

IE6600-Workshop

Visualizing with PCA

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1.Curse of dimensionality

True vs. Observed Dimensionality

What is the dimensionality of this dataset?

| ID | Observed Value (x ₁) - Height (cm) | Predictive Value (y) – Weight (kg) |
|----|---|---------------------------------------|
| 1 | 170 | 65 |
| 2 | 187 | 80 |
| 3 | 175 | 75 |
| 4 | 160 | 45 |
| 5 | 159 | 56 |



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Observed Dimensionality: 6



Features observed over time:

x₂: # of your waist (cm)
x₃: # of exams you have today
x₄: # of jokes you heard today
x₅: # of days left until summer

True vs. Observed Dimensionality

True Dimensionality: 3

| ID | Observed Value (x ₁) - Height (cm) | Observed value (x_2) - Waist (cm) | Predictive Value (y) – Weight (kg) |
|----|--|---|--|
| 1 | 170 | 67 | 65 |
| 2 | 187 | 86 | 80 |
| 3 | 175 | 76 | 75 |
| 4 | 160 | 59 | 45 |
| 5 | 159 | 66 | 56 |



Curse of Dimensionality

Example:

True Dimensionality

- 20x20 bitmap: {0, 1}⁴⁰⁰ potential events
- The handwritten number **THREE** may be only 2⁴⁰ events



1x400

Curse of Dimensionality

Example:

- 20x20 bitmap: {0, 1}⁴⁰⁰ potential events
- Randomly sampling 2400 events



Curse of Dimensionality

Data mining/Machine learning methods are statistical

Χ₁

- Use numbers to build up the predictor f(x)
- Use categorical variables to classify: {0, 1}

Dimensionality (d) increases, and fewer observations (n) per region In the case of d>>n

 X_2

- 1d: 3 regions
- 2d: 3^2 regions
- 100d well...



 X_3



2.Methods of dealing with high dimensionality

Methods *Dealing with high dimensionality*

Use domain knowledge

- Feature engineering
- e.g. Testosterone -> Muscle building

Assumption on dimensions

- Independence: count along each dimension, e.g. Naïve bayes.
 When counting the frequency of x₁ ignore the x₂
- Smoothness: nearby region should have similar distribution of classes
- Symmetry: e.g. invariance to order of the dimensions: order doesn't matter

Reduce the dimensionality

- Create a new set of variables



Dimensionality Reduction

Feature selection

- Select a subset of the original dimension: $x_1, x_2, x_3, x_4 x_5$
- Discriminative: Select class as predictor

Feature extraction

Construct a new set of dimensions:

$$E_i = f(x_1, x_2, x_3, x_4, x_5)$$



- e.g. Linear combination of original dimensions: PCA

3.Principal components analysis (PCA)

One of the most beautiful ALGORITHMs

Principal Components Analysis (PCA)

The goal of PCA is to find a new set of dimensions (attributes) that better captures the variability of the data.

Some Math

Standard Basis Vector

d-dimensional Cartesian coordinate space is specified via the d unit vectors, called the standard basis vectors, along each of the axes. The j-th standard basis vector e_j is the d-dimensional unit vector whose j-th component is 1 and the rest of the components are 0

$$e_j = \left(0, 1_j, \dots, 0\right)^T$$

Any other vector in \mathbb{R}^d can be written as linear combination of the standard basis vectors. For example, each of the points xi can be written as the linear combination d

$$x_i = x_{i1}e_1 + x_{i2}e_2 + \dots + x_{id}e_d = \sum_{j=1}^d x_{ij}e_j$$

where the scalar value x_{ij} is the coordinate value along the j-th axis or attribute

