

### Checkpoint

- Assignment 3 due tonight
- My office hours today at 4:30 in 17 Hillhouse Room 115
- Assignment 4 out tonight, on PCA and logistic regression
- Practice midterm will go out next Thursday
- Midterm in class on Tuesday, October 15
- Questions?

## **For Today**

- PCA (briefly)
- Latent variables
- Mixtures
- Bayes

### **PCA**

PCA finds the directions of greatest variation in the data, and represents the data in terms of those directions.

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### **PCA: Algorithm**

**1** Center the data:  $x_i \mapsto x_i - \overline{x}$ 

xi is d vector xiT is 1xd vector Product is dxd matrix

- ② Compute the  $d \times d$  sample covariance  $S = \frac{1}{n} \sum_{i=1}^{n} x_i x_i^T$
- Find the first k eigenvectors of S
- Project the data onto those k vectors

# Handwritten Digits (3s)

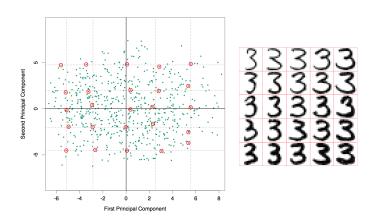
## Handwritten Digits (3s)

$$\hat{f}(\lambda) = \bar{x} + \lambda_1 v_1 + \lambda_2 v_2 
= + \lambda_1 \cdot + \lambda_2 \cdot \cdot \cdot$$

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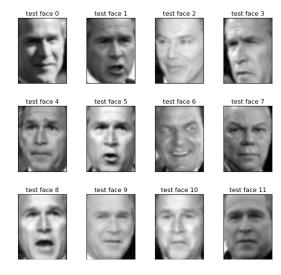
## Handwritten Digits (3s) – Top 2 components

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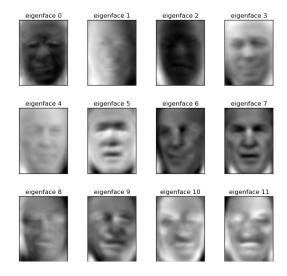


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### **Faces**



## **Eigenfaces**



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#### Genes mirror geography within Europe

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The publisher's final edited version of this article is available at Nature

This article has been corrected. See the correction in volume 456 on page 274.

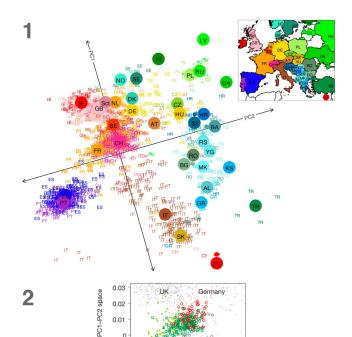
See commentary "Editorial comment should accompany hot papers online," in Nature, volume 455 on page 861.

See other articles in PMC that cite the published article.

#### Abstract

Go to: ☑

Understanding the genetic structure of human populations is of fundamental interest to medical, forensic and anthropological sciences. Advances in high-throughput genotyping technology have markedly improved our understanding of global patterns of human genetic variation and suggest the potential to use large samples to uncover variation among closely spaced populations<sup>1-2</sup>. Here we characterize genetic variation in a sample of 3,000 European individuals genotyped at over half a million variable DNA sites in the human genome. Despite low average levels of genetic differentiation among Europeans, we find a close correspondence between genetic and geographic distances; indeed, a geographical map of Europe arises naturally as an efficient two-dimensional summary of genetic variation in Europeans. The results emphasize that when mapping the genetic basis of a disease phenotype, spurious associations can arise if genetic structure is not properly accounted for. In addition, the results are relevant to the prospects of genetic ancestry testing<sup>6</sup>, an individual's DNA can be used to infer their geographic origin with surprising accuracy—often to within a few hundred kilometres.



