INTRODUCTION TO



FOR



USING



TUE VU

ADVANCED COMPUTING & DATA SCIENCE (ACDS)

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OUTLINES

- 1. Introduction to Machine Learning
- 2. Why R
- 3. Types of Machine Learning
- 4. Caret package
- 5. Supervised Learning
- 6. Unsupervised Learning







5. Supervised Learning 1. Regression

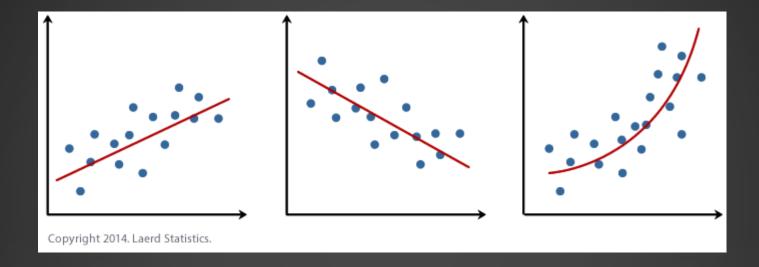
- - Linear Regression
 - Multi-Linear Regression (MLR)
 - Other typical Linear Regression Technique
 - Logistic Regression
- 2. Decision Tree
- 3. Ensemble Prediction
 - Random Forest
 - Bagging
 - Boosting
- 4. Model based Prediction
 - Naïve Bayes
 - Linear Discriminant Analysis
- 5. Regularization & Variable selection
 - Ridge Regression
 - LASSO
 - ELASTIC-NET
- 6. Dimension Reduction
 - Principal Component Analysis
- 7. Neural Network
- 8. Support Vector Machine
- 9. K-Nearest Neighbor





5.1. Regression based method

5.1.1. Linear Regression



$$y = ax + b$$

5.1. Regression based method

5.1.2. Multi-Linear Regression – Ordinary Least Square Regression

$$y = a_1 x_1 + a_2 x_2 + \dots + a_n x_n + b$$



- 5.1. Regression based method
- 5.1.3. Other typical Linear Regression Technique

Stepwise Linear Regression:

It's a step by step Regression to determine which covariates set best match with the dependent variable. Using AIC as criteria:

method = 'ImStepAIC'

Principal Component Regression:

Linear Regression using the output of a Principal Component Analysis (PCA). PCR is skillful when data has lots of highly correlated predictors method = 'pcr'



5.1. Regression based method

5.1.4. Logistic Regression

Logistic regression is another technique borrowed by machine learning from the field of statistics. It is the goto method for binary classification problems (problems with two class values).

Typical binary classification: True/False, Yes/No, Pass/Fail, Spam/No Spam, Male/Female



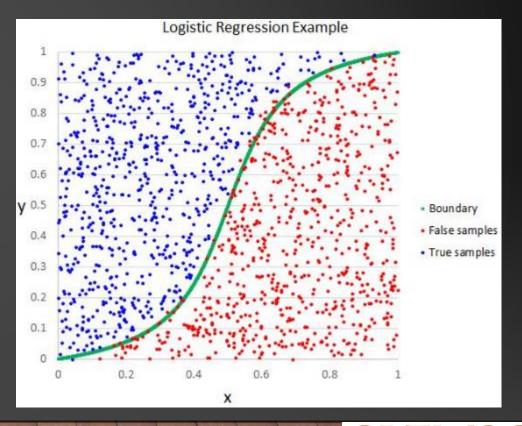
5.1. Regression based method

5.1.4. Logistic Regression

Unlike linear regression, the prediction for the output is transformed using a non-linear function called the logistic function.

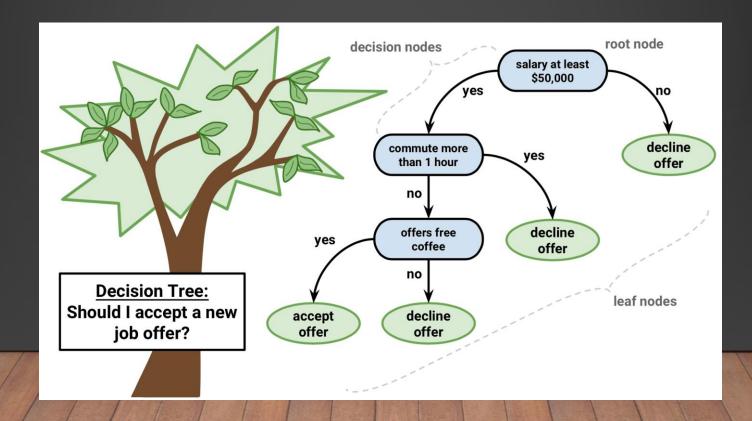
The standard logistic function has formulation:

$$f(x) = \frac{1}{1 + e^{-x}}$$



5.2. Decision Tree

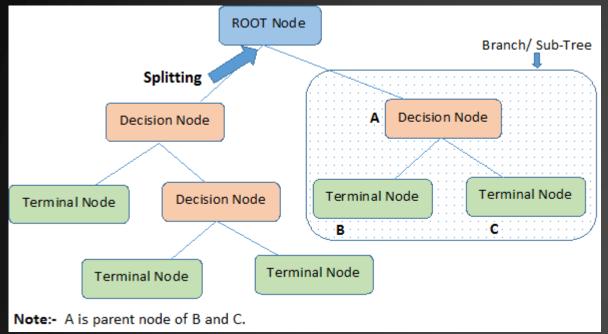
- Tree based learning algorithms are considered to be one of the best and mostly used supervised learning methods.
- Tree based methods empower predictive models with high accuracy, stability and ease of interpretation
- Non-parametric and non-linear relationships
- Types: Categorical and Continuous

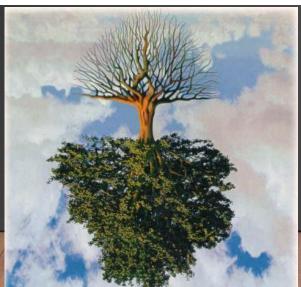






5.2. Decision Tree





Terminology

- 1. Root Node: It represents entire population or sample and this further gets divided into two or more homogeneous sets.
- 2. Splitting: It is a process of dividing a node into two or more sub-nodes.
- 3. Decision Node: When a sub-node splits into further sub-nodes, then it is called decision node.
- 4. Leaf/ Terminal Node: Nodes with no children (no further split) is called Leaf or Terminal node.
- 5. Pruning: When we reduce the size of decision trees by removing nodes (opposite of Splitting), the process is called pruning.
- 6. Branch / Sub-Tree: A sub section of decision tree is called branch or sub-tree.
- 7. Parent and Child Node: A node, which is divided into sub-nodes is called parent node of sub-nodes where as sub-nodes are the child of parent node