| (| $_{ m sepal}$ | $_{ m sepal}$ width | petal length | $\begin{array}{c} { m petal} \\ { m width} \end{array}$ | class | |
|-----------------------|---------------|---------------------|-----------------|---|-----------------|--|
| | X_1 | X_2 | X_2 | X_4 | X_5 | |
| x ₁ | 5.9 | 3.0 | 4.2 | 1.5 | Iris-versicolor | |
| \mathbf{x}_2 | 6.9 | 3.1 | 4.9 | 1.5 | Iris-versicolor | |
| \mathbf{x}_3 | 6.6 | 2.9 | 4.6 | 1.3 | Iris-versicolor | |
| \mathbf{x}_4 | 4.6 | 3.2 | 1.4 | 0.2 | Iris-setosa | |
| \mathbf{x}_5 | 6.0 | 2.2 | 4.0 | 1.0 | Iris-versicolor | |
| \mathbf{x}_6 | 4.7 | 3.2 | 1.3 | 0.2 | Iris-setosa | |
| \mathbf{x}_7 | 6.5 | 3.0 | 5.8 | 2.2 | Iris-virginica | |
| \mathbf{x}_8 | 5.8 | 2.7 | 5.1 | 1.9 | Iris-virginica | |
| 1 | 11.1 | ÷ | 4 | : | : | |
| x ₁₄₉ | 7.7 | 3.8 | 6.7 | 2.2 | Iris-virginica | |
| \mathbf{x}_{150} | 5.1 | 3.4 | 1.5 | 0.2 | Iris-setosa | |

Standard Basis Vector

For example:

Consider the Iris data

$$x_1 = (5.9, 3.0, 4.2)$$

$$x_1 = 5.9e_1 + 3.0e_2 + 4.2e_3 = 5.9\begin{pmatrix}1\\0\\0\end{pmatrix} + 3.0\begin{pmatrix}0\\1\\0\end{pmatrix} + 4.2\begin{pmatrix}0\\0\\1\end{pmatrix} = \begin{pmatrix}5.9\\3.0\\4.2\end{pmatrix}$$

Geometric View

For example:

 $x_1 = (5.9, 3.0)$ $x_1 \in \mathbb{R}^2$ $x_1 = (5.9, 3.0, 4.2)$ $x_1 \in \mathbb{R}^3$ X_3 3dvector.html X_2 4 3 $\mathbf{x}_1 = (5.9, 3.0, 4.2)$ $\mathbf{x}_1 = (5.9, 3.0)$ 3 2 2 1 1 X_2 6⁵4³ X_1 0 2 3 $\mathbf{5}$ 6 0 1 4 X_1

Let the data **D** consist of *n* points over *d* attributes, i.e., it is an $n \times d$ matrix, given as

| | 1 | X_1 | X_2 | $\mathcal{T}_{\mathcal{T}}^{(n)}$ | X_d |
|------------|----------------|------------------------|----------|-----------------------------------|----------|
| | \mathbf{x}_1 | <i>x</i> ₁₁ | x_{12} | ••• | x_{1d} |
| D = | \mathbf{x}_2 | x_{21} | x_{22} | ••• | x_{2d} |
| | ÷ | : | ÷ | ••• | |
| | \mathbf{x}_n | x_{n1} | x_{n2} | | x_{nd} |
| | | | | | |

Each point $\mathbf{x}_i = (x_{i1}, x_{i2}, \cdots, x_{id})^T$ is a vector in the ambient *d*-dimensional vector space spanned by the *d* standard basis vectors $\mathbf{e}_1, \mathbf{e}_2, \cdots, \mathbf{e}_d$, where \mathbf{e}_i corresponds to the *i*-th attribute X_i . Recall that the standard basis is an orthonormal basis for the data space, i.e., the basis vectors are pair-wise orthogonal, $\mathbf{e}_i^T \mathbf{e}_j = 0$, and have unit length $\|\mathbf{e}_i\| = 1$.



As such, given any other set of d orthonormal vectors $\mathbf{u}_1, \mathbf{u}_2, \cdots, \mathbf{u}_d$, with $\mathbf{u}_i^T \mathbf{u}_j = 0$ and $\|\mathbf{u}_i\| = 1$ (or $\mathbf{u}_i^T \mathbf{u}_i = 1$), we can re-express each point \mathbf{x} as the linear combination

$$x = a_1u_1 + a_2u_2 + \dots + a_du_d$$

where the vector $\mathbf{a} = (a_1, a_2, \dots, a_d)^T$ represents the coordinates of \mathbf{x} in the new basis. The above linear combination can also be expressed as a matrix multiplication

x = Ua

where U is the $d \times d$ matrix, whose *i*-th column comprises the *i*-th basis vector \mathbf{u}_i

$$U = \begin{pmatrix} | & | & | \\ u_1 u_2 \cdots u_d \\ | & | & | \end{pmatrix}$$

The matrix **U** is an *orthogonal* matrix, whose columns, the basis vectors, are *or*thonormal, i.e., they are pairwise orthogonal and have unit length

$$\mathbf{u}_i^T \mathbf{u}_j = \begin{cases} 1 & \text{if } i = j \\ 0 & \text{if } i \neq j \end{cases}$$