## Where are we going?

- So far: Mostly "surface representations" & explicit features
  - Listing information for Zillow
  - Political blogs word counts

## Where are we going?

- So far: Mostly "surface representations" & explicit features
  - Listing information for Zillow
  - Political blogs word counts
- Next: "Hidden" representations and latent variables

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Among the many interesting objects which will engage your attention that of providing for the common defense will merit particular regard. To be prepared for war is one of the most effectual means of preserving peace.

### **Barack Obama**

The latent variable is "war and peace"

### 2009



PCA doesn't pull out a semantic representation, although it does something close to it.

And for the first time, that includes the full cost of fighting in Iraq and Afghanistan. For 7 years, we have been a nation at war. No longer will we hide its price.

## The elephants in the room

When we see image: it is two elephants playing with a ball. The notion of playing is a hidden variable For a computer, it is challenging. So we need latent variables/probabilistic models



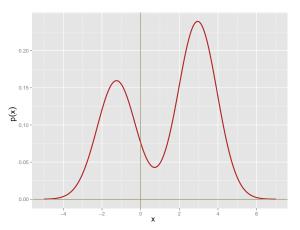
### **Mixtures**

All latent variable models are some sort of mixture model

- Key technique: Mixture models
- Mixtures have latent variables
- Flexible tool
- Simple and difficult at the same time

### **Gaussian Mixture**

#### A mixture of two gaussians



The latent variable comes from the probabilistic interpretation of this model

$$p(x) = \frac{2}{5}\phi(x; -1.25, 1) + \frac{3}{5}\phi(x; 2.95, 1)$$

The two gaussians that were mixed

### **Mixtures**

#### n is the latent variable

Mixture of f and g:

Or similarly, You flip a coin. If it is heads (with probability  $\eta$ ) then sample from f(x). Else, sample from g(x). The coin flip is the latent variable, since it is unseen (you only see the final x that you got from a gaussian).

$$p(x) = \eta f(x) + (1 - \eta)g(x)$$

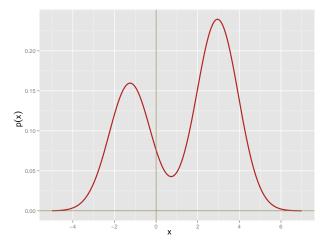
Simplest, most common kind of latent variable model

• Hidden variable represention: Define  $Z \sim \text{Bernoulli}(\eta)$  and

$$p(x) = \sum_{z=0,1} p(x \mid z) p(z)$$

with 
$$p(x | 0) = f(x)$$
,  $p(x | 1) = g(x)$ ,  $p(z) = \eta^{z} (1 - \eta)^{(1-z)}$ .

### **Gaussian Mixture: All the Key Concepts**



### **Bayesian Inference**

Is a coin flip prob of heads /  $\theta$  always 0.5? What if ridges in the coin make it change?

The parameter  $\theta$  of a model is viewed as a random variable. Inference usually carried out as follows:

Assume a distribution over your param  $\theta$  i.e. no longer fixed Flip the coin many times to gather more evidence, and update your beliefs about  $\theta$ 

- Choose a *generative model*  $p(x | \theta)$  for the data.
- Choose a *prior distribution*  $\pi(\theta)$  that expresses beliefs about the parameter before seeing any data.
- After observing data  $\mathcal{D}_n = \{x_1, \dots, x_n\}$ , update beliefs and calculate the *posterior distribution*  $p(\theta \mid \mathcal{D}_n)$ .

```
1. Form some prior beliefs, i.e. \pi(\theta)
```

Bayes rule:  $P(A \mid B) = P(A \& B) / P(B)$  $P(B \mid A) = P(A \mid B) * P(B) / P(A)$ 

<sup>2.</sup> Then, consider model:  $p(x|\theta)$ 

<sup>3.</sup> Compute posterior distribution over  $\boldsymbol{\theta}$  using Bayes rule