5.2. Decision Tree

Types

- Categorical Regression: "Yes/No", "TRUE/FALSE", "Male/Female", "Colors"
- Continuous Regression: when the dependent variable is continuous



5.2. Decision Tree

Splitting algorithm

- Gini Impurity: (Categorical)
- Chi-Square index (Categorical)
- Cross-Entropy & Information gain (Categorical)
- Reduction Variance (Continuous)

Different packages/methods use different splitting algorithm

- CART (Classification And Regression Tree) use Gini index & Entropy
- ID3 & C5.0 Tree use Entropy & Information gain
- CHAID use Chi-Square





5.2. Decision Tree

Gini impurity example: (Categorical)

$$Gini = 1 - \sum_{i} p(i)^2$$

A sample of 30 students with three variables Gender (Boy/ Girl), Class (IX/ X) and Height (5 to 6 ft).

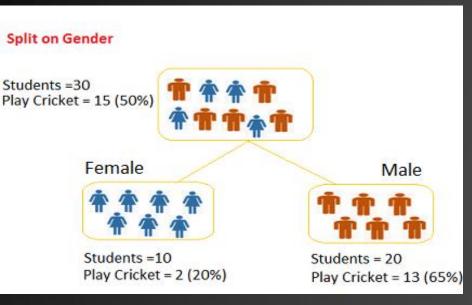
- 15 out of these 30 play cricket in leisure time.
- predict who will play cricket during leisure period?

In this problem, we need to segregate students who play cricket in their leisure time based on highly significant input variable among all three.



5.2. Decision Tree

Gini impurity example: (Categorical)



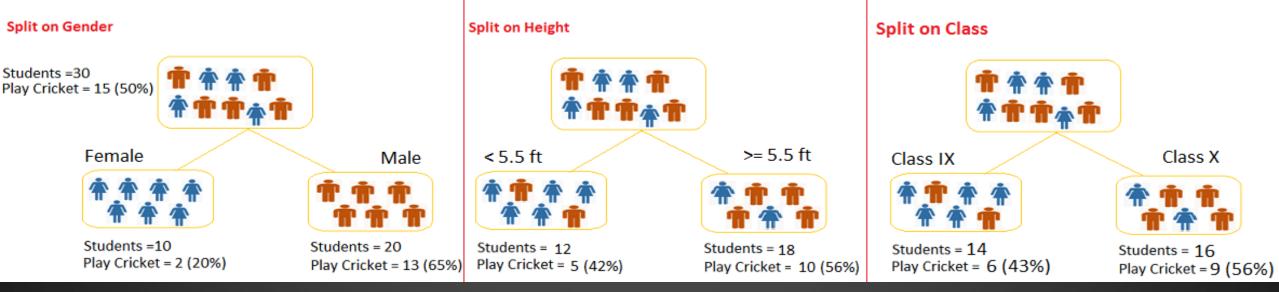
$$G_1 = 1 - 0.2^2 - (1 - 0.2)^2 = 0.32$$
 $G_2 = 1 - 0.65^2 - (1 - 0.65)^2 = 0.45$

GG=10/30*.32+20/30*.45=0.4



5.2. Decision Tree

Gini impurity example: (Categorical) – The smaller the Gini the better



 $G_A = 1 - 0.56^2 - (1 - 0.56)^2 = 0.50$ $G_5=1-0.43^2-(1-0.43)^2=0.49$

GG=10/30*.32+20/30*.45=**0.4**

GH = 12/30*.5+18/30*.5=0.5

 $G_A=1-0.56^2-(1-0.56)^2=0.5$

 $GC = 14/30^*.49 + 16/30^*.5 = 0.49$

Gini=0 when all observation belongs to one label.

 $G_2=1-0.65^2-(1-0.65)^2=0.45$ $G_3=1-0.42^2-(1-0.42)^2=0.5$



 $G_1=1-0.2^2-(1-0.2)^2=0.32$



5.2. Decision Tree

Chi-Square example: (Categorical)

Split by Gender

| | | Not Play | | Expected | Expected Not | Chi-Square | |
|--------|--------------|----------|-------|--------------|---------------|------------|----------|
| Node | Play Cricket | Cricket | Total | Play Cricket | Play Cricket | Play | Not Play |
| | | | | , | ,, | Cricket | Cricket |
| Female | 2 | 8 | 10 | 5 | 5 | 1.34 | 1.34 |
| Male | 13 | 7 | 20 | 10 | 10 | 0.95 | 0.95 |
| | | | | Tot | al Chi-Square | 4. | 58 |

 $\chi^2 = \sqrt{\frac{(Actual - Expected)^2}{Expected}}$

Gender is more significant than Class

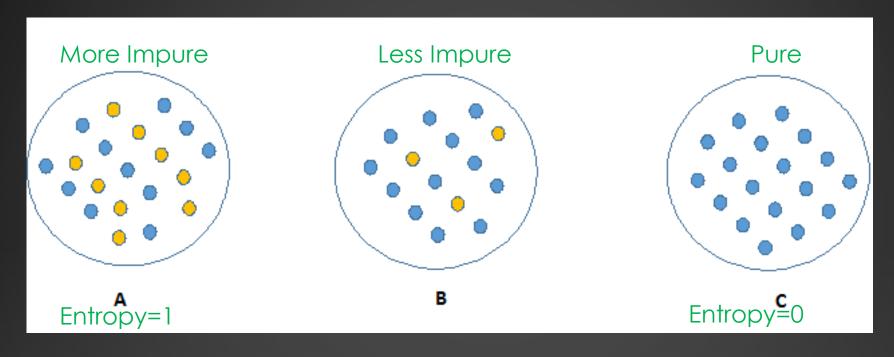
Split by Class

| | | Not Play | | Expected | Expected Not | Chi-Square | |
|------|--------------|----------|---|----------|----------------|------------|----------|
| Node | Play Cricket | Cricket | Total | | Play Cricket | Play | Not Play |
| | 1 | | <u> </u> | | , | Cricket | Cricket |
| IX | 6 | 8 | 14 | 7 | 7 | 0.38 | 0.38 |
| X | 9 | 7 | 16 | 8 | 8 | 0.35 | 0.35 |
| | | | | Tof | tal Chi-Square | 1.4 | 46 |