Since U is orthogonal, this means that its inverse equals its transpose

 $\mathbf{U}^{-1} = \mathbf{U}^T$

which implies that $\mathbf{U}^T \mathbf{U} = \mathbf{I}$, where \mathbf{I} is the $d \times d$ identity matrix.

$$x = Ua$$
$$U^T x = U^T Ua$$
$$a = U^T x$$

For example:





Original Basis

Optimal Basis

Mohammed J. Zaki, Wagner Meira, Jr., 2014

For example:

$$u_1 = \begin{pmatrix} -0.390\\ 0.089\\ -0.916 \end{pmatrix} \quad u_2 = \begin{pmatrix} -0.639\\ -0.742\\ 0.200 \end{pmatrix} \quad u_3 = \begin{pmatrix} -0.663\\ 0.664\\ 0.346 \end{pmatrix}$$

The new coordinates of the centered point $x = (-0.343, -0.754, 0.241)^T$ can be computed as:

$$a = U^{T}x = \begin{pmatrix} -0.390 & 0.089 & -0.916 \\ -0.639 & -0.742 & 0.200 \\ -0.663 & 0.664 & 0.346 \end{pmatrix} \begin{pmatrix} -0.390 \\ 0.089 \\ -0.916 \end{pmatrix} = \begin{pmatrix} -0.154 \\ 0.828 \\ -0.190 \end{pmatrix}$$

x can be written as the linear combination

 $x = -0.154u_1 + 0.828u_2 - 0.190u_3$



Mohammed J. Zaki, Wagner Meira, Jr., 2014

PCA

- 1. Find the first dimension to capture as much of the variability as possible
- 2. The second dimension is orthogonal to the first, and subject to that constraint, captures as much of the remaining variability.
- 3. And so on...until the dth dimension.



PCA Why find greatest variability?

Example:

We have a 2 dimensional data here to project all the data points to 1 dimension axis u_1 , u_2 : {x₁, x₂} -> u_1 , u_2

The data points in u_1 -space are more expanded (greater variability) than in u_2 -space.

- 1. Points are close in u_2 -space but far in (x₁, x₂)-space which is not so ideal to represent the original dataset
- 2. The overall distances in u_1 -space, with the highest variability, can represent the original distribution and variability



PCA Find mean and center the data

1. Center the data at zero: $Z = D - 1 \cdot \mu^T$

$$\mathbf{Z} = \mathbf{D} - \mathbf{1} \cdot \boldsymbol{\mu}^{T} = \begin{pmatrix} \mathbf{x}_{1}^{T} \\ \mathbf{x}_{2}^{T} \\ \vdots \\ \mathbf{x}_{n}^{T} \end{pmatrix} - \begin{pmatrix} \boldsymbol{\mu}^{T} \\ \boldsymbol{\mu}^{T} \\ \vdots \\ \boldsymbol{\mu}^{T} \end{pmatrix} = \begin{pmatrix} \mathbf{x}_{1}^{T} - \boldsymbol{\mu}^{T} \\ \mathbf{x}_{2}^{T} - \boldsymbol{\mu}^{T} \\ \vdots \\ \mathbf{x}_{n}^{T} - \boldsymbol{\mu}^{T} \end{pmatrix} = \begin{pmatrix} \mathbf{z}_{1}^{T} \\ \mathbf{z}_{2}^{T} \\ \vdots \\ \mathbf{z}_{n}^{T} \end{pmatrix}$$

Mohammed J. Zaki, Wagner Meira, Jr., Data Mining and Analysis: Fundamental Concepts and Algorithms, Cambridge University Press

PCA Direction

- 1. Center the data at zero: $Z = D 1 \cdot \mu^T$
- 2. Covariance matrix $\sum = \frac{1}{n} (Z^T Z)$
 - covariance of x_1, x_2 :
 - x_1, x_2 increase or decrease together or when one decreases the other one increases

Example



Towards the greatest variance direction

- 4. Look for a vector always keep in the same direction: $\sum u = \lambda u$
 - **u**: eigenvectors
 - Σ : covariance matrix
 - λ : scalar variable
 - Principal components = eigenvectors with largest eigenvalues



