

CO902  
Supporting Lecture Notes:  
Principal Components Analysis Redux  
Lecture 6 — 11 Feb 2013

Due to some inconsistencies in notations and one derivation malfunction, this handout summarizes the key results for Principal Components Analysis (PCA) as a data reduction technique.

## 1 Preliminaries

First some notation and key results from linear algebra. Breaking from the slides and my own boardwork, here I will use bold capital Greek letters for matrices (e.g.  $n \times n$  matrix  $\mathbf{A}$ ) and bold lower case Greek letters for vectors (e.g.  $n$ -vector  $\mathbf{u}$ ).

**Eigendecomposition of a square matrix.** For a  $n \times n$  matrix  $\mathbf{A}$ , a length- $n$  column vector  $\mathbf{u}$  and scalar  $\lambda$  that satisfy

$$\mathbf{A}\mathbf{u} = \lambda\mathbf{u}$$

are an eigenvector-eigenvalue pair. For real symmetric  $\mathbf{A}$  (the only kind we're concerned with) the eigenvectors can be chosen to be real, orthogonal and to have unit length, i.e.  $\mathbf{u}_j^\top \mathbf{u}_{j'} = 1$  for  $j \neq j'$  and  $\mathbf{u}_j^\top \mathbf{u}_j = 1$ . Collecting all  $n$  eigenvectors into a  $n \times n$  matrix

$$\mathbf{U} = [\mathbf{u}_1 \ \mathbf{u}_2 \ \cdots \ \mathbf{u}_n],$$

and putting the corresponding eigenvalues into a  $n \times n$  diagonal matrix  $\mathbf{\Lambda}$  gives

$$\mathbf{\Lambda} = \text{diag}(\lambda_1, \lambda_2, \dots, \lambda_n).$$

The set of eigenvectors and eigenvalues then gives

$$\mathbf{A}\mathbf{U} = \mathbf{U}\mathbf{\Lambda}.$$

The orthogonality and unit length<sup>1</sup> means that  $\mathbf{U}$  is orthonormal and  $\mathbf{U}^\top \mathbf{U} = \mathbf{U}\mathbf{U}^\top = \mathbf{I}$ , where  $\mathbf{I}$  is the identity matrix. Premultiplying by  $\mathbf{U}^\top$  shows that

$$\mathbf{U}^\top \mathbf{A}\mathbf{U} = \mathbf{\Lambda}.$$

the matrix  $\mathbf{A}$  can be diagonalized with its eigenvectors, and postmultiplying by  $\mathbf{U}^\top$  shows

$$\mathbf{A} = \mathbf{U}\mathbf{\Lambda}\mathbf{U}^\top = \sum_{i=1}^n \lambda_i \mathbf{u}\mathbf{u}^\top \tag{1}$$

which explicitly shows that  $\mathbf{A}$  is the sum of  $n$  rank-1 matrices.

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<sup>1</sup>And an assumption of non-repeated eigenvalues, which is going to be the case for real data.

## 2 Unsupervised Learning - Notation

Let the data of interest be length- $d$  column vectors  $x_i, i = 1, \dots, n$ , assembled into a  $d \times n$  data matrix  $\mathbf{X}$ ,

$$\mathbf{X} = [\mathbf{x}_1 \mathbf{x}_2 \cdots \mathbf{x}_n].$$

The average data vector is

$$\bar{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^n \mathbf{x}_i = \frac{1}{n} \mathbf{X} \mathbf{1},$$

where  $\mathbf{1}$  is a length- $n$  column vector of ones, and

$$\mathbf{X} - \bar{\mathbf{X}} = [\mathbf{x}_1 - \bar{\mathbf{x}}, \mathbf{x}_2 - \bar{\mathbf{x}}, \dots, \mathbf{x}_n - \bar{\mathbf{x}}] = \mathbf{X} - \frac{1}{n} \mathbf{X} \mathbf{1} \mathbf{1}^\top$$

is the centered data matrix. The  $d \times d$  sample covariance matrix of  $\mathbf{X}$  is

$$\mathbf{S} = \frac{1}{n} \sum_{i=1}^n (\mathbf{x}_i - \bar{\mathbf{x}})(\mathbf{x}_i - \bar{\mathbf{x}})^\top = \frac{1}{n} (\mathbf{X} - \bar{\mathbf{X}})(\mathbf{X} - \bar{\mathbf{X}})^\top.$$

(Please make sure you're comfortable with all these alternative ways of writing the same expression!)

## 3 Principal Components Analysis - The crux of it

PCA amounts to finding a  $k \times d$  matrix  $\mathbf{U}$  such that the data projected into  $k$  dimensions,

$$\mathbf{y}_i = \mathbf{U}^\top \mathbf{x}_i,$$

is as “variable” as possible<sup>2</sup>. Here, variability is defined by the average squared L2 norm of  $\mathbf{y}_i$  from its mean, or, equivalently, as the sum of variances in each of the  $d$  dimensions.