5.2. Decision Tree

Information Gain: Specify by Entropy



$$Entropy = -p\log_2 p - q\log_2 q$$

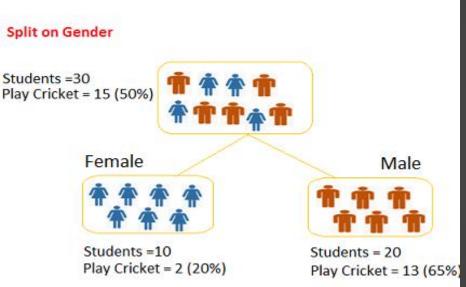


5.2. Decision Tree

Cross-Entropy example: (Categorical) – the smaller the Entropy the better

$$Entropy = -p\log_2 p - q\log_2 q$$

 $Information \ Gain = 1 - Entropy$



$$E_1 = -0.2\log_2 0.2 - 0.8\log_2 0.8 = 0.72$$

$$E_2 = -0.65\log_2 0.65 - 0.35\log_2 0.35 = 0.93$$

$$E_{\rm G} = \frac{10}{30} * 0.72 + \frac{20}{30} * 0.93 = 0.86$$

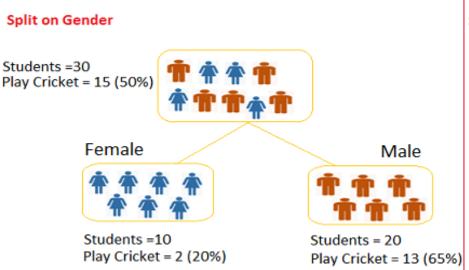


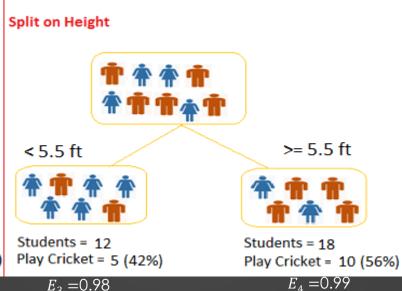
5.2. Decision Tree

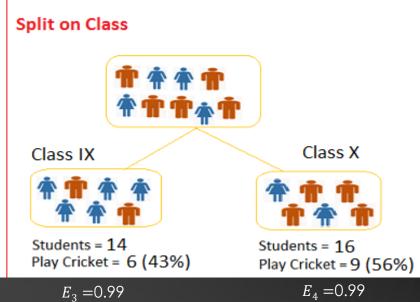
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$$E_2 = -0.65\log_2 0.65 - 0.35\log_2 0.35 = 0.93$$

$$E_{\rm G} = \frac{10}{30} * 0.72 + \frac{20}{30} * 0.93 = 0.86$$

$$E_{\rm C} = 0.98$$

 $E_{\rm C} = 0.99$

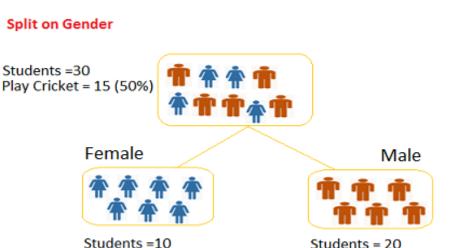


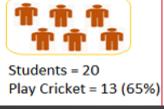


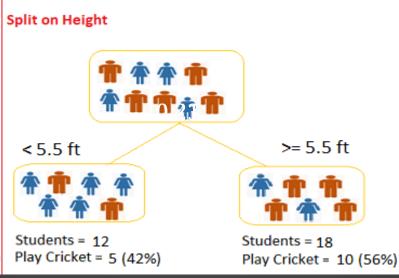
5.2. Decision Tree

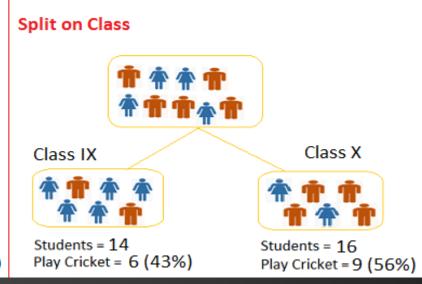
Reduction-Variance example: (Continuous) – the smaller the Variance the better

$$Variance = \frac{\sum (X - \bar{X})^2}{n}$$









$$\overline{V}_1 = \frac{(2*1+8*0)}{10} = 0.2$$

$$V_1 = \frac{2*(1-0.2)^2 + 8*(0-0.2)^2}{10} = 0.16$$

$$V_2 = 0.23$$

$$VG = \frac{10}{30} * 0.16 + \frac{20}{30} * 0.23 = 0.21$$

Play Cricket = 2 (20%)

$$VC = \frac{14}{30} * 0.24 + \frac{16}{30} * 0.25 = 0.25$$





| S. No. | Factors | Variables | Algorit hms | Methods | Splitting Criteria | Type of Tree | Missing Value | Noisy Data | Analysis | Performance | Pruning Support |
|-----------|---------------------------|---|-----------------|--|---|-----------------------------------|--------------------------------------|----------------------------------|--|--------------------------|---|
| 1 | Entropy | Continuous Discrete Categorical | ID3 and C4.5 | Exploratory Analysis (Impurity Measure) | Grouping of Classes that include up to 50% of the Data | | | | | Little slower to compute | |
| 2 | Information Gain | Categorical , discrete or Nominal | ID3 and C4.5 | Use Entropy for Analysis | Split dataset into large number of partitions | Consider Binary Split | Can't handle missing values | Can't handle noisy Data | Biased towards multivalued attributes | | Cannot handle Over Fitting |
| 3 | Information Gain Ratio | Numerical Continuous Discrete and Nominal | C4.5 | Use Information Gain and split information value or Intrinsic Information for Analysis | Prefer unbalanced splits in which one partition is much smaller than the other | Consider Multivalu ed Split | Can handle Missing Values | Can handle noisy Data | Reduce Biased towards multivalued attributes | | Reduce Over Fitting by using Pruning Methods |
| 4 | Gini Index | Discrete Numerical continuous | CART | Use Weighted average of each branch index (Impurity Measure) | Grouping of largest class from other classes (Twoing Criteria) | Consider Binary Split | Can handle Missing Values | Can handle noisy Data | Biased towards multivalued attributes | Little Faster to compute | Reduce Over Fitting by using Pruning Methods |