Towards the greatest variance direction

PCA Find eigenvalues, eigenvector, and PCs

1. Find eigenvalues by solving : $\sum u = \lambda u \rightarrow |\sum -\lambda I| = 0$. (*NOTE*: *Determinant of matrix A*: |A|)

$$-\begin{vmatrix} 2.0 - \lambda & 0.8 \\ 0.8 & 0.6 - \lambda \end{vmatrix} = (2 - \lambda)(0.6 - \lambda) - 0.8 * 0.8 = \lambda^2 - 2.6\lambda + 0.56 = 0$$

-
$$\{\lambda_1, \lambda_2\} = \frac{1}{2} (2.6 \pm \sqrt{2.6^2 - 4 * 0.56}) = \{2.36, 0.23\}$$

2. Find i^{th} eigenvector by solving: $\sum u_i = \lambda_i u_i$

$$\begin{array}{l} - \left(\begin{array}{cc} 2.0 & 0.8 \\ 0.8 & 0.6 \end{array} \right) \left(\begin{array}{c} u_{11} \\ u_{12} \end{array} \right) = 2.36 \left(\begin{array}{c} u_{11} \\ u_{12} \end{array} \right) \\ \Rightarrow \begin{array}{c} 2u_{11} + 0.8u_{12} = 2.36u_{11} \\ 0.8u_{11} + 0.6u_{12} = 2.36u_{12} \end{array} \Rightarrow u_{11} = 2.2u_{12} \Rightarrow u_{1} \sim \left(\begin{array}{c} 2.2 \\ 1 \end{array} \right), \\ \Rightarrow make ||u_{1}|| = 1, u_{1} \frac{1}{\sqrt{2.2^{2}+1}}, then u_{1} = \left(\begin{array}{c} 0.91 \\ 0.41 \end{array} \right), slope = 0.454 \\ - \left(\begin{array}{c} 2.0 & 0.8 \\ 0.8 & 0.6 \end{array} \right) \left(\begin{array}{c} u_{21} \\ u_{22} \end{array} \right) = 0.23 \left(\begin{array}{c} u_{21} \\ u_{22} \end{array} \right) \Rightarrow then u_{2} = \left(\begin{array}{c} -0.41 \\ 0.91 \end{array} \right) \\ 3.1^{st} PC: \left(\begin{array}{c} 0.91 \\ 0.41 \end{array} \right), 2^{nd} PC: \left(\begin{array}{c} -0.41 \\ 0.91 \end{array} \right) \end{array} \right) \end{array}$$

PCA Fraction of total variance, and choose dimensionality

Often we may not know how many dimensions, r, to use for a good approximation. One criteria for choosing r is to compute the fraction of the total variance captured by the first r principal components, computed as

$$f(r) = \frac{\lambda_1 + \lambda_2 + \dots + \lambda_r}{\lambda_1 + \lambda_2 + \dots + \lambda_d} = \frac{\sum_{i=1}^r \lambda_i}{\sum_{i=1}^d \lambda_i} = \frac{\sum_{i=1}^r \lambda_i}{var(D)}$$

Given a certain desired variance threshold, say α , starting from the first principal component, we keep on adding additional components, and stop at the smallest value r, for which $f(r) \ge \alpha$ (α can be 0.9, 0.95 as purposes).

In practice, α is usually set to 0.9 or higher, so that the reduced dataset captures at least 90% of the total variance.

PCA (\mathbf{D}, α) : $\mu = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i / / \text{ compute mean}$ $\mathbf{Z} = \mathbf{D} - \mathbf{1} \cdot \boldsymbol{\mu}^T$ // center the data $\Sigma = \frac{1}{n} (\mathbf{Z}^T \mathbf{Z}) / /$ compute covariance matrix $(\lambda_1, \lambda_2, \dots, \lambda_d) = \text{eigenvalues}(\Sigma) // \text{compute eigenvalues}$ $\mathbf{U} = (\mathbf{u}_1 \ \mathbf{u}_2 \ \cdots \ \mathbf{u}_d) = \text{eigenvectors}(\mathbf{\Sigma}) // \text{ compute eigenvectors}$ $f(r) = \frac{\sum_{i=1}^{r} \lambda_i}{\sum_{i=1}^{d} \lambda_i}$, for all $r = 1, 2, \cdots, d$ // fraction of total variance 7 Choose smallest r so that $f(r) \ge \alpha //$ choose dimensionality $\mathbf{s} \ \mathbf{U}_r = egin{pmatrix} \mathbf{u}_1 & \mathbf{u}_2 & \cdots & \mathbf{u}_r \end{pmatrix} // \ ext{reduced basis} \end{pmatrix}$ $\mathbf{A} = \{\mathbf{a}_i \mid \mathbf{a}_i = \mathbf{U}_r^T \mathbf{x}_i, \text{ for } i = 1, \cdots, n\}$ // reduced dimensionality data