Writing the  $k \times n$  reduced data matrix

$$\mathbf{Y} = [\mathbf{y}_1 \mathbf{y}_2 \cdots \mathbf{y}_n].$$

and its mean

$$\bar{\mathbf{y}} = \frac{1}{n} \sum_{i=1}^n \mathbf{y}_i = \frac{1}{n} \mathbf{Y} \mathbf{1},$$

this aforementioned variance is

$$\frac{1}{n} \sum_{i=1}^n \|\mathbf{y}_i - \bar{\mathbf{y}}\|^2 = \frac{1}{n} \sum_{i=1}^n (\mathbf{y}_i - \bar{\mathbf{y}})^\top (\mathbf{y}_i - \bar{\mathbf{y}}) = \frac{1}{n} \text{tr}((\mathbf{Y} - \bar{\mathbf{Y}})(\mathbf{Y} - \bar{\mathbf{Y}})^\top).$$

Using the tricks shown in the class notes, you can show that the  $k$  dimensional transform  $\mathbf{U}$  that maximizes this variance are the *first  $k$  eigenvectors* of  $\mathbf{S}$ , where eigenvectors are sorted by eigenvalues, largest to smallest. See, also, the class notes for the derivation of how this same choice of  $\mathbf{U}$  minimizes the approximation error for a  $k$  dimensional basis set.

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<sup>2</sup>Note that  $\mathbf{y}_i$  no longer refers to “outcome” or response, as it did with supervised learning.

## 4 Principal Components Analysis - Useful Observations

The class notes show that the optimized variance measure for a  $k$ -dimensional PCA is

$$\frac{1}{n} \sum_{i=1}^n \|\mathbf{y}_i - \bar{\mathbf{y}}\|^2 = \sum_{j=1}^k \lambda_j.$$

That is, the contribution of the variance of the  $j$ th component is precisely  $\lambda_j$ . This tells us about the relative importance of each component.

**Eigenvalue plots.** To understand the relative contribution of each component  $\mathbf{u}_j$ , we make plots of the ordered eigenvalues  $\lambda_j$  (largest to smallest) and their cumulative variance. The eigenvalue plot (sometimes called a “scree plot”) shows how much variance explained by each component. The cumulative variance plot, a graph of

$$\sum_{j=1}^k \lambda_j / \sum_{j=1}^d \lambda_j$$

versus  $k$ , shows the proportion of variance is explained by the first  $k$  components. This plot is useful when thinking of PCA as a  $k$ -dimensional representation of the full  $d$  dimensional data.

**PCA as a transformation?** What if we choose  $k = d$ , i.e., made a full-rank transformation, no data reduction. Then all we’ve done is change coordinates. What is the  $d \times d$  sample covariance matrix of this  $\mathbf{Y}$ ? Using the fact  $\bar{\mathbf{Y}} = \mathbf{Y}\mathbf{1}\mathbf{1}^\top/n = \mathbf{U}^\top \mathbf{X}\mathbf{1}\mathbf{1}^\top/n = \mathbf{U}^\top \bar{\mathbf{X}}$ ,

$$\begin{aligned} \frac{1}{n}(\mathbf{Y} - \bar{\mathbf{Y}})(\mathbf{Y} - \bar{\mathbf{Y}})^\top &= \frac{1}{n}(\mathbf{U}^\top \mathbf{X} - \mathbf{U}^\top \bar{\mathbf{X}})(\mathbf{U}^\top \mathbf{X} - \mathbf{U}^\top \bar{\mathbf{X}})^\top \\ &= \frac{1}{n} \mathbf{U}^\top (\mathbf{X} - \bar{\mathbf{X}})(\mathbf{X} - \bar{\mathbf{X}})^\top \mathbf{U} \\ &= \mathbf{U}^\top \mathbf{S} \mathbf{U}. \\ &= \mathbf{\Lambda}, \end{aligned}$$

where, since  $k = d$ ,  $\mathbf{\Lambda}$  is the  $d \times d$  diagonal matrix of eigenvalues, using result (1) above. This shows that when you use a full rank transformation you “diagonalise” or “whiten” the data, as the covariances between the  $k = d$  variables are now zero.