5.2. Decision Tree

| Pros | Cons |
|---|--|
| Work with categorical/continuous Easy to implement & explain Easy to display graphically No need to clean data | Low Predictive accuracyHigh chance of overfitting |



5.2. Decision Tree

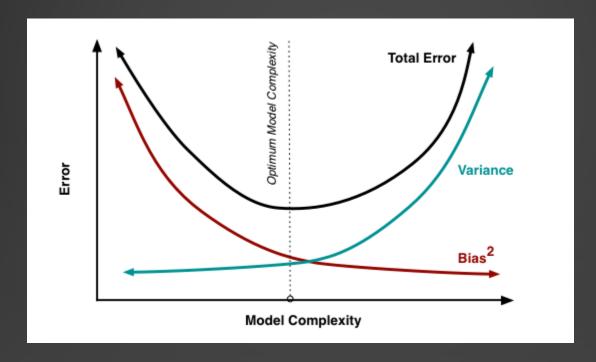
| Pros | Cons |
|---|--|
| Work with categorical/continuous Easy to implement & explain Easy to display graphically No need to clean data | Low Predictive accuracyHigh chance of overfitting |

By aggregating many Decision Trees using Ensemble methods, the Predictive Performance can be substantially improved



5.3. Ensemble

- Ensemble is to "group" many predictive model to achieve a better accuracy and model stability
- How to balance between Bias & Variance to get optimal Error?
- Trade-off management
- Solved by Ensemble method





5.3. Ensemble

5.3.1. Random Forest

Random Forest is considered to be a **panacea** of all data science problems. On a funny note, when you can't think of any algorithm (irrespective of situation), use random forest!

Opposite to Decision Tree, Random Forest use bootstrapping technique to grow multiple tree

Random Forest is a versatile machine learning method capable of performing both regression and classification tasks.

It is a type of ensemble learning method, where a group of weak models combine to form a powerful model.

The end output of the model is like a black box and hence should be used judiciously.



5.3. Ensemble

- 5.3.1. Random Forest
 - 1. If there are M input variables, a number m<M is specified such that at each node, m variables are selected at random out of the M. The best split on these m is used to split the node. The value of m is held constant while we grow the forest.
 - 2. Each tree is grown to the largest extent possible and there is no pruning.

3. Predict new data by aggregating the predictions of the ntree trees (i.e., majority votes for

classification, average for regression).





5.3. Ensemble

5.3.1. Random Forest

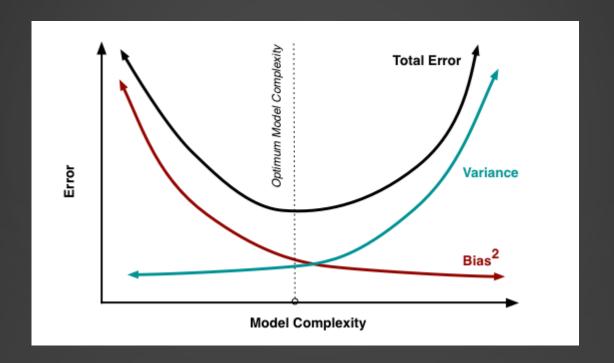
| Pros | Cons |
|--|--|
| Solve classification & regression Power to handle large data with high dimension Handy at missing data | Not very good for regression (limited in predicting) Like blackbox model and hard to control parameter Slow processing speed |



5.3. Ensemble

5.3.2. Bagging (Bootstrap Aggregating)

Reduce variance & Avoid Overfitting by combining results of multiple classifiers on different sub-samples



Classifier: Decision Trees, Regression models, etc.





5.3. Ensemble

5.3.2. Bagging (Bootstrap Aggregating)

What is Bootstrap?

The bootstrap method is a resampling technique used to estimate statistics on a population by sampling a dataset with replacement.



5.3. Ensemble

5.3.2. Bagging (Bootstrap Aggregating)

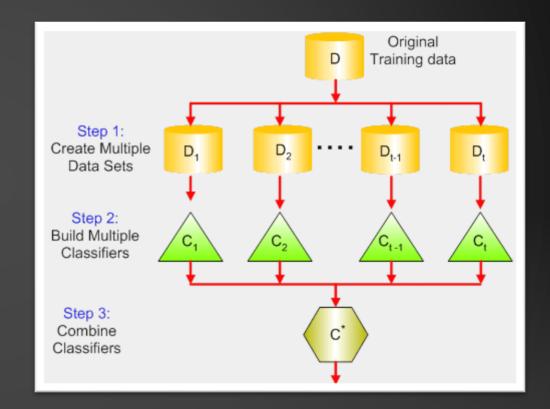
3 steps in Baggings:

Step 1: Here you replace the original data with new sub-sample data using bootstrapping.

Step 2: Train each sub-sample data using ML algorithm

Step 3: Lastly, you use an average value to combine the predictions of all the classifiers, depending on the problem. Generally, these combined values are more robust than a single model.

Bagging has been demonstrated to give impressive improvements in accuracy by combining together hundreds or even thousands of models into a single procedure.





5.3. Ensemble

5.3.2. Bagging (Bootstrap Aggregating)

Bagging in R can be used in many different model:

- ctreebag: used for Decision Tree
- bagFDA: used for Flexible Discriminant Analysis
- IdaBag: Bagging for Linear Discriminant Analysis
- plsBag: Bagging for Principal Linear Regression
- etc





5.3. Ensemble

5.3.3. Boosting

Approach to convert weak predictors to get stronger predictors.





5.3. Ensemble

5.3.3. Boosting

Step 1: A set of classifiers: h_1, \dots, h_k

Classifier h: Decision Trees, Regression models, etc.