Mohammed J. Zaki, Wagner Meira, Jr., Data Mining and Analysis: Fundamental Concepts and Algorithms

PCA





Novembre, John et al. "Genes mirror geography within Europe." Nature vol. 456,7218 (2008): 98-101. doi:10.1038/nature07331 IE6600 Visualization and Computation for Analytics, NEU | ©2022 Zhenyuan Lu

$$\sum = \begin{pmatrix} 0.681 & -0.039 & 1.265 \\ -0.039 & 0.187 & -0.320 \\ 1.265 & -0.32 & 3.092 \end{pmatrix}$$

$$\lambda_1 = 3.662$$
 $\lambda_2 = 0.239$ $\lambda_3 = 0.059$

Question

$$u_1 = \begin{pmatrix} -0.39\\ 0.089\\ -0.916 \end{pmatrix} \qquad u_2 = \begin{pmatrix} -0.639\\ -0.742\\ 0.200 \end{pmatrix} \qquad u_3 = \begin{pmatrix} -0.663\\ 0.664\\ 0.346 \end{pmatrix}$$

What is the total variance? What is the fraction of total variance for each PC? If let $\alpha = 0.95$ how many PCs we need to keep?

4.PCA Implementation in R

PCA (\mathbf{D}, α) : $\mu = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$ // compute mean $\mathbf{Z} = \mathbf{D} - \mathbf{1} \cdot \mu^T$ // center the data $\mathbf{\Sigma} = \frac{1}{n} (\mathbf{Z}^T \mathbf{Z})$ // compute covariance matrix $(\lambda_1, \lambda_2, \dots, \lambda_d) = \text{eigenvalues}(\mathbf{\Sigma})$ // compute eigenvalues $\mathbf{U} = (\mathbf{u}_1 \quad \mathbf{u}_2 \quad \dots \quad \mathbf{u}_d) = \text{eigenvectors}(\mathbf{\Sigma})$ // compute eigenvectors $f(r) = \frac{\sum_{i=1}^{r} \lambda_i}{\sum_{i=1}^{d} \lambda_i}$, for all $r = 1, 2, \dots, d$ // fraction of total variance 7 Choose smallest r so that $f(r) \ge \alpha$ // choose dimensionality $\mathbf{U}_r = (\mathbf{u}_1 \quad \mathbf{u}_2 \quad \dots \quad \mathbf{u}_r)$ // reduced basis $\mathbf{A} = \{\mathbf{a}_i \mid \mathbf{a}_i = \mathbf{U}_r^T \mathbf{x}_i, \text{ for } i = 1, \dots, n\}$ // reduced dimensionality data

1 data("iris")

```
2 head(iris)
```

```
1 irisCent <- iris %>% transmute(sL=Sepal.Length-mean(Sepal.Length),
2 sW=Sepal.Width-mean(Sepal.Width),
3 pL=Petal.Length-mean(Petal.Length),
4 pW=Petal.Width-mean(Petal.Width))
```

```
1 ic <- cov(irisCent)</pre>
```

- 1 # use eigen() function to compute eigenvalues and eigenvectors
- 2 ieigen <- eigen(ic)
- 3 ie <- ieigen\$values
- 4 iv <- ieigen\$vectors
- 5 row.names(iv) <- names(iris %>% select(-Species))
- 6 colnames(iv) <- paste0(rep("PC",ncol(iv)),1:ncol(iv)) # or sprintf("PC%d",1:4)

```
# Fraction of the total variance
```

2 fr <- ie/sum(ie)</pre>

```
# Choose number of dimensionality
      threshold <- function(x, th) {</pre>
        sum <- 0
        seq <- 0
       for (i in 1:length(x)) {
 6
          sum <- sum + x[i]</pre>
 7
          if (sum >= th) {
 8
            seq <- i
 9
            break
10
11
12
        return(seq)
13
14
15
16
      threshold(x=fr, 0.95)
```

