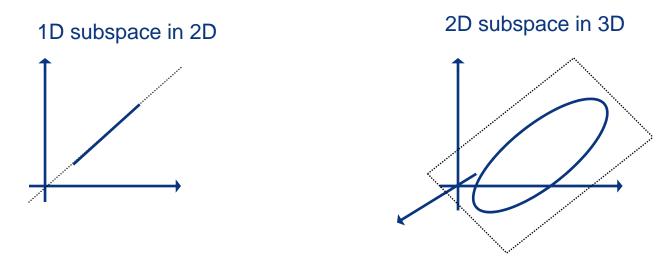
PCA and LDA

Nuno Vasconcelos ECE Department, UCSD

Principal component analysis

▶ basic idea:

 if the data lives in a subspace, it is going to look very flat when viewed from the full space, e.g.

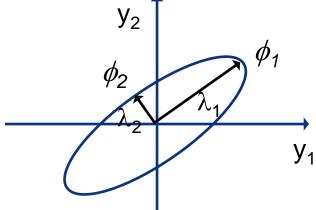


 this means that if we fit a Gaussian to the data the equiprobability contours are going to be highly skewed ellipsoids

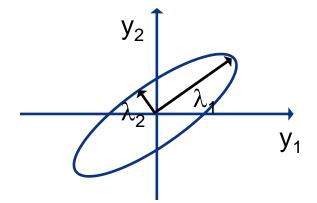
Principal component analysis

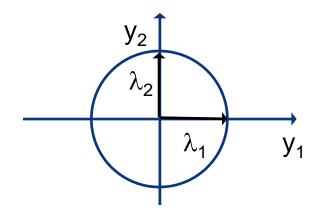
▶ If y is Gaussian with covariance Σ , the equiprobability contours are the ellipses whose y_2 ↑

- principal components ϕ_i are the eigenvectors of Σ
- principal lengths λ_i are the eigenvalues of Σ



by computing the eigenvalues we know if the data is flat $\lambda_1 >> \lambda_2$: flat $\lambda_1 = \lambda_2$: not flat





Principal component analysis (learning)

- ▶ Given sample $\mathcal{D} = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}, \ x_i \in \mathcal{R}^d$
 - compute sample mean: $\hat{\mu} = \frac{1}{n} \sum_{i} (\mathbf{x}_i)$
 - compute sample covariance: $\hat{\Sigma} = \frac{1}{n} \sum_{i} (\mathbf{x}_i \hat{\mu}) (\mathbf{x}_i \hat{\mu})^T$
 - \bullet compute eigenvalues and eigenvectors of $\hat{\Sigma}$

$$\hat{\Sigma} = \Phi \Lambda \Phi^T, \ \Lambda = diag(\sigma_1^2, \dots, \sigma_n^2) \ \Phi^T \Phi = I$$

- order eigenvalues $\sigma_1^2 > ... > \sigma_n^2$
- if, for a certain k, $\sigma_k << \sigma_1$ eliminate the eigenvalues and eigenvectors above k.

Principal component analysis

- Given principal components $\phi_i, i \in 1, ..., k$ and a test sample $\mathcal{T} = \{\mathbf{t}_1, ..., \mathbf{t}_n\}, \ t_i \in \mathcal{R}^d$
 - subtract mean to each point $\mathbf{t}_i' = \mathbf{t}_i \hat{\mu}$
 - ullet project onto eigenvector space $\mathbf{y}_i = \mathbf{A}\mathbf{t}_i'$ where

$$\mathbf{A} = \left[\begin{array}{c} \phi_1^T \\ \vdots \\ \phi_k^T \end{array} \right]$$

• use $T' = \{y_1, \dots y_n\}$ to estimate class conditional densities and do all further processing on \mathbf{y} .

