

Displays for Statistics 5401/8401

Lecture 24

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Class Web Page

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Principal components

Principal components are specific linear combinations $z_j \equiv \mathbf{v}_j' \mathbf{x} = \sum_{1 \leq \ell \leq p} v_{\ell j} x_\ell$ of variables x_1, \dots, x_p . The vectors $\mathbf{v}_j = [v_{1j}, \dots, v_{pj}]'$, $1 \leq j \leq p$, of coefficients are chosen to have certain properties.

There are at least two ways to motivate principal components.

- Principal components are linear combinations $\mathbf{v}_j' \mathbf{x}$ of variables which have the largest variances subject to constraints on the coefficients $v_{\ell j}$:

$$\|\mathbf{v}_j\|^2 = \mathbf{v}_j' \mathbf{v}_j = \sum_{\ell} v_{\ell j}^2 = 1 \text{ (normalized)}$$

$$\mathbf{v}_j' \mathbf{v}_k = \sum_{\ell} v_{\ell j} v_{\ell k} = 0, j \neq k \text{ (orthogonal)}$$

- Principal components are linear combinations from which you can approximately reconstruct a data matrix \mathbf{X} using a least squares criterion.

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The two approaches agree in an important way.

Both view the first few principal components as a set, preferably small, of *new* variables z_1, z_2, \dots, z_m , linearly related to x_1, x_2, \dots, x_p , and which lose as little as possible "important information" in the complete data.

- The variance maximization approach equates "important information" with high variability.
- The data matrix approximation approach equates "important information" with having an approximation with small errors.

I focus on their use in *approximating* a N by p data matrix \mathbf{X} , because I think that is closer to the way principal components are usually used.

In many cases, the single most important description or summary of a data matrix \mathbf{X} is its sample mean vector $\bar{\mathbf{x}} = \sum_i \mathbf{x}_i / N$.

I think this is because the N by p matrix

$$\mathbf{1}_N \bar{\mathbf{x}}' = \begin{bmatrix} \bar{\mathbf{x}}' \\ \bar{\mathbf{x}}' \\ \dots \\ \bar{\mathbf{x}}' \end{bmatrix}$$

often "explains" or predicts \mathbf{X} well in the following sense:

Elements of $\tilde{\mathbf{X}} \equiv \mathbf{X} - \mathbf{1}_N \bar{\mathbf{x}}'$ are often much smaller than the elements of \mathbf{X} .

Principal components are actually derived from a process which attempts to approximate $\tilde{\mathbf{X}}$ rather than \mathbf{X} . By adding $\mathbf{1}_N \bar{\mathbf{x}}'$ to an approximation $\hat{\tilde{\mathbf{X}}}$ to $\tilde{\mathbf{X}}$, you can then get an approximation $\hat{\mathbf{X}}$ for \mathbf{X} :

$$\hat{\mathbf{X}} = \hat{\tilde{\mathbf{X}}} + \mathbf{1}_N \bar{\mathbf{x}}'$$

The matrix of residuals of \mathbf{X} from $\bar{\mathbf{x}}$ is,

$$\tilde{\mathbf{X}} = \mathbf{X} - \mathbf{1}_N \bar{\mathbf{x}}' = \begin{bmatrix} (\mathbf{x}_1 - \bar{\mathbf{x}})' \\ (\mathbf{x}_2 - \bar{\mathbf{x}})' \\ (\mathbf{x}_3 - \bar{\mathbf{x}})' \\ \vdots \\ (\mathbf{x}_N - \bar{\mathbf{x}})' \end{bmatrix}, \text{ N by p}$$

When $N > p$, usually $\text{rank}(\tilde{\mathbf{X}}) = p$. That is, you can always find p pairs of vectors,

$$\mathbf{U}_k \text{ (N} \times \text{1) and } \mathbf{v}_k \text{ (p} \times \text{1)}$$

so that $\tilde{\mathbf{X}}$ is the sum of p *outer products* $\mathbf{U}_k \mathbf{v}_k'$ of \mathbf{U}_k and \mathbf{v}_k :

$$\tilde{\mathbf{X}} = \sum_{1 \leq k \leq p} \mathbf{U}_k \mathbf{v}_k'$$

The element in row i (case i) and column l (variable l) of $\tilde{\mathbf{X}}$ is

$$\tilde{x}_{il} \equiv x_{il} - \bar{x}_l = \sum_{1 \leq k \leq p} u_{ik} v_{lk} = \sum_{1 \leq k \leq p} v_{lk} u_{ik}$$

There are an infinite number of such sets of vectors $\{\mathbf{U}_k\}$ and $\{\mathbf{v}_k\}$.

