

Topic # 06: Data Analytics

Dimension Reduction, Principal Components

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Dimension Reduction

- The setting: we have a high-dimensional matrix of data \mathbf{X} . We would like to reduce this to a few **important** factors.
- We will do this by building a simple **linear** model for \mathbf{X} and use this model to represent \mathbf{X} in a **lower dimensional** space.
- Factor modeling is a **super useful** framework, whether you get a deep understanding or just learn how they work in practice.

Principal Component Analysis

- A **large** set of **correlated** variables
- **Principal Component** summarizes this set with a **smaller number of representative** variables
- **Principal Components** explain **most of the variability** in the original set
- **Principal Component Regression** use principal components as **predictors** in the regression model

What are Principal Components?

- Wish to **visualize** n observations with p features X_1, X_2, \dots, X_p
- Produce **two-dimensional** scatter plots: $p(p-1)/2$ plots
- Mostly likely, **none** of them will be **informative**
- We need a better method
- We need a **low-dimensional** representation that captures **as much of the information** as possible

Principal Components Analysis (PCA)

- PCA seeks a **small number** of dimensions that are as informative as possible
- The concept “informative” is measured by the amount of **variability**
- Each of the dimensions found by PCA is a linear combination of the p features
- The first **principal component** is the normalized linear combination of the features

$$Z_1 = \phi_{11}X_1 + \phi_{21}X_2 + \dots + \phi_{p1}X_p$$

that has the largest variance. By normalized, we mean

$$\sum_{j=1}^p \phi_{j1}^2 = 1.$$

Principal Components Analysis (PCA), Cont.

- We look for **linear combination** of the sample feature values of the form

$$z_{i1} = \phi_{11}x_{i1} + \phi_{21}x_{i2} + \dots + \phi_{p1}x_{ip}$$

that has the **largest sample variance**.

- We refer $z_{11}, z_{21}, \dots, z_{n1}$ as the **scores** of the first principal component.
- The PCA loading vector solves the **optimization problem**

$$\begin{aligned} \max_{\phi_{11}, \dots, \phi_{p1}} \quad & \left\{ \frac{1}{n} \sum_{i=1}^n \left(\sum_{j=1}^p \phi_{j1} x_{ij} \right)^2 \right\} \\ \text{s.t.,} \quad & \sum_{j=1}^p \phi_{j1}^2 = 1. \end{aligned}$$

Principal Components Analysis (PCA), Cont.

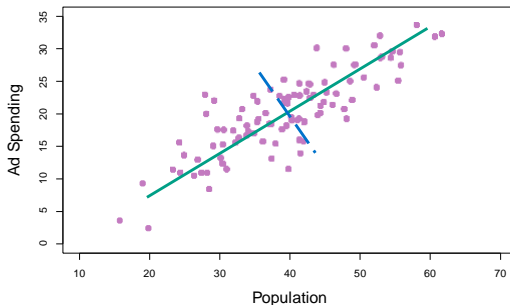
- We look for the **second principal component** Z_2

$$z_{i2} = \phi_{12}x_{i1} + \phi_{22}x_{i2} + \dots + \phi_{p2}x_{ip}$$

that has **largest sample variance** under constraint Z_2 is **uncorrelated** with Z_1 .

- The **uncorrelated** constraint is equivalent to constraining the direction ϕ_2 to be **orthogonal** to the direction ϕ_1 .
- Each principal component loading vector is **unique**.

Principal Components



- The population size (pop) and ad spending (ad) for 100 different cities are shown as purple circles.
- The green solid line indicates the first principal component
- The blue dashed line indicates the second principal component.

Proportion of Variance Explained

- Question: how much of the information in a given data set is **lost**?
- We consider the **proportion of variance explained** (PVE) by each principal component
- The total variance present in a data set (assuming variables have been centered to have mean zero)

$$\sum_{j=1}^p \text{Var}(X_j) = \sum_{j=1}^p \frac{1}{n} \sum_{i=1}^n x_{ij}^2$$

Proportion of Variance Explained, Cont.

- The variance explained by the m -th principal component is

$$\frac{1}{n} \sum_{i=1}^n z_{im}^2 = \frac{1}{n} \sum_{i=1}^n \left(\sum_{j=1}^p \phi_{jm} x_{ij} \right)^2$$

- Therefore, the PVE of the m -th principal component is given by

$$\frac{\sum_{i=1}^n \left(\sum_{j=1}^p \phi_{jm} x_{ij} \right)^2}{\sum_{j=1}^p \sum_{i=1}^n x_{ij}^2}$$

Proportion of Variance Explained, Cont.

- We want to use the **smallest number of principal components** to get a **good** understanding of the data.
- **How many** principal component are needed?
- **No single** answer!
- Produce a plot, choose the number of principal components in order to explain a **sizeable amount of variation** in the data.

- Data set USArrests

```
> states=row.names(USArrests)
> states
> names(USArrests)
> tail(USArrests)
> apply(USArrests,2,mean)
> apply(USArrests,2,var)
```

- Function **apply()**: arguments (dataset, row or column, function)

- Finding: variables are in **different scales**.