Principal component analysis

- ▶ there is an alternative manner to compute the principal components, based on singular value decomposition
- ► SVD:
 - any real n x m matrix (n>m) can be decomposed as

$$A = M\Pi N^T$$

- where M is a n x m column orthonormal matrix of left singular vectors (columns of M)
- Π a m x m diagonal matrix of singular values
- N^T a m x m row orthonormal matrix of right singular vectors (columns of N)

$$M^T M = I$$
 $N^T N = I$

▶ to relate this to PCA, we consider the data matrix

$$X = \begin{bmatrix} | & & | \\ x_1 & \dots & x_n \\ | & & | \end{bmatrix}$$

▶ the sample mean is

$$\mu = \frac{1}{n} \sum_{i} x_{i} = \frac{1}{n} \begin{bmatrix} 1 & & & & \\ x_{1} & \dots & x_{n} \\ 1 & & & \end{bmatrix} \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix} = \frac{1}{n} X 1$$

- and we can center the data by subtracting the mean to each column of X
- ▶ this is the centered data matrix

$$X_{c} = \begin{bmatrix} | & | & | \\ X_{1} & \dots & X_{n} \\ | & | \end{bmatrix} - \begin{bmatrix} | & | & | \\ \mu & \dots & \mu \\ | & | \end{bmatrix}$$
$$= X - \mu \mathbf{1}^{T} = X - \frac{1}{n} X \mathbf{1} \mathbf{1}^{T} = X \left(I - \frac{1}{n} \mathbf{1} \mathbf{1}^{T} \right)$$

▶ the sample covariance is

$$\Sigma = \frac{1}{n} \sum_{i} (x_{i} - \mu)(x_{i} - \mu)^{T} = \frac{1}{n} \sum_{i} x_{i}^{c} (x_{i}^{c})^{T}$$

where x_i^c is the ith column of X_c

▶ this can be written as

$$\Sigma = \frac{1}{n} \begin{bmatrix} | & & | \\ x_1^c & \dots & x_n^c \\ | & | \end{bmatrix} \begin{bmatrix} - & x_1^c & - \\ & \vdots & \\ - & x_n^c & - \end{bmatrix} = \frac{1}{n} X_c X_c^T$$

▶ the matrix

$$\boldsymbol{X}_{c}^{T} = \begin{bmatrix} - & \boldsymbol{X}_{1}^{c} & - \\ & \vdots & \\ - & \boldsymbol{X}_{n}^{c} & - \end{bmatrix}$$

is real n x d. Assuming n > d it has SVD decomposition

$$X_c^T = M\Pi N^T$$

$$M^T M = I$$
 $N^T N = I$

and

$$\Sigma = \frac{1}{n} X_c X_c^T = \frac{1}{n} N \Pi M^T M \Pi N^T = \frac{1}{n} N \Pi^2 N^T$$

$$\Sigma = \mathbf{N} \left(\frac{1}{n} \Pi^2 \right) \mathbf{N}^T$$

- ▶ noting that N is d x d and orthonormal, and Π^2 diagonal, shows that this is just the eigenvalue decomposition of Σ
- ▶ it follows that
 - the eigenvectors of Σ are the columns of N
 - the eigenvalues of Σ are

$$\lambda_i = \frac{1}{n} \pi_i^2$$

▶ this gives an alternative algorithm for PCA

- computation of PCA by SVD
- ▶ given X with one example per column
 - 1) create the centered data-matrix

$$\boldsymbol{X}_{c}^{T} = \left(\boldsymbol{I} - \frac{1}{n} \boldsymbol{1} \boldsymbol{1}^{T}\right) \boldsymbol{X}^{T}$$

• 2) compute its SVD

$$X_c^T = M\Pi N^T$$

• 3) principal components are columns of N, eigenvalues are

$$\lambda_i = \frac{1}{n} \pi_i^2$$

Limitations of PCA

- ▶ PCA is not optimal for classification
 - note that there is no mention of the class label in the definition of PCA
 - keeping the dimensions of largest energy (variance) is a good idea, but not always enough
 - certainly improves the density estimation, since space has smaller dimension
 - but could be unwise from a classification point of view
 - the discriminant dimensions could be thrown out
- ▶ it is not hard to construct examples where PCA is the worst possible thing we could do

Example

- consider a problem with
 - two n-D Gaussian classes with covariance $\Sigma = \sigma^2 I$, $\sigma^2 = 10$

$$X \sim N(\mu_i, 10I)$$

we add an extra variable which is the class label itself

$$X' = [X, i]$$

• assuming that $P_Y(0)=P_Y(1)=0.5$

$$E[Y] = 0.5 \times 0 + 0.5 \times 1 = 0.5$$

$$var[Y] = 0.5 \times (0 - 0.5)^{2} + 0.5 \times (1 - 0.5)^{2}$$

$$= 0.125 < 10$$

 dimension n+1 has the smallest variance and is the first to be discarded!