5.3. Ensemble

5.3.3. Boosting

Step 2: Create a classifier that combine all classification functions

$$f(x) = sign\left(\sum_{i=1}^{n} \alpha_i h_i(x)\right)$$

- Minimize error "\varepsilon"
- Iterative, each classifier h at a step
- (Re)calculate weight "w" based on error
- Upweight misclassification
- Select next classifier h for the next step

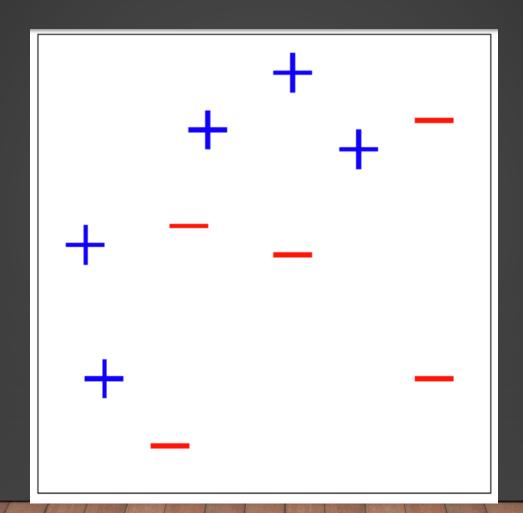
Weight
$$w_{new} = \begin{cases} \frac{1}{2} \frac{1}{1 - \varepsilon} w_{old} & \text{If right} \\ \frac{1}{2} \frac{1}{\varepsilon} w_{old} & \text{If wrong} \end{cases}$$

Error rate:
$$\varepsilon = \sum_{i=1}^{n} w_i$$

Breiman Voting power:
$$\alpha = \frac{1}{2} \ln \left(\frac{1 - \varepsilon}{\varepsilon} \right)$$

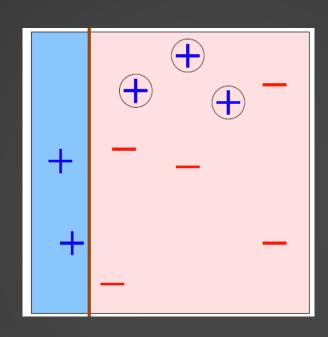
5.3. Ensemble

5.3.3. Boosting



5.3. Ensemble

5.3.3. Boosting



- Upweight misclassification

$$w_{new} = egin{cases} rac{1}{2} rac{1}{1-arepsilon} w_{old} & ext{If right} \\ rac{1}{2} rac{1}{arepsilon} w_{old} & ext{If wrong} \end{cases}$$

$$w_i = 1/10$$

Error rate:
$$\varepsilon = \sum w_i$$

Breiman Voting power:
$$\alpha = \frac{1}{2} \ln \left(\frac{1 - \varepsilon}{\varepsilon} \right)$$

5.3. Ensemble

5.3.3. Boosting

$$D_2$$
 + - 0.167
+ - 0.07 -

Error rate: $\varepsilon = \sum w_i$

$$\varepsilon_2 = 0.21$$
$$\alpha_2 = 0.65$$

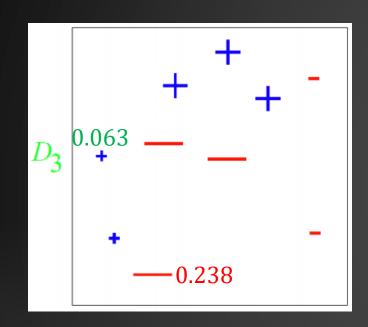
Breiman Voting power: $\alpha = \frac{1}{2} \ln \left(\frac{1 - \varepsilon}{\varepsilon} \right)$

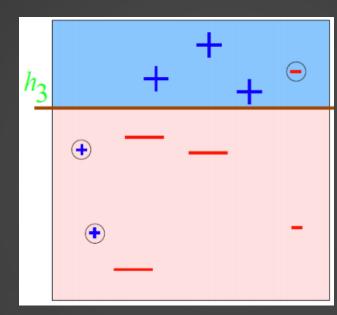
- Upweight misclassification

$$w_{new} = egin{cases} rac{1}{2} rac{1}{1-arepsilon} w_{old} & ext{If right} \ rac{1}{2} rac{1}{arepsilon} w_{old} & ext{If wrong} \end{cases}$$

5.3. Ensemble

5.3.3. Boosting





- Upweight misclassification

$$w_{new} = egin{cases} rac{1}{2} rac{1}{1-arepsilon} w_{old} & ext{If right} \\ rac{1}{2} rac{1}{arepsilon} w_{old} & ext{If wrong} \end{cases}$$

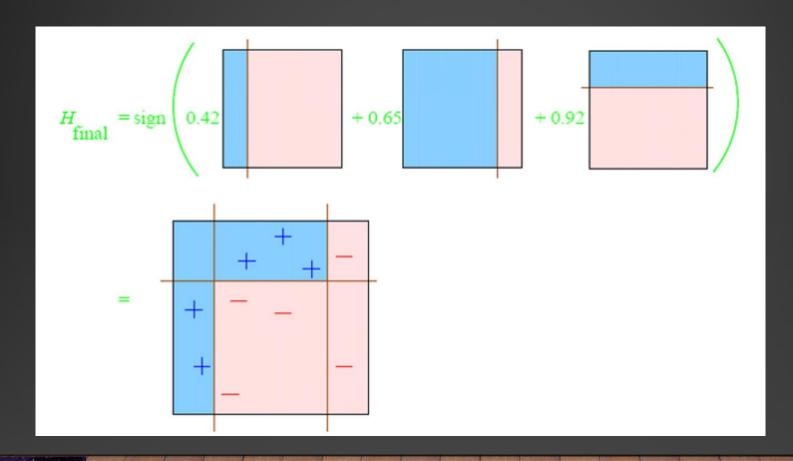
Error rate:
$$\varepsilon = \sum w_i$$

$$\varepsilon_3 = 0.14$$
$$\alpha_2 = 0.92$$

Breiman Voting power: $\alpha = \frac{1}{2} \ln \left(\frac{1 - \varepsilon}{\varepsilon} \right)$

5.3. Ensemble

5.3.3. Boosting



5.3. Ensemble

5.3.3. Boosting-AdaBoost (Adaptive Boosting)

Adaptive: weaker learners are tweaked by misclassify from previous classifier

AdaBoost is best used to boost the performance of decision trees on binary classification problems.

Better for classification rather than regression.