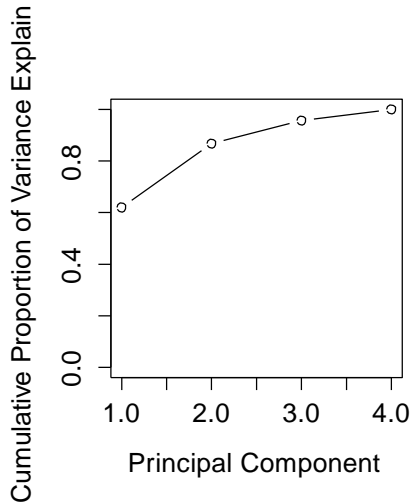
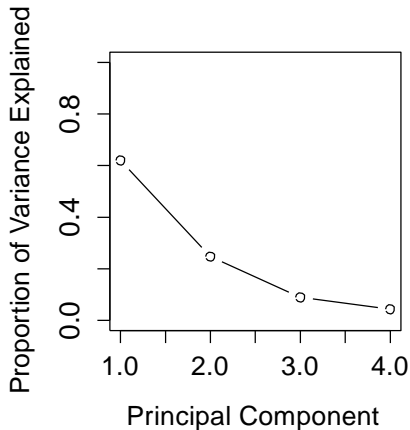
Principal Component Analysis in R

• Function `prcomp()`

```
> pr.out=prcomp(USArrests,scale=T)
> names(pr.out)
> pr.out$center
> pr.out$scale
> pr.out$rotation
> summary(pr.out)
> pr.var=pr.out$sdev^2
> pve=pr.var/sum(pr.var)
> plot(cumsum(pve))
```

• Finding: two principal variables explain **87% variability**.

Principal Components



- Play with other data sets: e.g., the Boston data set
- How much variability can be explained by the first and second principal variables?

Principal Components Regression

- A **dimension reduction** technique for **regression**
- The **Principal Components Regression** (PCR) approach involves constructing the first M principal components Z_1, \dots, Z_M , and then use these components as the predictors in a **linear regression**
- The key idea: a **small number** of principal components suffice to explain **most** of the variability in the data, as well as the **relationship** with the response.
 - PCA reduces dimension, which is always good.
 - Higher variance covariates are good in regression, and we choose the top PCs to have highest variance.
 - The PCs are independent: no multicollinearity

- **Test Error** rate and **Training error** rate can be very different.
- Hold out a subset of the data; apply statistical learning method to those hold-out data
- Randomly divide data into two parts
- Validation error rate is assessed using MSE

k-fold Cross-Validation

- k -fold CV randomly divide data into k folds of approximately equal size
- The first fold is treated as a validation set, and the method is fit on the remaining $k - 1$ folds.
- The mean squared error, MSE_1 , is computed in the hold-out fold.
- Repeat k times: each time, a different fold is treated as a validation set.
- The k -fold CV estimate is computed by averaging these values

$$CV_{(k)} = \frac{1}{k} \sum_{i=1}^k MSE_i.$$

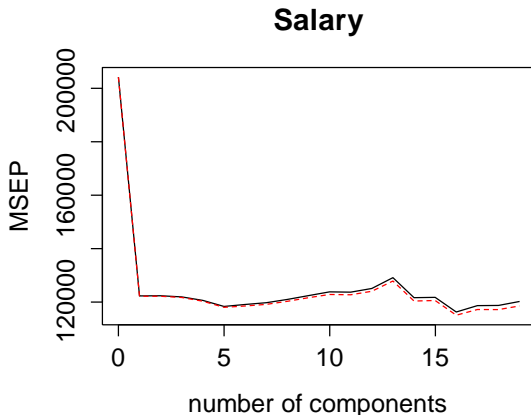
Principal Components Regression in R

- Principal Components Regression can be performed using function `pcr()` in library `pls`
- Determine number of principal components using eyes.
- In this data set, 7 may be a good size for principal components.
- In function `pcr()`, specify number of principal components by using `ncomp=`

```
> library(pls)
> library(ISLR)
> pcr.fit=pcr(Salary~.,data=Hitters,scale=T,validation="CV")
> summary(pcr.fit)

> pcr.fit2=pcr(Salary~.,data=Hitters,scale=T,validation="CV",ncomp=7)
> summary(pcr.fit2)
```
- Argument `validation="CV"` causes `pcr()` to compute the ten-fold cross-validation error for each possible M, the number of principal components used.

Principal Components Regression



Note that the smallest error occurs when $M = 16$, but it is roughly the same when **only one component** is included.

Principal Components Regression in R, cont.

- Perform `pcr()` on a training subset
- Make predictions on test subset
- Calculate prediction errors

```
set.seed(1)
Hitters=na.omit(Hitters)
x = model.matrix(Salary~., Hitters)[,-1]
y=Hitters$Salary
train = sample(1:nrow(x), nrow(x)/2)
test=(-train)
y.test=y[test]
```

```
pcr.fit = pcr(Salary~., data= Hitters, subset=train, scale=T,
validation = "CV")
pcr.pred = predict(pcr.fit, x[test,], ncomp=7)
```

```
mean((pcr.pred - y.test)^2)
[1] 96556.22
```

Questions, Comments?

See you next time.