**Approximating individual cases.** While usually our interest will be working with the transformed  $k$ -dimensional data  $\{\mathbf{y}_i\}$ , we may want to see what we’ve lost, i.e. how close the PCA is approximating the data  $\{\mathbf{x}_i\}$ . The class notes show that data for the  $i$ -th case can be approximated

$$\hat{\mathbf{x}}_i = \bar{\mathbf{x}} + \sum_{j=1}^k \alpha_{ij} \mathbf{u}_j$$

where  $\alpha_{ij}$  are the approximating coefficients for case  $i$  basis element  $j$ ,

$$\alpha_{ij} = \mathbf{u}_j^\top (\mathbf{x}_i - \bar{\mathbf{x}}).$$

Putting these two together in “matrix mode” gives the approximation for all  $n$  cases

$$\hat{\mathbf{X}} = \bar{\mathbf{X}} + \mathbf{U}\mathbf{U}^\top (\mathbf{X} - \bar{\mathbf{X}})$$

## 5 PCA via SVD

The two “take away” messages about PCA are (1) it is the best  $k$ -dimensional approximation ( $k \times n$   $\mathbf{Y}$ ) to  $d$  dimensional dataset ( $d \times n$   $\mathbf{X}$ ), and (2) it is obtained through the eigenvectors of the  $d \times d$  sample covariance matrix of  $\mathbf{X}$ ,

$$\mathbf{S} = \frac{1}{n}(\mathbf{X} - \bar{\mathbf{X}})(\mathbf{X} - \bar{\mathbf{X}})^\top.$$

Notice that we haven’t made any comment about whether  $\mathbf{S}$  is invertable? That’s because it doesn’t matter; we never need to invert  $\mathbf{S}$  and if  $d \geq n$   $\mathbf{S}$  won’t be full rank and one or more eigenvalues will be zero. No big deal... those would be the last eigenvectors to consider in an approximation anyway.

A practical issue arises, however, if  $d \gg n$ , particularly when  $n$  is very small (10-20) and  $d$  is very large 100 or more. In that case, we are forming a gigantic  $d \times d$  matrix  $\mathbf{S}$  when we know the rank can be no more than  $n - 1$  (not  $n$ , due to the centering).

This is where a Singular Value Decomposition (SVD) comes in handy. Often people are sloppy and interchangeably refer to SVD and PCA, which is wrong. A SVD is a factorisation of an arbitrary (not necessarily square) matrix; for  $m \times n$  matrix  $\mathbf{A}$ , the SVD is

$$\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top$$

where

$\mathbf{U}$ : Eigenvectors of  $\mathbf{A}\mathbf{A}^\top$ ,

$\mathbf{V}$ : Eigenvectors of  $\mathbf{A}^\top\mathbf{A}$ , and

$\mathbf{\Sigma}$ : Diagonal matrix, common eigenvalues of  $\mathbf{A}\mathbf{A}^\top$  &  $\mathbf{A}^\top\mathbf{A}$  *squared*.

So, now consider the SVD of the centered data matrix  $\mathbf{X} - \bar{\mathbf{X}}$ ,

$$\mathbf{X} - \bar{\mathbf{X}} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top$$

Plug this into the sample variance:

$$\begin{aligned}\mathbf{S} &= \frac{1}{n}(\mathbf{X} - \bar{\mathbf{X}})(\mathbf{X} - \bar{\mathbf{X}})^\top \\ &= \frac{1}{n}(\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top)(\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top)^\top \\ &= \frac{1}{n}\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top\mathbf{V}\mathbf{\Sigma}\mathbf{U}^\top \\ &= \mathbf{U}\left(\frac{1}{n}\mathbf{\Sigma}^2\right)\mathbf{U}^\top.\end{aligned}$$

This shows that the the  $\mathbf{U}$  from the SVD of  $\mathbf{X} - \bar{\mathbf{X}}$  is exactly the eigenvectors of  $\mathbf{S}$ , and the eigenvalues of  $\mathbf{S}$  relate to the SVD as per

$$\mathbf{\Lambda}_{jj} = \mathbf{\Sigma}_{jj}^2/n.$$

For PCA, this means instead of computing  $\mathbf{S}$ , we can just submit  $\mathbf{X} - \bar{\mathbf{X}}$  to a SVD *as long* as we get the details right:

1. Compute centered data matrix,  $\mathbf{X} - \bar{\mathbf{X}}$ ;
2. Compute SVD of  $\mathbf{X} - \bar{\mathbf{X}}$ ; only  $\mathbf{U}$  and  $\mathbf{\Sigma}$  are needed; then
3.  $\mathbf{U}$  are the eigenvectors of  $\mathbf{S}$ , and
4.  $\text{diag}(\mathbf{\Sigma}^2)/n$  are the eigenvalues of  $\mathbf{S}$ ,  $\text{diag}(\mathbf{\Lambda})$

In matlab-ese, this means you can do either

```
[U, D] = eig(S);
```

or

```
[U, Dc] = svd(Xcenter); D=Dc.^2/n;
```

*Watch out!* The function `eig` sorts the eigenvalues from smallest to largest, while `svd` sorts them largest to smallest.