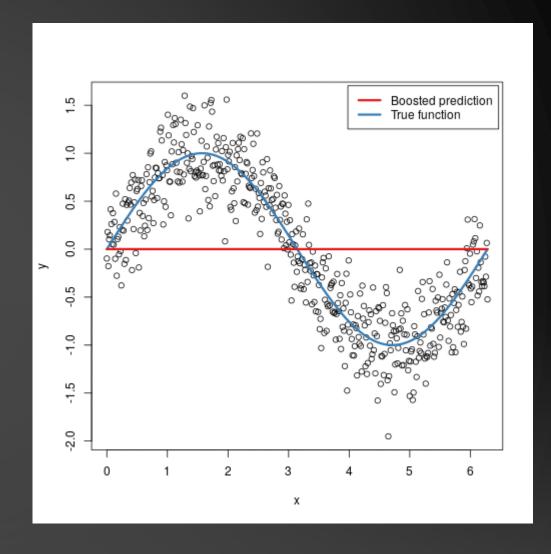
Sensitive to noise





5.3. Ensemble

- 5.3.4. Boosting-Gradient Boosting Machines
 - Extremely popular ML algorithm
 - Widely used in Kaggle competition
 - Ensemble of shallow and weak successive tree, with each tree learning and improving on the previous





- 5.3. Ensemble
- 5.3.4. Boosting-Gradient Boosting Machines
 - Extremely popular ML algorithm
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5.4. Model Based Prediction

- 5.4.1. Naïve Bayes
 - Assuming data follow a probabilistic model
 - Assuming all predictors are independent (Naïve assumption)
 - Use Bayes's theorem to identify optimal classifiers







5.4. Model Based Prediction

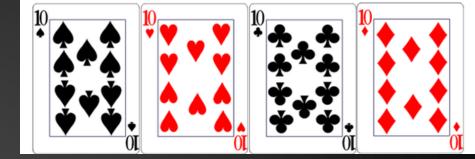
Bayes Theorem

5.4.1. Naïve Bayes

Marginal Probability: P(A); P(B)

Joint Probability: P(A,B) or $P(A \cap B)$

Conditional Probability: P(A | B); P(B | A)





5.4. Model Based Prediction

5.4.1. Naïve Bayes

Marginal Probability: P(A); P(B)

Joint Probability: P(A,B) or $P(A \cap B)$

Conditional Probability: P(A | B); P(B | A)

$$P(A \cap B) = P(A)P(B \mid A) = P(B)P(A \mid B)$$





5.4. Model Based Prediction

5.4.1. Naïve Bayes

Bayes Theorem:

Bayes Theorem

$$P(A|B) = \frac{P(A)P(B|A)}{P(B)}$$



Bayes Theorem

5.4. Model Based Prediction

5.4.1. Naïve Bayes

Example: Picnic Day:

You are planning a picnic today, but the morning is cloudy

- •Oh no! 50% of all rainy days start off cloudy!
- •But cloudy mornings are common (about 40% of days start cloudy)
- •And this is usually a dry month (only 3 of 30 days tend to be rainy, or 10%)
- => What is the chance of rain during the day?





5.4. Model Based Prediction

5.4.1. Naïve Bayes

Example: Picnic Day:

You are planning a picnic today, but the morning is cloudy

- •Oh no! 50% of all rainy days start off cloudy! $P(Cloud \mid Rain) = 0.5$
- •But cloudy mornings are common (about 40% of days start cloudy): P(Cloud)=0.4
- •And this is usually a dry month (only 3 of 30 days tend to be rainy, or 10%): P(Rain)=0.1
- => What is the chance of rain during the day? P(Rain|Cloud)

$$P(Rain|Cloud) = \frac{P(Rain) P(Cloud|Rain)}{P(Cloud)} = \frac{0.1 * 0.5}{0.4} = 0.125 = 12.5\%$$





Bayes Theorem

5.4. Model Based Prediction

5.4.1. Naïve Bayes

Example: Allergy or Not?

Hunter says she is itchy. There is a test for Allergy to Cats, but this test is not always right:

- For people that really do have the allergy, the test says "Yes" 80% of the time
- For people that do not have the allergy, the test says "Yes" 10% of the time ("false positive")

If 1% of the population have the allergy, and Hunter's test says "Yes", what are the chances that Hunter really has the allergy?





Bayes Theorem

5.4. Model Based Prediction

5.4.1. Naïve Bayes

Example: Allergy or Not?

Hunter says she is itchy. There is a test for Allergy to Cats, but this test is not always right:

- For people that really do have the allergy, the test says "Yes" 80% of the time P(Yes | Allergy)=0.8
- For people that do not have the allergy, the test says "Yes" 10% of the time ("false positive")
 P(Yes | NoAllergy)=0.1
- 1% of the population have the allergy: P(Allergy)=0.01
- 99% of the population have NO allergy: P(NoAllergy)=0.99

Hunter's test says "Yes", what are the chances that Hunter really has the allergy?: P(Allergy | Yes)

$$P(Allergy|Yes) = \frac{P(Allergy) P(Yes|Allergy)}{P(Yes)}$$

P(Yes) = P(Yes|Allergy)*P(Allergy) + P(Yes|NoAllergy)*P(NoAllergy)





5.4. Model Based Prediction

5.4.1. Naïve Bayes

$$P(Allergy|Yes) = \frac{P(Allergy) P(Yes|Allergy)}{P(Allergy)*P(Yes|Allergy) + P(NoAllergy)*P(Yes|NoAllergy)}$$

$$= \frac{0.01*0.8}{0.01*0.8+0.99*0.1} = 0.0747 \sim 7\%$$



Bayes Theorem

5.4. Model Based Prediction

5.4.1. Naïve Bayes

Special Bayes Theorem:

$$P(A|B) = \frac{P(A)P(B|A)}{P(A)P(B|A) + P(notA)P(B|notA)}$$

Suppose we have many predictors Ai:

$$P(A_1|B) = \frac{P(A_1)P(B|A_1)}{P(A_1)P(B|A_1) + P(A_2)P(B|A_2) + P(A_3)P(B|A_3)}$$

$$P(A_1)+P(A_2)+P(A_3)=1$$





5.4. Model Based Prediction

- 5.4.1. Naïve Bayes
 - A classification technique based on Bayes Theorem
 - Assume Independence among predictors: A₁, A₂, A_{3,...} A_n
 - Outcome is Independence variable Class B