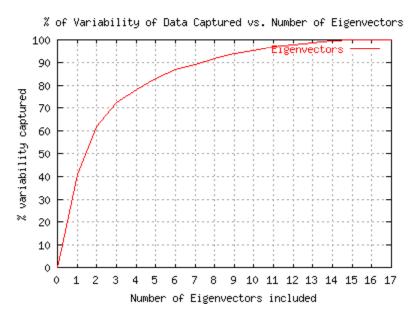
Example

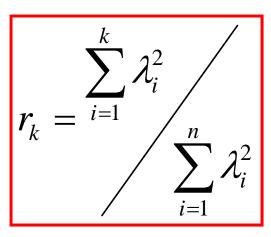
▶ this is

- a very contrived example
- but shows that PCA can throw away all the discriminant info
- does this mean you should never use PCA?
 - no, typically it is a good method to find a suitable subset of variables, as long as you are not too greedy
 - e.g. if you start with n = 100, and know that there are only 5
 variables of interest
 - picking the top 20 PCA components is likely to keep the desired 5
 - your classifier will be much better than for n = 100, probably not much worse than the one with the best 5 features
- ▶ is there a rule of thumb for finding the number of PCA components?

Principal component analysis

- a natural measure is to pick the eigenvectors that explain p % of the data variability
 - can be done by plotting the ratio r_k as a function of k

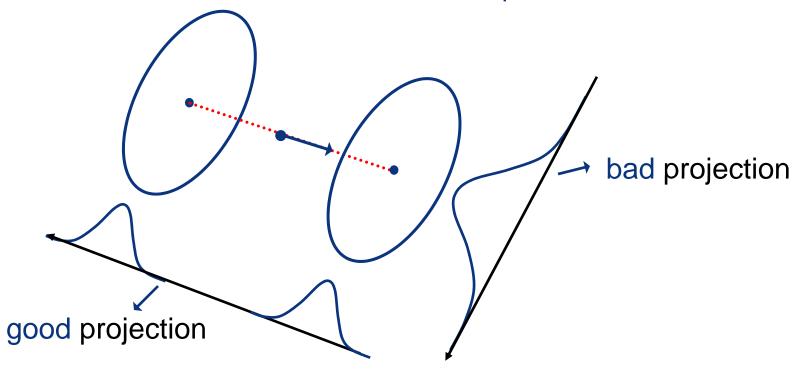




 e.g. we need 3 eigenvectors to cover 70% of the variability of this dataset

Fischer's linear discriminant

- what if we really need to find the best features?
 - harder question, usually impossible with simple methods
 - there are better methods at finding discriminant directions
- ▶ one good example is linear discriminant analysis (LDA)
 - the idea is to find the line that best separates the two classes



we have two classes such that

$$\begin{split} E_{X|Y}\big[X\mid Y=i\big] &= \mu_i \\ E_{X|Y}\big[\big(X-\mu_i\big)\big(X-\mu_i\big)^T\mid Y=i\big] &= \Sigma_i \end{split}$$

and want to find the line

$$z = w^T x$$

that best separates them

one possibility would be to maximize

$$(E_{Z|Y}[Z|Y=1] - E_{Z|Y}[Z|Y=0])^{2} =$$

$$(E_{X|Y}[w^{T}x|Y=1] - E_{X|Y}[w^{T}x|Y=0])^{2} = (w^{T}[\mu_{1} - \mu_{0}])^{2}$$

• however, this $(w^T [\mu_1 - \mu_0])^2$

can be made arbitrarily large by simply scaling w

- ▶ we are only interested in the direction, not the magnitude
- need some type of normalization
- Fischer suggested

$$\max \frac{between\ class\ scatter}{within\ class\ scatter} = \\ \max_{w} \frac{\left(E_{Z|Y}[Z\mid Y=1] - E_{Z|Y}[Z\mid Y=0]\right)^{2}}{\text{var}[Z\mid Y=1] + \text{var}[Z\mid Y=0]}$$

we have already seen that

$$(E_{Z|Y}[Z|Y=1] - E_{Z|Y}[Z|Y=0])^{2} = (w^{T}[\mu_{1} - \mu_{0}])^{2}$$
$$= w^{T}[\mu_{1} - \mu_{0}][\mu_{1} - \mu_{0}]^{T} w$$