### Bayes' Theorem

The posterior distribution can be written as

Posterior 
$$p(\theta \mid x_1, \dots, x_n) = \frac{p(x_1, \dots, x_n \mid \theta)\pi(\theta)}{p(x_1, \dots, x_n)} = \frac{p(x_1, \dots, x_n \mid \theta)\pi(\theta)}{p(x_1, \dots, x_n)} = \frac{\mathcal{L}_n(\theta)\pi(\theta)}{c_n} \propto \mathcal{L}_n(\theta)\pi(\theta)$$

where  $\mathcal{L}_n(\theta) = \prod_{i=1}^n p(x_i \mid \theta)$  is the *likelihood function* and

$$c_n = p(x_1, \ldots, x_n) = \int p(x_1, \ldots, x_n | \theta) \pi(\theta) d\theta = \int \mathcal{L}_n(\theta) \pi(\theta) d\theta$$

is the normalizing constant, which is also called evidence.

Sometimes also called marginal likelihood?

### **Example**



Assume the data comes from a Bernoulli distribution over Θ

 $X \sim \text{Bernoulli}(\theta)$  with data  $\mathcal{D}_n = \{x_1, \dots, x_n\}$ . Prior Beta $(\alpha, \beta)$  distribution

$$\pi_{\alpha,\beta}(\theta) = \frac{\Gamma(\alpha+\beta)}{\Gamma(\alpha)\Gamma(\beta)} \theta^{\alpha-1} (1-\theta)^{\beta-1}$$

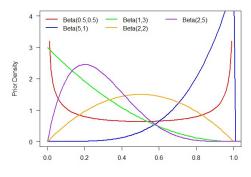
Let  $s = \sum_{i=1}^{n} x_i$  be the number of "successes."

Posterior distribution  $\theta \mid \mathcal{D}_n$  is Beta( $\alpha + s, \beta + n - s$ ). Posterior mean is a mixture:

$$\bar{\theta} = \frac{\alpha + s}{\alpha + \beta + n} = \left(\frac{n}{\alpha + \beta + n}\right)\widehat{\theta} + \left(\frac{\alpha + \beta}{\alpha + \beta + n}\right)\theta_0$$

where  $\hat{\theta} = s/n$  is the MLE and  $\theta_0 = \alpha/(\alpha + \beta)$  is the prior mean.

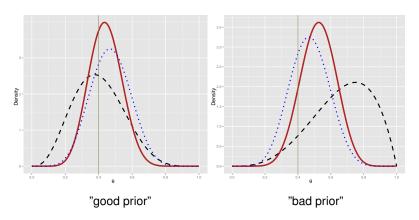
#### **Prior Distributions**



#### Example of beta priors

### **Example**

n= 15 points sampled as  $X\sim \text{Bernoulli}(\theta=0.4)$ , with s= 7 heads.



Prior distribution (black-dashed), likelihood function (blue-dotted), posterior distribution (red-solid).

### **Dirichlet**

Fancier coin flipping models i.e. rolling a die

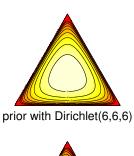
Multinomial model with Dirichlet prior is generalization of the Bernoulli/Beta model.

$$\mathsf{Dirichlet}_{\alpha}(\theta) = \frac{\Gamma(\sum_{j=1}^{K} \alpha_j)}{\prod_{j=1}^{K} \Gamma(\alpha_j)} \prod_{j=1}^{K} \theta_j^{\alpha_j - 1}$$

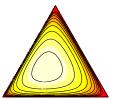
where  $\alpha = (\alpha_1, \dots, \alpha_K) \in \mathbb{R}_+^K$  is a non-negative vector.

### **Example**

#### Likelihood function is just p(x I Θ)









posterior distribution with n = 20

posterior distribution with n = 200

### Reading

- Please be sure to read the detailed notes on Bayesian inference that have been posted to Canvas
- Not necessary to understand everything—but you should be able to do some of the basic "coin flipping" calculations
- We'll need this when discussing topic models

### **Summary**

- PCA is classical and modern—key tool for ML
- Mixtures are latent variable models
- The mixing weight encodes a hidden variable
- Computing with mixtures uses basic probabilistic reasoning