WWH 102 (same as class)
  - Instructor: Rachel Hodos (hodos@cims.nyu.edu)
  - Required attendance; no exceptions.
- Grader: Prasoon Goyal (pg1338@nyu.edu)

#### • Prerequisite:

- DS-GA-1003/CSCI-GA.2567 (Machine Learning and Computational Statistics)
- **Grading:** not finalized problem sets (55%) + in class midterm exam (20%) + in class final exam (20%) + participation (5%)
  - I would love to see an active Piazza with students asking & responding to each other's questions. Will contribute to your participation grade.
  - 6 assignments (every 1–2 weeks). Both theory and programming.
  - First homework out tomorrow, due Friday Sept. 18 at 5pm (via email)
  - Important: See collaboration policy on class webpage
- Solutions to the theoretical questions must be rigorous.
- For the programming assignments, I recommend Python.

- Variable for each **symptom** (e.g. "fever", "cough", "fast breathing", "shaking", "nausea", "vomiting")
- Variable for each **disease** (e.g. "pneumonia", "flu", "common cold", "bronchitis", "tuberculosis")
- Diagnosis is performed by **inference** in the model:

 $p(\text{pneumonia} = 1 \mid \text{cough} = 1, \text{fever} = 1, \text{vomiting} = 0)$ 

• One famous model, Quick Medical Reference (QMR-DT), has 600 diseases and 4000 findings

- Naively, could represent multivariate distributions with table of probabilities for each outcome (assignment)
- How many outcomes are there in QMR-DT? 2<sup>4600</sup>
- Estimation of joint distribution would require a huge amount of data
- Inference of conditional probabilities, e.g.

 $p(\text{pneumonia} = 1 \mid \text{cough} = 1, \text{fever} = 1, \text{vomiting} = 0)$ 

would require summing over exponentially many variables' values

• Moreover, defeats the purpose of probabilistic modeling, which is to make predictions with *previously unseen observations* 

## Structure through independence

• If  $X_1, \ldots, X_n$  are independent, then

$$p(x_1,\ldots,x_n)=p(x_1)p(x_2)\cdots p(x_n)$$

- $2^n$  entries can be described by just *n* numbers (if  $|Val(X_i)| = 2$ )!
- However, this is not a very useful model observing a variable X<sub>i</sub> cannot influence our predictions of X<sub>j</sub>
- If  $X_1, \ldots, X_n$  are conditionally independent given Y, denoted as  $X_i \perp \mathbf{X}_{-i} \mid Y$ , then

$$p(y, x_1, ..., x_n) = p(y)p(x_1 | y) \prod_{i=2}^n p(x_i | x_1, ..., x_{i-1}, y)$$
  
=  $p(y)p(x_1 | y) \prod_{i=2}^n p(x_i | y).$ 

• This is a simple, yet *powerful*, model

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## Example: naive Bayes for classification

• Classify e-mails as spam (Y = 1) or not spam (Y = 0)

- Let 1 : *n* index the words in our vocabulary (e.g., English)
- $X_i = 1$  if word *i* appears in an e-mail, and 0 otherwise
- E-mails are drawn according to some distribution  $p(Y, X_1, \ldots, X_n)$
- Suppose that the words are conditionally independent given Y. Then,

$$p(y, x_1, \ldots x_n) = p(y) \prod_{i=1}^n p(x_i \mid y)$$

Estimate the model with maximum likelihood. Predict with:

$$p(Y = 1 \mid x_1, \dots, x_n) = \frac{p(Y = 1) \prod_{i=1}^n p(x_i \mid Y = 1)}{\sum_{y \in \{0,1\}} p(Y = y) \prod_{i=1}^n p(x_i \mid Y = y)}$$

- Are the independence assumptions made here reasonable?
- Philosophy: Nearly all probabilistic models are "wrong", but many are nonetheless useful

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- A Bayesian network is specified by a directed *acyclic* graph G = (V, E) with:
  - **①** One node  $i \in V$  for each random variable  $X_i$
  - One conditional probability distribution (CPD) per node, p(x<sub>i</sub> | x<sub>Pa(i)</sub>), specifying the variable's probability conditioned on its parents' values
- Corresponds 1-1 with a particular factorization of the joint distribution:

$$p(x_1,\ldots,x_n) = \prod_{i\in V} p(x_i \mid \mathbf{x}_{\operatorname{Pa}(i)})$$

- Powerful framework for designing *algorithms* to perform probability computations
- Enables use of prior knowledge to specify (part of) model structure

## Example

• Consider the following Bayesian network:



• What is its joint distribution?

$$p(x_1,...x_n) = \prod_{i \in V} p(x_i \mid \mathbf{x}_{\operatorname{Pa}(i)})$$
  
$$p(d, i, g, s, l) = p(d)p(i)p(g \mid i, d)p(s \mid i)p(l \mid g)$$

$$p(x_1,\ldots x_n) = \prod_{i\in V} p(x_i \mid \mathbf{x}_{\operatorname{Pa}(i)})$$

Will my car start this morning?