Predictand Prior Likelihood
Probability $P(B|A) = \frac{P(B)P(A|B)}{P(A|B)}$

Predictor Prior Prob

Posterior Prob



5.4. Model Based Prediction

5.4.1. Naïve Bayes

- Assuming data follow a probabilistic model
- Assuming all predictors are independent (Naïve assumption)
- Use Bayes's theorem to identify optimal classifiers



| Pros | Cons |
|---|--|
| Easy to predict With independent variables, Naive Bayes performs better compared to other Computationally convenient Perform well in categorical compared to numerical variables | Have to make assumption about independence predictors If variable has not observed in training data sets then model will assign 0 probability and unable to make decision |





- 5.4. Model Based Prediction
- 5.4.2. Linear Discriminant Analysis
 - LDA is a supervised learning model that is similar to logistic regression in that the outcome variable is categorical and can therefore be used for classification.
 - LDA is useful with two or more class of objects



5.4. Model Based Prediction

5.4.2. Linear Discriminant Analysis

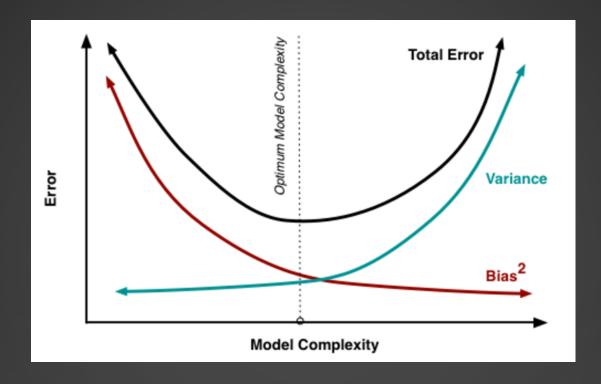
| Logistic Regression | Linear Discriminant Analysis |
|---|---|
| Supervised Learning with Categorical outcome Deal with only binary outcome Use logistic function to model probability | Supervised Learning with Categorical outcome Deal with more than 2 outcomes Used Bayes Theorem to model probability |

| Naïve Bayes | Linear Discriminant Analysis |
|---|---|
| Categorical outcome Assume Independence input (cross-correlation is ~ 0) | Categorical outcomeAssume Gaussian Distribution for inputSensitive to Overfitting |





5.5. Regularization and Variable selection



In order to reduce the Model Complexity or to avoid Multi-Collinearity, one needs to reduce the number of covariates (or set the coefficient to be zero). If the coefficients are too large, let's penalize them to enforce them to be smaller





5.5. Regularization and Variable selection

5.4.1. Regularization

Recap about Linear Regression:

- Observation (output): y
- Independent variable (input): X
- Linear Regression model: $y = \beta_0 + \beta x + \varepsilon$
- Multi-Linear Regression: $y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 \dots + \beta_n x_n + \varepsilon$



5.5. Regularization and Variable selection

5.4.1. Regularization

To determine the optimal β , we resolve the least square problem using the "Lost Function":

$$\hat{\beta} = \underset{\beta}{\operatorname{argmin}} \sum_{i=1}^{n} (y_i - \beta x_i)^2 = \underset{\beta}{\operatorname{argmin}} \|y - \beta x\|_2^2$$

Need to add some penalty term



5.5. Regularization and Variable selection

5.4.1. Regularization

Ridge Regression
$$\hat{\beta}_{ridge} = \sum_{i=1}^{n} (y_i - \beta_i x_i)^2 + \lambda \sum_{j=1}^{m} \beta_j^2$$

Lost Function:

- Shrink the coefficient that contribute most error
- Reduce magnitude of coef that contribute the most to increasing L

 λ : Regularization Penalty, to be selected that the model minimized the error β : vector with many components as predictors

• As
$$\lambda \to 0$$
, $\hat{\beta}_{ridge} \to \hat{\beta}_{OLS}$;

• As
$$\lambda \to \infty$$
, $\hat{\beta}_{ridge} \to 0$.



5.5. Regularization and Variable selection

5.4.1. Regularization

LASSO Lost Function

$$\sum_{i=1}^{n} (y_i - \beta_i x_i)^2 + \lambda \sum_{i=1}^{m} |\beta_i|$$

- Set these coefficient to 0
- Retain only most important coef

- λ : Regularization Penalty, to be selected that the model minimized the error
- β : vector with many components as predictors



5.5. Regularization and Variable selection

5.4.1. Regularization

Package GLMNET that minimizes the Lost Function:

$$\sum_{i=1}^{n} (y_i - \beta_i x_i)^2 + \lambda \left[(1 - \alpha) \sum_{j=1}^{n} |\beta_j| \right]$$

 $\alpha = 0$: pure Ridge Regression

 $\alpha = 1$: pure LASSO

 $0 < \alpha < 1$: Elastic Nets



5.5. Regularization and Variable selection

5.4.1. Regularization

Lcavol: log(cancer volume)
Lweight: log(prostate weight)

Age: age

Lbph: log(benign prostatic hyperplasia amount)

Svi: seminal vesicle invasion

Lcp: log(capsular penetration)

Gleason: Gleason score

Pgg45: percentage Gleason scores 4 or 5

Lpsa: log(prostate specific antigen)



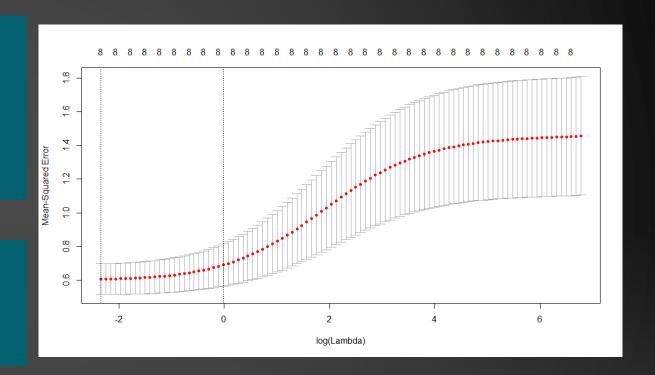


5.5. Regularization and Variable selection

5.4.2. Ridge Regression

```
library(glmnet)
y <- training$lpsa
x <- training[,-c(9,10)]
x <- as.matrix(x)
Ridge     <- cv.glmnet(x,y,alpha=0)
plot(Ridge)</pre>
```

```
> cvfit_Ridge$lambda.min
[1] 0.09645702
> cvfit_Ridge$lambda.1se
[1] 1.189167
```