That is, you *may* be able closely to *approximate* $\tilde{\mathbf{X}}$ by a sum of $m < p$ outer products:

$$\tilde{\mathbf{X}} \approx \sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k' \text{ or } \tilde{\mathbf{X}}_l = \sum_{1 \leq k \leq m} v_{lk} \mathbf{U}_k, l = 1, \dots, p$$

in the sense that the elements of $\tilde{\mathbf{X}} - \sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k'$ are small.

This would be a *reduced rank approximation* (specifically a rank m approximation) to $\tilde{\mathbf{X}}$ because

$$\text{rank}(\sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k') = m < p$$

How do you find such \mathbf{U}_i 's and \mathbf{v}_i 's?

That is, for a specific $m < p$, how do you find

- Variables \mathbf{U}_k , $k = 1, \dots, m$
- Coefficient vectors \mathbf{v}_k , $k = 1, \dots, m$

such that $\tilde{\mathbf{X}} - \sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k'$ is small?

In this representation, $\tilde{\mathbf{X}} = \sum_{1 \leq k \leq p} \mathbf{U}_k \mathbf{v}_k'$, think of \mathbf{U}_k and \mathbf{v}_k in the following way:

- Each N by 1 \mathbf{U}_k is a new *variable* with a value for each of the N cases
- Element v_{lk} of the p by 1 vector \mathbf{v}_k is a *coefficient* of \mathbf{U}_k , in a representation for column $\tilde{\mathbf{X}}_l$ of $\tilde{\mathbf{X}}$.

Specifically, if $\tilde{\mathbf{X}}_l$ is column l of $\tilde{\mathbf{X}}$,

$$\tilde{\mathbf{X}}_l = \sum_{1 \leq k \leq p} v_{lk} \mathbf{U}_k$$

Except for there being no constant term (intercept), this looks a little like a multiple regression of $\tilde{\mathbf{X}}_l$ on $\mathbf{U}_1, \dots, \mathbf{U}_p$.

When $\text{rank}(\tilde{\mathbf{X}}) = p$, when $m < p$ you can't exactly reproduce $\tilde{\mathbf{X}}$ by $\sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k'$, but it may be possible to get a good fit that is, have $\tilde{\mathbf{X}} - \sum_{1 \leq k \leq m} \mathbf{U}_k \mathbf{v}_k'$.

Let's start with $m = 1$, that is, find a $N \times 1$ \mathbf{U} and $p \times 1$ \mathbf{v} such that

$$\tilde{\mathbf{X}}^{(1)} = \mathbf{U}\mathbf{v}' = [v_1\mathbf{U} \ v_2\mathbf{U} \ \dots \ v_p\mathbf{U}] \approx \tilde{\mathbf{X}} \text{ (rank 1)}$$

that is, find numbers

- $\{u_i\}$, $i = 1, \dots, N$
- $\{v_l\}$, $l = 1, \dots, p$

so that $\tilde{x}_{il}^{(1)} - u_i v_l$ is small.

Let's drop the \approx and just use \mathbf{X} , since what we are doing does not depend on working with a matrix with 0 means, although that is the usual case.

If you view \mathbf{U} as a new variable, this is like finding one predictor variable \mathbf{U} so that the regressions (without intercept) of each column of \mathbf{X} on \mathbf{U} is a good fit.

Contrast this with the usual regression situation where you are *given* a predictor variable \mathbf{Z} and seek to find coefficients. Here you need to find both predictor and coefficients.

The first thought many statisticians would have would be to find \mathbf{U} and \mathbf{v} so that \mathbf{Uv}' is close to \mathbf{X} by the *least squares* criterion.