PCA (\mathbf{D}, α) : $\mu = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i //$ compute mean $\mathbf{Z} = \mathbf{D} - \mathbf{1} \cdot \mu^T //$ center the data $\mathbf{\Sigma} = \frac{1}{n} (\mathbf{Z}^T \mathbf{Z}) //$ compute covariance matrix $(\lambda_1, \lambda_2, \dots, \lambda_d) = \text{eigenvalues}(\mathbf{\Sigma}) //$ compute eigenvalues $\mathbf{U} = (\mathbf{u}_1 \quad \mathbf{u}_2 \quad \dots \quad \mathbf{u}_d) = \text{eigenvectors}(\mathbf{\Sigma}) //$ compute eigenvectors $f(r) = \frac{\sum_{i=1}^{r} \lambda_i}{\sum_{i=1}^{d} \lambda_i}$, for all $r = 1, 2, \dots, d //$ fraction of total variance 7 Choose smallest r so that $f(r) \ge \alpha //$ choose dimensionality $\mathbf{U}_r = (\mathbf{u}_1 \quad \mathbf{u}_2 \quad \dots \quad \mathbf{u}_r) //$ reduced basis $\mathbf{A} = \{\mathbf{a}_i \mid \mathbf{a}_i = \mathbf{U}_r^T \mathbf{x}_i, \text{ for } i = 1, \dots, n\} //$ reduced dimensionality data

- 1 biPCA <- prcomp(iris[1:4], scale = TRUE)</pre>
- 2 biPCA\$sdev^2/sum(biPCA\$sdev^2)
- 3 biPCA\$rotation

```
barplot(
 1
 2
       fr,
 3
       ylim = c(0, 1),
 4
       col = "sandybrown",
       xlab = "Principal Component",
 5
       ylab = "Explained Variances",
 6
 7
       axes = TRUE
 8
     axis(1, c(0.7, 1.9, 3.1, 4.3),
 9
10
          labels = sprintf("PC%d", 1:4))
     lines(cumsum(fr), type = 's', col = "darkgreen")
11
12
     legend(
13
       x = 2.5,
14
       y = 0.5,
       legend = c("Explained Variance", "Cumulative
15
     Explained Variance"),
16
17
       pch = c(15, 15),
18
       col = c("sandybrown", "darkgreen"),
19
       bty = 'n'
```



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Exercise:

- Try three datasets: mpg, BostonHousing (mlbench), BreastCancer (mlbench) on two scale methods: a) mean=0, b)mean=0, variance=1.
- 2. Compare to the built-in PCA function prcomp()

PCA VS Linear Discriminant Analysis



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5.PCA: Pros and Cons

PCA *Pros and Cons*

Pros

- 1. Good performance on processing speed
- 2. Reflects intuition on the data
- 3. Efficient reduction in size of data

Cons

- 1. Doesn't consider class separability since it doesn't take into account the class labels
- 2. PCA simply performs a coordinate rotation that aligns the transformed axes with the directions of maximum variance
- 3. There is no guarantee that the directions of maximum variance will contain good features for discrimination
 - PCA cannot recognize the class lables

Resources

Resource

Textbook:

Galit Shmueli, Peter C. Bruce, Inbal Yahav, Nitin R. Patel, Kenneth C. Lichtendahl Jr., Data Mining for Business <u>Analytics:</u> Concepts, Techniques, and Applications in R **(DMBA)**, Wiley, 1st Edition, ISBN-10: 1118879368, ISBN-13: 978-1118879368.

Additional Textbooks:

R For Data Science (open license, R4DS), Wickham, Hadley, and Garrett Grolemund

R Markdown (<u>open license</u>, **RMD**), Xie, Yihui, et al.

James, Gareth, et al. An Introduction to Statistical Learning: with Applications in R. Springer, 2017. (open license, ISL)

Mohammed J. Zaki, Wagner Meira, Jr., Data Mining and Analysis: Fundamental Concepts and Algorithms, Cambridge University Press, May 2014. ISBN: 9780521766333.

David Hand, Heikki Mannila, Padhraic Smyth. Principles of Data Mining, The MIT Press, 2001, ISBN-10: 026208290X, ISBN-13: 978-0262082907.

Tan, Pang-Ning, et al. Introduction to Data Mining (DM). Pearson Education, 2006.

Materials

@Victor Lavrenko

Lu, Z. (2022). Data Visualization Tutorial in R. zhenyuanlu.github.io.