Lecture 2

David A. Levin

University of Oregon

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- In this set-up, we assume the parameter(s) θ has a **prior** distribution $\pi(\theta)$.
- ► Given θ , the random variables $(X_1, ..., X_n)$ have a distribution $f(x_1, ..., x_n | \theta)$.
- ightharpoonup If the distributions of X_i 's are conditionally independent, then

$$f(x_1,\ldots,x_n\mid\theta)=\prod_{i=1}^n f(x_i\mid\theta).$$

$$f(\theta \mid x_1, \dots, x_n) = \frac{f(x_1, \dots, x_n \mid \theta) \pi(\theta)}{\int f(x_1, \dots, x_n \mid \theta) \pi(\theta) d\theta}$$

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► The issue is often computing the normalizing constant in the posterior:

$$\int f(x_1,\ldots,x_n\,|\,\theta)\pi(\theta)\,d\theta$$

- If θ is high-dimensional, this especially can be difficult. Modern Bayesian statistics uses many methods including Markov Chain Monte Carlo to evaluate this constant.
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Suppose that

$$\pi(\mu) = \frac{1}{\sqrt{2\pi\tau^2}} \exp\left(-\frac{1}{2\tau^2}(\mu - \nu)^2\right)$$

That is $\theta \sim N(\nu, \tau^2)$.

Suppose that $X_1, ..., X_n$ are i.i.d.

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

that is, X_i are $N(\mu, \sigma^2)$

- ► Here v, τ^2, σ^2 are assumed known. They are called **hyperparameters**.
- ► Calculating the posterior seems messy but it all works out because the prior is conjugate.

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$$f(\mu \mid x_{1},...,x_{n}) \propto f(x_{1},...,x_{n} \mid \mu)\pi(\mu;\nu,\tau)$$

$$= c(\sigma,\tau) \exp\left(-\frac{1}{2\sigma^{2}} \left(\sum_{i=1}^{n} (x_{i} - \mu)^{2} - \frac{1}{2\tau^{2}} (\mu - \nu)^{2}\right)\right)$$

$$= c(\sigma,\tau,\mathbf{x}) \exp\left(\mu \left(\frac{S_{n}}{2\sigma^{2}} + \frac{\nu}{2\tau^{2}}\right) - \mu^{2} \left(\frac{n}{2\sigma^{2}} + \frac{1}{2\tau^{2}}\right)\right)$$

$$= c(\sigma,\tau,\mathbf{x}) \exp\left[-\frac{1}{2} \frac{n\tau^{2} + \sigma^{2}}{\sigma^{2}\tau^{2}} \left(\mu^{2} - \mu \frac{\sigma^{2}\tau^{2}}{n\tau^{2} + \sigma^{2}} \left(\frac{\bar{x}n\tau^{2} + \nu\sigma^{2}}{\sigma^{2}\tau^{2}}\right)\right)\right]$$

$$= c(\sigma,\tau,\mathbf{x}) \exp\left[-\frac{1}{2\nu(\sigma,\tau)} \left(\mu - \frac{n\bar{x}\tau^{2} + \nu\sigma^{2}}{n\tau^{2} + \sigma^{2}}\right)^{2}\right]$$

All the exponent that does not depend on μ is thrown into the multiplicative constant $c(\sigma, \tau, x)$.

But the only distribution this can be is Normal, with variance $v(\sigma, \tau) = \sigma^2 \tau^2 / (n\tau^2 + \sigma^2)$, and with mean

$$\bar{x} \frac{\tau^2}{\tau^2 + \sigma^2/n} + v \frac{\sigma^2/n}{\tau^2 + \sigma^2/n}$$
.



$$\mathbb{E}[(\mu - T)^2 \mid \mathbf{x}]$$

among statistics T depending on x, then the minimizer is $T = \mathbb{E}[\mu \mid x]$.