That is, find \mathbf{U} and \mathbf{v} so as to minimize

$$\sum_{1 \leq i \leq N} \sum_{1 \leq l \leq p} (x_{i,l} - u_i v_l)^2 = \|\mathbf{X} - \mathbf{Uv}'\|^2$$

That's what we're going to do.

Notation: When $\mathbf{A} = [a_{ij}]$ is a matrix

$$\|\mathbf{A}\|^2 = \sum_i \sum_j a_{ij}^2 = \text{trace}(\mathbf{A}'\mathbf{A})$$

The **Singular Value Decomposition** (SVD) of \mathbf{X} is the key to finding \mathbf{U} and \mathbf{v} to minimize $\|\mathbf{X} - \mathbf{Uv}'\|^2$.

The SVD of \mathbf{X} is a mathematical representation of \mathbf{X} which is useful in many contexts.

Vocabulary

$\mathbf{X} = \mathbf{LTR}'$ is the *singular value decomposition* (SVD) of \mathbf{X}

Facts

- $\mathbf{X} = \mathbf{LTR}' = \sum_{1 \leq k \leq p} t_k \mathbf{L}_k \mathbf{r}_k' = \sum_{1 \leq k \leq p} (t_k \mathbf{L}_k) \mathbf{r}_k'$, a sum of p *outer products* of $t_k \mathbf{L}_k$ and \mathbf{r}_k
- The t_i 's are unique
- When $t_i \neq t_j$, all $j \neq i$, \mathbf{L}_i and \mathbf{r}_i are unique (except for multiplication of both by -1 : $(-\mathbf{L}_i)(-\mathbf{r}_i)' = \mathbf{L}_i \mathbf{r}_i'$)

Thus the SVD $\mathbf{X} = \mathbf{LTR}'$ of \mathbf{X} is essentially unique.

- The N by 1 vectors \mathbf{L}_j , $j = 1, \dots, p$ are the **left singular vectors** of \mathbf{X} .
- The p by 1 vectors \mathbf{r}_j , $j = 1, \dots, p$ are the **right singular vectors** of \mathbf{X} .
- The p scalars $t_1 \geq t_2 \geq \dots \geq t_p \geq 0$ are the **singular values** of \mathbf{X} .

The Singular Value Decomposition

For *any* $N \times p$ matrix \mathbf{X} with $N \geq p$, there are always three matrices \mathbf{L} , \mathbf{R} and \mathbf{T} such that $\mathbf{X} = \mathbf{LTR}'$ where

- $\mathbf{L} = [\mathbf{L}_1, \mathbf{L}_2, \dots, \mathbf{L}_p]$ is $N \times p$ with $\mathbf{L}'\mathbf{L} = \mathbf{I}_p$. That is, the columns $\mathbf{L}_1, \dots, \mathbf{L}_p$ of \mathbf{L} are *orthonormal*:

$$\mathbf{L}_j' \mathbf{L}_j = 1, \mathbf{L}_j' \mathbf{L}_k = 0, j \neq k$$

Note: When $N > p$, this does not mean that $\mathbf{L}^{-1} = \mathbf{L}'$, since \mathbf{L} is not square.

- $\mathbf{R} = [\mathbf{r}_1, \mathbf{r}_2, \dots, \mathbf{r}_p]$ is $p \times p$ *square* with $\mathbf{R}'\mathbf{R} = \mathbf{I}_p$, that is, the columns $\mathbf{r}_1, \dots, \mathbf{r}_p$ of \mathbf{R} are *orthonormal*:

$$\mathbf{r}_j' \mathbf{r}_j = 1, \mathbf{r}_j' \mathbf{r}_k = 0, j \neq k$$

Since \mathbf{R} is $p \times p$, this means that $\mathbf{R}^{-1} = \mathbf{R}'$ and $\mathbf{R}\mathbf{R}' = \mathbf{I}_p$. \mathbf{R} is an orthogonal matrix.

- $\mathbf{T} = \text{diag}[t_1, \dots, t_p]$, $p \times p$, *diagonal* with $t_1 \geq t_2 \geq \dots \geq t_p \geq 0$

When there are only $s < p$ singular values $t_i \neq 0$, so that $t_{s+1} = \dots = t_p = 0$,

$$\mathbf{X} = \sum_{1 \leq k \leq s} (t_k \mathbf{L}_k) \mathbf{r}_k'$$

a sum of only s outer products.