Heckerman et al., Decision-Theoretic Troubleshooting, 1995

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$$p(x_1,\ldots,x_n) = \prod_{i\in V} p(x_i \mid \mathbf{x}_{\operatorname{Pa}(i)})$$

What is the differential diagnosis?



Fig. 1 The ALARM network representing causal relationships is shown with diagnostic (1), intermediate (0) and measurement (0) nodes. Co cardiac actypt. CVP; entribute nervous pressure, IVED volume: left ventricular enddiastolic volume, IV failure: left ventricular failure, MV: minute ventilation, PA Sat: pulmonary artery angen saturation, PAP: pulmonary artery pressure, RCWP: pulmonary capillary wedge pressure, Pres: breathing pressure, RR: respiratory rater, PR: Vola performary ensistance, TV: Vidal volume

### Beinlich et al., The ALARM Monitoring System, 1989

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- Evidence is denoted by shading in a node
- Can interpret Bayesian network as a **generative process**. For example, to *generate* an e-mail, we

**1** Decide whether it is spam or not spam, by samping  $y \sim p(Y)$ 

**2** For each word i = 1 to n, sample  $x_i \sim p(X_i | Y = y)$ 

# Bayesian network structure implies conditional independencies!



- The joint distribution corresponding to the above BN factors as  $p(d, i, g, s, l) = p(d)p(i)p(g \mid i, d)p(s \mid i)p(l \mid g)$
- However, by the chain rule, any distribution can be written as
  p(d, i, g, s, l) = p(d)p(i | d)p(g | i, d)p(s | i, d, g)p(l | g, d, i, g, s)

• Thus, we are assuming the following additional independencies:  $D \perp I$ ,  $S \perp \{D, G\} \mid I$ ,  $L \perp \{I, D, S\} \mid G$ . What else?

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# Bayesian network structure implies conditional independencies!

- Generalizing the above arguments, we obtain that a variable is independent from its non-descendants given its parents
- Common parent fixing B decouples A and C
- Cascade knowing B decouples A and C

- V-structure Knowing C couples A and B
  - This important phenomona is called **explaining away** and is what makes Bayesian networks so powerful





# A simple justification (for common parent)



We'll show that p(A, C | B) = p(A | B)p(C | B) for any distribution p(A, B, C) that factors according to this graph structure, i.e.

$$p(A, B, C) = p(B)p(A \mid B)p(C \mid B)$$

#### Proof.

$$p(A, C \mid B) = \frac{p(A, B, C)}{p(B)} = p(A \mid B)p(C \mid B)$$

# D-separation ("directed separated") in Bayesian networks

- Algorithm to calculate whether  $X \perp Z \mid \mathbf{Y}$  by looking at graph separation
- Look to see if there is **active path** between X and Z when variables Y are observed:



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- Look to see if there is **active path** between X and Z when variables Y are observed:



- If no such path, then X and Z are **d-separated** with respect to **Y**
- d-separation reduces statistical independencies (hard) to connectivity in graphs (easy)
- Important because it allows us to quickly prune the Bayesian network, finding just the relevant variables for answering a query

## D-separation example 1



Is  $X_6 \perp X_5 \mid X_2, X_3$ ? Is  $X_4 \perp X_5 \mid X_2, X_3$ ?

## D-separation example 2



Is  $X_4 \perp X_5 \mid X_1, X_6$ ?

What about if  $X_6$  is not observed? I.e., is  $X_4 \perp X_5 \mid X_1$ ?

## Independence maps

- Let *I*(*G*) be the set of all conditional independencies implied by the directed acyclic graph (DAG) *G*
- Let I(p) denote the set of all conditional independencies that hold for the joint distribution p.
- A DAG G is an **I-map** (independence map) of a distribution p if  $I(G) \subseteq I(p)$ 
  - A fully connected DAG G is an I-map for any distribution, since  $I(G) = \emptyset \subseteq I(p)$  for all p
- *G* is a **minimal I-map** for *p* if the removal of even a single edge makes it not an I-map
  - A distribution may have several minimal I-maps
  - Each corresponds to a specific node-ordering
- G is a **perfect map** (P-map) for distribution p if I(G) = I(p)

- Different Bayesian network structures can be **equivalent** in that they encode precisely the same conditional independence assertions (and thus the same distributions)
- Which of these are equivalent?



- Different Bayesian network structures can be **equivalent** in that they encode precisely the same conditional independence assertions (and thus the same distributions)
- Are these equivalent?



## 2011 Turing Award was for Bayesian networks



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- Bayesian networks given by (G, P) where P is specified as a set of local conditional probability distributions associated with G's nodes
- One interpretation of a BN is as a **generative model**, where variables are sampled in topological order
- Local and global independence properties identifiable via **d-separation** criteria
- Computing the probability of any assignment is obtained by multiplying CPDs
  - Bayes' rule is used to compute conditional probabilities
  - Marginalization or inference is often computationally difficult