▶ In this case, the Bayes estimator is

$$\bar{x}\frac{\tau^2}{\tau^2 + \sigma^2/n} + v\frac{\sigma^2/n}{\tau^2 + \sigma^2/n}$$

- ▶ This is a convex combination of the data-only estimator \bar{X} and the prior mean ν . The weight of \bar{X} tend to 1 as $n \to \infty$.
- ▶ Other inferences are possible, e.g. *credible intervals*, so we can find *a* and *b* so that

$$\mathbb{P}(a < \mu < b \mid \mathbf{x}) = 0.95$$

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- A graphical model specifies a factorization which a joint density must obey.
- Example: If $X_1, ..., X_n$ are independent, then the joint density completely factors into marginals:

$$f(x_1, ..., x_n) = \prod_{i=1}^n f_{X_i}(x_i).$$
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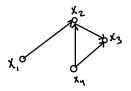
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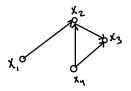
- nodes represent variables
- directed arrows represent dependencies
- Any variable with an arrow pointing to x_i is called a **parent** of x_i ; we denote by pa_i all the parents of x_i .



$$p(x_1,\ldots,x_n)=\prod_{i=1}^n p(x_i\mid \mathsf{pa}_i).$$

$$p(x_1, x_2, x_3, x_4) = p(x_1) p(x_2 \mid x_1, x_4) p(x_3 \mid x_2, x_4) p(x_4).$$

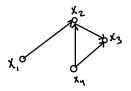
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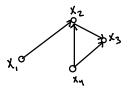
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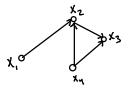


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- ► The graphical model encodes *conditional independence* statements.
- ► A set of variables *X* and *Y* are conditionally independent given *Z* if

$$\mathbb{P}(X \in A, Y \in B \mid Z) = \mathbb{P}(X \in A \mid Z)\mathbb{P}(X \in B \mid Z).$$

► In terms of pdfs/pmfs,

$$p(x,y \mid z) = p(x \mid z)p(y \mid z).$$

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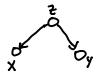
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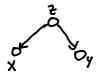




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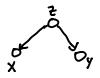
- ▶ Any time *X* and *Y* are separated by a **tail-to-tail** vertex *Z*, they are conditionally independent.
- Note that *X* and *Y* are not independent unconditionally, in general.
- Exercise: Show by example that *X* and *Y* are not necessarily independent.



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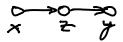


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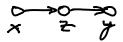




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Thus, if *x* and *y* are connected by a **head-to-tail** vertex *z*, then they are conditionally independent.

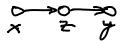
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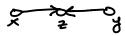
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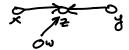
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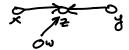
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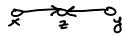
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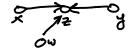
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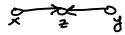




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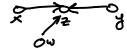
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- Let A, B, C be sets of vertices. The set C blocks A to B if every path from a vertex $a \in A$ to a vertex $b \in B$ contains either
 - a vertex $c \in C$ so that c is head-to-tail or tail-to-tail, or
 - ▶ a vertex *v* which neither belongs to *C* or is a decendent of any vertex in *C*.

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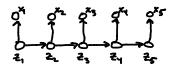
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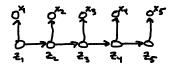
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- ► The above is a **Hidden Markov Model**.
- ► We can use *d*-separation to establish useful conditional independence relations:

$$(X_1,\ldots,X_k)\perp(X_{k+1},\ldots,X_n)\mid Z_k$$

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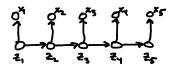


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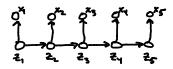
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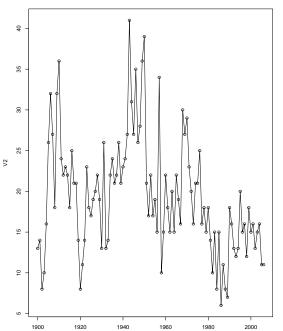
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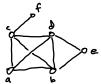
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V1

- Can we do something similar, but for undirected graphs? That is, can we use undirected graphs to specify classes of joint probability distributions, where variables are identified with vertices?
- ▶ A **Markov random field** satisfies a simpler version of the *d*-separation theorem: *A* and *B* are blocked by *C* if every path from *A* to *B* includes a vertex in *C*.
- ► We will first specify a structure for joint distributions whose components are identified with vertices of the graph.
- We will state a theorem which shows the equivalence of Markov random fields with the distributions having the specified product structure.