Fact:

- $\text{Rank}(\mathbf{X}) = s =$ number of non-zero singular values. When $s < p$, $t_s > t_{s+1} = t_{s+2} = \dots = t_p = 0$.

The SVD is often the best way numerically to determine the $\text{Rank}(\mathbf{X})$:

- Compute \mathbf{T} from \mathbf{X}
- Count how many diagonal elements are non-zero except for rounding error. This should be $\text{Rank}(\mathbf{X})$.

Computing the SVD in MacAnova: `svd()`

Suppose x is a REAL matrix. Then `svd(x)` computes the vector $[t_1, \dots, t_p]'$ of singular values of x (`diag(T)`, *not* T)

`svd(x, left:T)` computes a structure with two components:

- `values`, vector of singular values
- `leftvectors` matrix L whose columns are L_1, \dots, L_p , the left singular vectors

`svd(x, right:T)` computes a structure with two components:

- `values`, vector of singular values
- `rightvectors` matrix R whose columns are r_1, \dots, r_p , right singular vectors

`svd(x, right:T, left:T)` OR `svd(x, all:T)` computes a 3 component structure:

- `values`: singular values
- `leftvectors`: left singular vectors
- `rightvectors`: right singular vectors

You can construct the diagonal matrix T in the SVD by

```
Cmd> tmatrix <- dmat(results$values); tmatrix#Diagonal matrix T
(1,1) 1415.4 0 0 0
(2,1) 0 27.14 0 0
(3,1) 0 0 2.2961 0
(4,1) 0 0 0 0.41587
```

Here are numerical checks that the right and left singular vectors are orthonormal ($R'R = I_p$ and $L'L = I_p$):

```
Cmd> R <- results$rightvectors # right singular vectors
```

```
Cmd> list(R) # size is p by p
R REAL 4 4 (labels)
```

```
Cmd> R' %*% R # = I_4
(1) (2) (3) (4)
(1) 1 4.0533e-17 -6.9218e-17 4.9043e-17
(2) 4.0533e-17 1 6.2095e-17 4.8526e-17
(3) -6.9218e-17 6.2095e-17 1 -1.4827e-16
(4) 4.9043e-17 4.8526e-17 -1.4827e-16 1
```

This is I_4 .

```
Cmd> L <- results$leftvectors # left singular vectors
```

```
Cmd> list(L) # size is N by p
L REAL 10 4 (labels)
```

```
Cmd> L' %*% L # = I_4
(1) (2) (3) (4)
(1) 1 -6.9389e-18 -1.1102e-16 -2.3592e-16
(2) -6.9389e-18 1 1.6653e-16 3.8858e-16
(3) -1.1102e-16 1.6653e-16 1 -4.4409e-16
(4) -2.3592e-16 3.8858e-16 -4.4409e-16 1
```

This is also I_4 .

Example

```
Cmd> x <- run(10)^run(0,3)' # powers of run(10)
Cmd> setlabels(x, \
  structure("@", vector("i^0", "i^1", "i^2", "i^3")))
Cmd> x # 10 ny 4 matrix
      i^0      i^1      i^2      i^3
(1) 1 1 1 1
(2) 1 2 4 8
(3) 1 3 9 27
(4) 1 4 16 64
(5) 1 5 25 125
(6) 1 6 36 216
(7) 1 7 49 343
(8) 1 8 64 512
(9) 1 9 81 729
(10) 1 10 100 1000

Cmd> vals <- svd(x); vals #just Sing values
(1) 1415.4 27.14 2.2961 0.41587
```

X has rank 4, but since the two smallest singular value are so small, it is close to having rank 3, or possibly even rank 2.

```
Cmd> results <- svd(x, left:T, right:T) # or all:T
```

```
Cmd> compnames(results)
(1) "values"           Structure
(2) "leftvectors"     component
(3) "rightvectors"    names
```

- `values` p -vector, (t_1, t_2, \dots, t_p)
- `leftvectors` N by p $L = [L_1 \dots L_p]$
- `rightvectors` p by p square $R = [r_1 \dots r_p]$

Relationship with Eigenvalues and Eigenvectors

- Each right singular vector r_j is an eigenvector of $X'X$ with eigenvalue t_j^2 .

