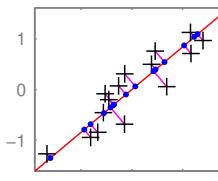


PCA: Principal Component Analysis

Iain Murray
<http://iainmurray.net/>

PCA: Principal Component Analysis

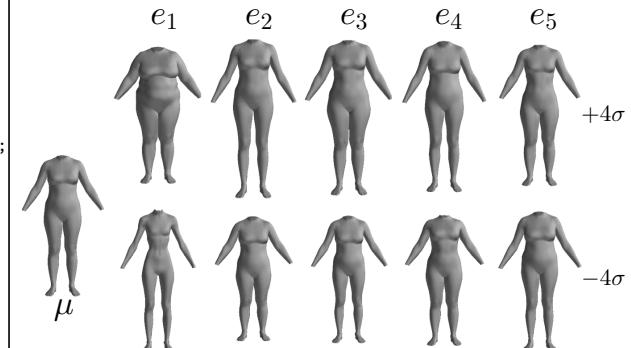


Code assuming X is zero-mean

```
% Find top K principal directions:  
[V, E] = eig(X'*X);  
[E,id] = sort(diag(E),1,'descend');  
V = V(:, id(1:K)); % DxK  
  
% Project to K-dims:  
X_kdim = X*V; % NxK  
  
% Project back:  
X_proj = X_kdim * V'; % NxD
```

K = 1
+ = X
• = Xproj
— = V(:,1)

PCA applied to bodies



Freifeld and Black, ECCV 2012

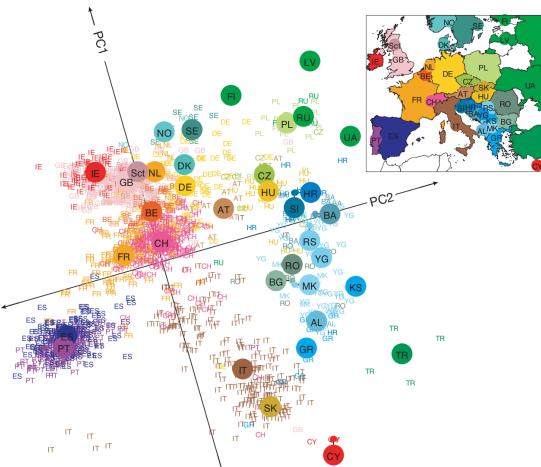
PCA applied to DNA

Novembre et al. (2008) — doi:10.1038/nature07331
Carefully selected both individuals and features

1,387 individuals

197,146 single nucleotide polymorphisms (SNPs)

Each person reduced to two(!) numbers with PCA



MSc course enrollment data

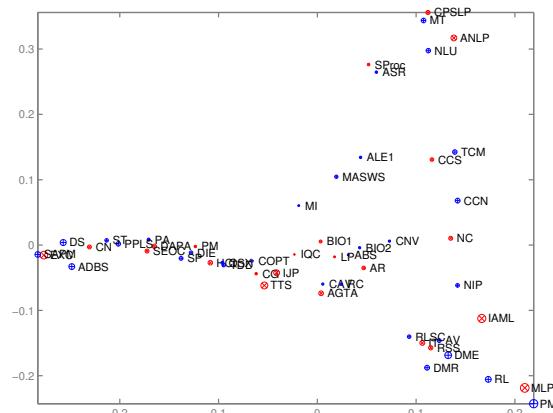
Binary $S \times C$ matrix M

$M_{sc} = 1$, if student s taking course c

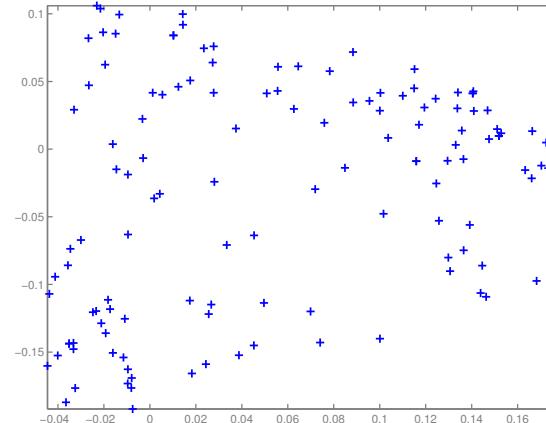
Each course is a length S vector

. . . OR each student is a length C vector

PCA applied to MSc courses



PCA applied to MSc students



Truncated SVD

$$\begin{aligned} &\% \text{PCA via SVD}, \\ &\% \text{for zero-mean } X: \\ &[U, S, V] = svd(X, 0); \\ &U = U(:, 1:K); \\ &S = S(1:K, 1:K); \\ &V = V(:, 1:K); \\ &X_kdim = U*S; \\ &X_proj = U*S*V'; \end{aligned}$$

$$\left[\begin{array}{cccc} X_{11} & X_{12} & \dots & X_{1D} \\ X_{21} & X_{22} & \dots & X_{2D} \\ X_{31} & X_{32} & \dots & X_{3D} \\ X_{41} & X_{42} & \dots & X_{4D} \\ X_{51} & X_{52} & \dots & X_{5D} \\ \vdots & \vdots & \ddots & \vdots \\ X_{N1} & X_{N2} & \dots & X_{ND} \end{array} \right] \approx \left[\begin{array}{c} U_{11} & \dots & U_{1K} \\ U_{21} & \dots & U_{2K} \\ U_{31} & \dots & U_{3K} \\ U_{41} & \dots & U_{4K} \\ U_{51} & \dots & U_{5K} \\ \vdots & \ddots & \vdots \\ U_{N1} & \dots & U_{NK} \end{array} \right] \left[\begin{array}{ccc} S_{11} & 0 & 0 \\ 0 & \ddots & 0 \\ 0 & 0 & S_{KK} \end{array} \right] \left[\begin{array}{ccc} V_{11} & V_{21} & \dots & V_{D1} \\ \vdots & \vdots & \ddots & \vdots \\ V_{1K} & V_{2K} & \dots & V_{DK} \end{array} \right]$$

$$X \approx U S V^\top$$

PCA summary

Project data onto major axes of covariance
 $X^\top X$ is covariance if make data zero mean

Low-dim coordinates can be useful:
— visualization
— if can't cope with high-dim data

Can project back into original space:
— detail is lost: still in K -dim subspace
— PCA minimizes the square error

PPCA: Probabilistic PCA

Gaussian model: $\Sigma = WW^\top + \sigma^2 I$

W is $D \times K$, σ^2 small \Rightarrow nearly low-rank

W is also orthogonal

As $\sigma^2 \rightarrow 0$, recover PCA.

Need $\sigma^2 > 0$ to explain data

Special case of factor analysis: $\Sigma = WW^\top + \Phi$, with Φ diagonal

Dim reduction in other models

Can replace x with Ax in any model

A is a $K \times D$ matrix of projection params

Large D : a lot of extra parameters

NB: Neural nets already have such projections

Practical tip

Scale features to have unit variance

Equivalently: find eigenvectors of correlation rather than covariance

Avoids issues with (arbitrary?) scaling.

If multiply feature by 10^9 , PC points along that feature

E.g., if change unit of feature from metres to nanometres