also

$$\begin{aligned} & \text{var} \big[Z \mid Y = i \big] = E_{Z \mid Y} \left\{ \! \left(z - E_{Z \mid Y} \big[Z \mid Y = i \big] \! \right)^2 \mid Y = i \right\} \\ &= E_{Z \mid Y} \left\{ \! \left(w^T \big[x - \mu_i \big] \! \right)^2 \mid Y = i \right\} \\ &= E_{Z \mid Y} \left\{ \! w^T \big[x - \mu_i \big] \! \big[x - \mu_i \big]^T w \mid Y = i \right\} \\ &= w^T \Sigma_i w \end{aligned}$$

and

$$J(w) = \frac{\left(E_{Z|Y}[Z|Y=1] - E_{Z|Y}[Z|Y=0]\right)^{2}}{\text{var}[Z|Y=1] + \text{var}[Z|Y=0]}$$
$$= \frac{w^{T}(\mu_{1} - \mu_{0})(\mu_{1} - \mu_{0})^{T} w}{w^{T}(\Sigma_{1} + \Sigma_{0})w}$$

which can be written as

between class scatter

$$J(w) = \frac{w^T S_B w}{w^T S_W w}$$

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$$S_B = (\mu_1 - \mu_0)(\mu_1 - \mu_0)^T$$

$$S_W = (\Sigma_1 + \Sigma_0)$$

within class scatter

maximizing the ratio

$$J(w) = \frac{w^T S_B w}{w^T S_W w}$$

• is equivalent to maximizing the numerator while keeping the denominator constant, i.e.

$$\max_{w} w^{T} S_{B} w \quad \text{subject to} \quad w^{T} S_{W} w = K$$

- and can be accomplished using Lagrange multipliers
- define the Lagrangian

$$L = w^T S_B w - \lambda (w^T S_W w - K)$$

and maximize with respect to both w and λ

setting the gradient of

$$L = w^{T} (S_{B} - \lambda S_{W}) w + \lambda K$$

with respect to w to zero we get

$$\nabla_{w} L = 2(S_{B} - \lambda S_{W}) w = 0$$

or

$$S_B w = \lambda S_W w$$

- ▶ this is a generalized eigenvalue problem
- ▶ the solution is easy when $S_w^{-1} = (\Sigma_1 + \Sigma_0)^{-1}$ exists

▶ in this case

$$S_W^{-1}S_B w = \lambda w$$

▶ and, using the definition of S_B

$$S_W^{-1}(\mu_1 - \mu_0)(\mu_1 - \mu_0)^T w = \lambda w$$

▶ noting that $(\mu_1 - \mu_0)^T w = \alpha$ is a scalar this can be written as

$$S_W^{-1}(\mu_1 - \mu_0) = \frac{\lambda}{\alpha} w$$

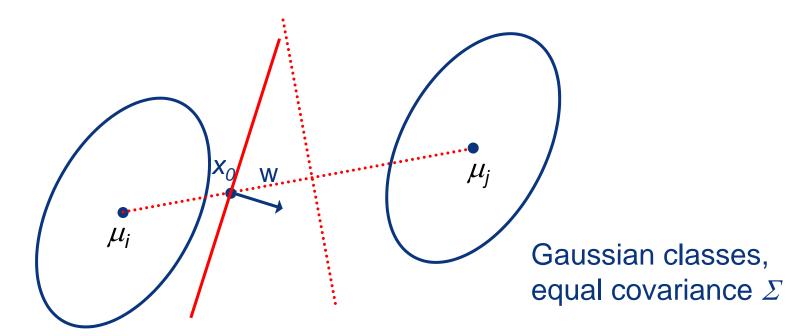
▶ and since we don't care about the magnitude of w

$$w^* = S_W^{-1}(\mu_1 - \mu_0) = (\Sigma_1 + \Sigma_0)^{-1}(\mu_1 - \mu_0)$$

- ▶ note that we have seen this before
 - for a classification problem with Gaussian classes of equal covariance $\Sigma_i = \Sigma$, the BDR boundary is the plane of normal

$$w = \Sigma^{-1} \left(\mu_i - \mu_j \right)$$

• if $\Sigma_1 = \Sigma_0$, this is also the LDA solution



- ▶ this gives two different interpretations of LDA
 - it is optimal if and only if the classes are Gaussian and have equal covariance
 - better than PCA, but not necessarily good enough
 - a classifier on the LDA feature, is equivalent to
 - the BDR after the approximation of the data by two Gaussians with equal covariance