- A **clique** in a graph is a set of vertices that are maximally connected.
- ► A **maximal** clique in a graph cannot be enlarged by adding another vertex to obtain another clique.
- Below {a, b, c} is a clique, but not maximal, because {a, b, c, d} is a clique containing it. {a, b, c, d} is maximal, because including e or f does not create a clique.



- ► To each maximal clique S in a graph, associate a **potential**, defined as $e^{-H_S(x_S)}$.
- Define a probability by

$$p(\mathbf{x}) = Z^{-1} \prod_{S} e^{H_S(x_S)} = Z^{-1} e^{-\sum_{S} H_S(x_S)}.$$

Theorem 2 (Hammersley-Clifford).

A joint distribution is a Markov random field if and only if it has the product form

$$p(\mathbf{x}) = Z^{-1} \prod_{S} e^{-H_{S}(x_{S})}$$

- The product is over maximal cliques S;
- $x_S = (x_{s_1}, ..., x_{s_r})$ where $S = \{s_1, ..., s_r\};$
- $ightharpoonup Z = \sum_x \prod_S e^{-H_S(x_S)}$ is a normalizing constant; computing Z can be quite difficult or expensive!

- Each variable is ±1;
- \triangleright β is parameter (inverse temperature) representing interaction strength between variables in "base graph";
- Y_v is a "noisy" observation of X_v ; the parameter η determines the "flip" probability.
- ► In image analysis applications, the *Y*'s are observed, and the *X*'s are unobserved.
- ► Total energy is

$$H(\mathbf{x}, \mathbf{y}) = -\beta \sum_{v \sim w} x_v x_w + h \sum_{v} x_v - \eta \sum_{v} x_v y_v$$



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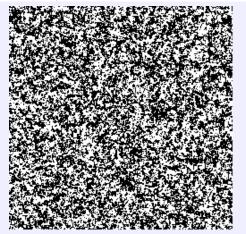
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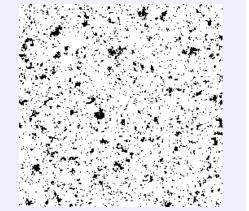
Three regimes

High temperature ($\beta < \beta_c$):



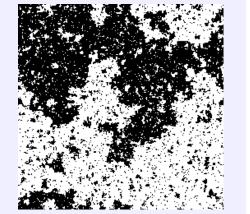
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- ▶ We take a Maximum Likelihood approach; regard the x as parameters, and pick the value x which makes the data y most likely. That is, maximize p(x, y) over all x, holding the observed y fixed.
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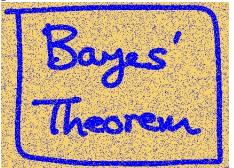
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Example from C. Bishop, *Machine Learning and Pattern Recognition*:
Original image

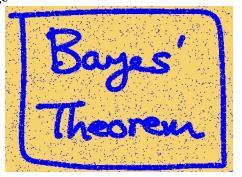


Example from C. Bishop, *Machine Learning and Pattern Recognition*:

Corrupted image



Example from C. Bishop, *Machine Learning and Pattern Recognition*:
Restored image



Message passing and the sum-product algorithm

In many algorithms we will need an effective way to compute marginals in a graphical model:

$$p(x_k) = \sum_{x_j: j \neq k} p(x_1, \dots, x_n).$$

- Note if each variable has say K possible values then this requires a sum over K^{n-1} terms; this grows exponentially as n grows, so does not scale.
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