That is, r_j satisfies $X'Xr_j = t_j^2 r_j$.

Check: $X'Xr_j = RTL'LTR'r_j = t_j^2 r_j$, because $L'L = I_p$ and $R'R = I_p$.

```
Cmd> sqrt(eigenvals(x' %*% x)) # numerical check
(1) 1415.4 27.14 2.2961 0.41587
```

$X'X$ is the $p \times p$ matrix of sums of squares (SS) and sums of products (SP) of the *columns* of X .

And, for the case we apply this to, $\tilde{X}'\tilde{X}$ consists of sums of squares $\sum_i (x_{i\ell} - \bar{x}_\ell)^2$ and products $\sum_i (x_{i\ell} - \bar{x}_\ell)(x_{ik} - \bar{x}_k)$. An, of course, $S_x = (1/(N-1))\tilde{X}'\tilde{X}$.

- Each left singular vector L_j is an eigenvector of the N by N matrix XX' with eigenvalue t_j^2 .

That is, L_j satisfies $XX'L_j = t_j^2 L_j$.

Check: $XX'L_j = LTR'RTL'L_j = t_j^2 L_j$ since $R'R = I_p$ and $L'L = I_p$.

```
Cmd> sqrt(round(eigenvals(x %*% x'),9)) # numerical check
(1)      1415.4      27.14      2.2961      0.41587      0
(6)         0         0         0         0         0
```

There are $N - p = 6$ zero eigenvalues. XX' is the $N \times N$ matrix of SS and SP of the rows of X .

Now define p linear combinations of the columns of X with coefficients from the right singular vectors r_j , $j = 1, \dots, p$:

$$U_j \equiv Z_j = Xr_j = LTR'r_j = t_j L_j, \quad j = 1, \dots, p$$

$Z_j = Xr_j = \sum_{1 \leq k \leq p} r_{kj} X_k$ is a linear combination of the columns of X .

The coefficients (weights) are the elements of

$$r_j = \begin{aligned} & j^{\text{th}} \text{ right singular vector of } X \\ & = j^{\text{th}} \text{ eigenvector of } XX'. \end{aligned}$$

Because $R'R = I_p$, $R'r_j = [0 \ 0 \ \dots \ 1 \ \dots \ 0]'$ and so

$$Z_j = LTR'r_j = t_j L_j$$

is proportional to a left singular vector of X .

Summary

If v_1, v_2, \dots, v_p are eigenvectors of XX' with eigenvalues $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_p \geq 0$ then

- the j^{th} *singular value* is $t_j = \sqrt{\lambda_j}$
- the j^{th} *right singular vector* is $r_j = v_j$ (could be $-v_j$), $j = 1, \dots, p$

If l_1, l_2, \dots, l_N (all N by 1) are eigenvectors of XX' with eigenvalues $\lambda_1 \geq \lambda_2 \geq \lambda_p \geq \dots \geq \lambda_N$ then

- the j^{th} *singular value* is $t_j = \sqrt{\lambda_j}$, $j = 1, \dots, p$
- the j^{th} *left singular vector* is $L_j = l_j$ (could be $-l_j$), $j = 1, \dots, p$
- The remaining eigenvalues $\lambda_{p+1}, \dots, \lambda_N$ are 0
- The remaining eigenvectors l_{p+1}, \dots, l_N are irrelevant

Back to low rank approximations

Q. With $m = 1$, what $N \times 1$ U and $p \times 1$ v minimize (make smallest) the "residual SS"

$$\|X - Uv'\|^2 = \sum_{1 \leq i \leq N} \sum_{1 \leq l \leq p} (x_{il} - u_i v_l)^2?$$

A. $U = Z_1 = t_1 L_1$ and $v = r_1$

That is

$$\hat{X}^{(1)} \equiv Z_1 r_1' = t_1 L_1 r_1' = X r_1 r_1'$$

is the best rank 1 approximation to X in the least squares sense.

This generalizes to rank $m > 1$:

$$\begin{aligned} \hat{X}^{(m)} &= \sum_{1 \leq j \leq m} Z_j r_j' = \sum_{1 \leq j \leq m} t_j L_j r_j' \\ &= X \left(\sum_{1 \leq j \leq m} r_j r_j' \right) \end{aligned}$$

is the best rank m approximation to X in the least squares sense.

How good is the approximation?

- The "residual sum of squares" is

$$\begin{aligned} \| \mathbf{X} - \hat{\mathbf{X}}^{(m)} \|^2 &= \sum_{1 \leq i \leq N} \sum_{1 \leq \ell \leq p} (x_{i\ell} - \hat{x}_{i\ell}^{(m)})^2 \\ &= \sum_{m+1 \leq k \leq p} t_k^2 = \sum_{m+1 \leq k \leq p} \lambda_k \end{aligned}$$

= the sum of the squared smallest $p - m$ singular values of \mathbf{X}

= sum of the smallest $p - m$ eigenvalues of $\mathbf{X}'\mathbf{X}$.

- The "total sum of squares" is

$$\| \mathbf{X} \|^2 = \sum_{\ell} \sum_i x_{i\ell}^2 = \sum_{1 \leq k \leq p} t_k^2 = \sum_{1 \leq k \leq p} \lambda_k = \text{tr } \mathbf{X}'\mathbf{X}$$

Therefore, when the ratio

$$\frac{\| \mathbf{X} - \hat{\mathbf{X}}^{(m)} \|^2}{\| \mathbf{X} \|^2} = \frac{\sum_{1 \leq i \leq N} \sum_{1 \leq \ell \leq p} (x_{i\ell} - \hat{x}_{i\ell}^{(m)})^2}{\sum_{1 \leq i \leq N} \sum_{1 \leq \ell \leq p} x_{i\ell}^2} = \frac{\sum_{m+1 \leq k \leq p} t_k^2}{\sum_{1 \leq k \leq p} t_k^2}$$

is small, the approximation is pretty good. This ratio is analogous to

$$SS_{\text{residual}} / SS_{\text{total}} = 1 - R^2$$

in regression.

In the rank m approximation,

$$\hat{\mathbf{X}}^{(m)} = \sum_{1 \leq k \leq m} \mathbf{z}_k \mathbf{r}_k' = \sum_{1 \leq k \leq m} t_k \mathbf{L}_k \mathbf{r}_k'$$

column \mathbf{X}_{ℓ} of \mathbf{X} is approximated by

$$\hat{\mathbf{X}}_{\ell}^{(m)} = \sum_{1 \leq k \leq m} \mathbf{z}_k r_{\ell k} = \sum_{1 \leq k \leq m} r_{\ell k} \mathbf{z}_k$$

a linear combination of $\mathbf{z}_1, \mathbf{z}_2, \dots, \mathbf{z}_m$.

Since the \mathbf{z}_k 's themselves are linear combinations of the columns of \mathbf{X} ($\mathbf{z}_k = \mathbf{X} \mathbf{r}_k$), so are the columns of $\hat{\mathbf{X}}^{(m)}$:

$$\begin{aligned} \hat{\mathbf{X}}_{\ell}^{(m)} &= \sum_{1 \leq k \leq m} r_{\ell k} \mathbf{X} \mathbf{r}_k \\ &= \sum_{1 \leq \ell \leq m} \left(\sum_{1 \leq k \leq m} r_{\ell k} r_{jk} \right) \mathbf{X}_j \end{aligned}$$

$\sum_{1 \leq k \leq m} r_{\ell k} r_{jk}$ is a partial sum of squares ($j = \ell$) or sum of products ($j \neq \ell$) of rows of \mathbf{R} . Since $\mathbf{R}\mathbf{R}' = \mathbf{I}_p$,

$$\sum_{1 \leq k \leq m} r_{\ell k}^2 = 1 - \sum_{m+1 \leq k \leq p} r_{\ell k}^2$$

$$\sum_{1 \leq k \leq m} r_{\ell k} r_{jk} = - \sum_{m+1 \leq k \leq p} r_{\ell k} r_{jk}$$