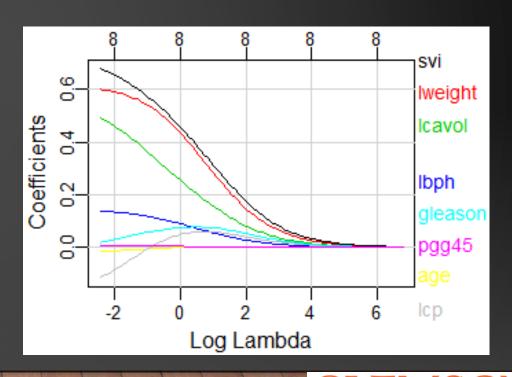




5.5. Regularization and Variable selection

5.4.2. Ridge Regression

```
> coef(cvfit_Ridge,s=cvfit_Ridge$lambda.min)
9 x 1 sparse Matrix of class "dgCMatrix"
(Intercept)
            0.075575516
lcavol
         0.486389709
lweight
        0.599589889
           -0.014473630
age
1bph
           0.137304129
svi
      0.674693010
           -0.110437780
1cp
gleason
         0.019949363
            0.006929742
pgg45
```



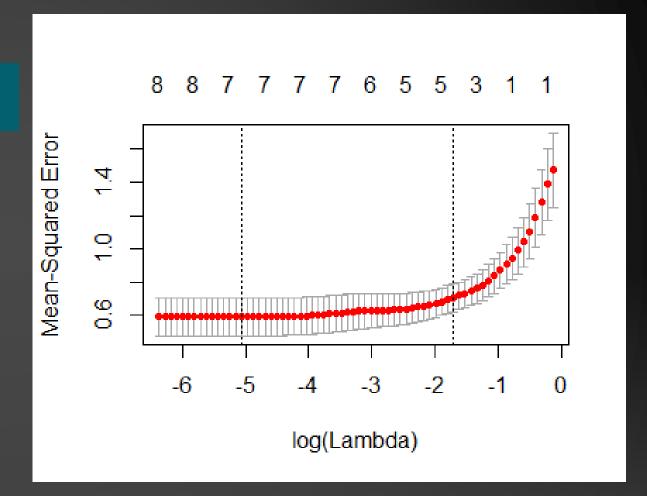




5.5. Regularization and Variable selection

5.4.3. LASSO

cvfit_LASSO c- cv.glmnet(x,y,alpha=1)
plot(cvfit_LASSO)



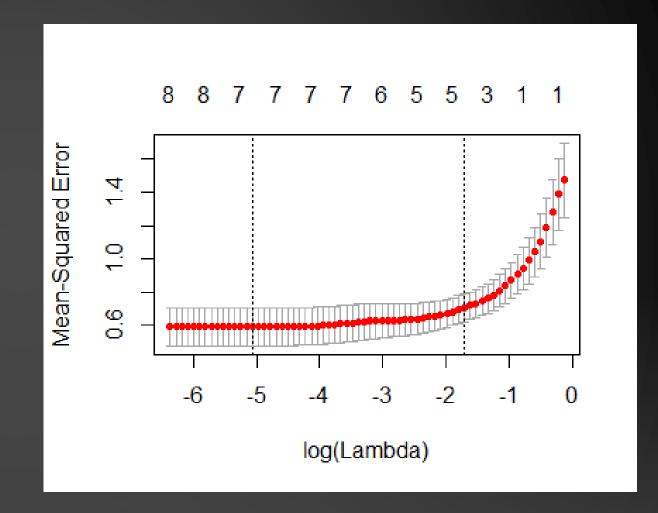




5.5. Regularization and Variable selection

5.4.3. LASSO

```
> cvfit_LASSO$lambda.min
[1] 0.006346228
> cvfit_LASSO$lambda.1se
[1] 0.1807428
```



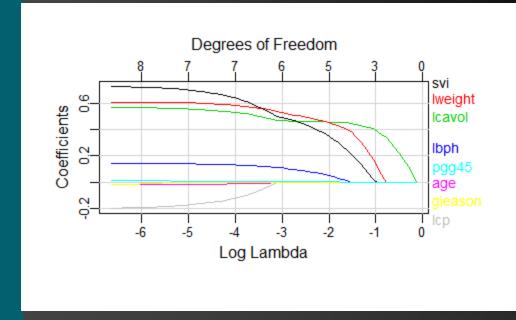




5.5. Regularization and Variable selection

5.4.3. LASSO

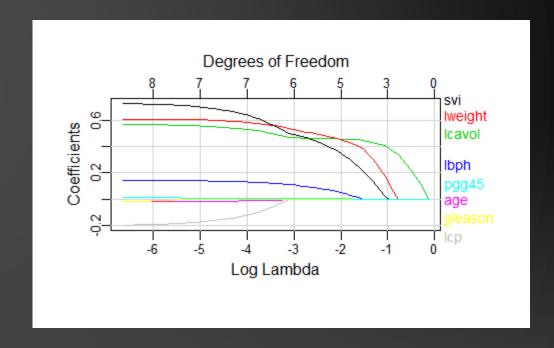
```
> coef(cvfit_LASSO,s=cvfit_LASSO$lambda.1se)
9 x 1 sparse Matrix of class "dgCMatrix"
(Intercept) 0.2602328553
lcavol
            0.4549883478
lweight
            0.4183048097
age
1bph
            0.0199337240
            0.2745819504
svi
1cp
gleason
            0.0005623797
pgg45
```





5.5. Regularization and Variable selection

5.4.3. LASSO





5.5. Regularization and Variable selection

5.4.4. ELASTICNET?





5.6. Dimension Reduction

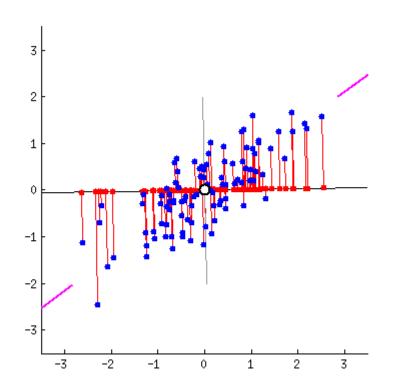
- When there are lots of covariates
- Computing resources limited
- Reduce covariates based on coherence
- Reduce noise



5.6. Dimension Reduction

Principal Component Analysis

- Handy with large data
- Where many variables correlate with one another, they will all contribute strongly to the same principal component
- Each principal component sums up a certain percentage of the total variation in the dataset
- More PCs, more summarization of the original data sets





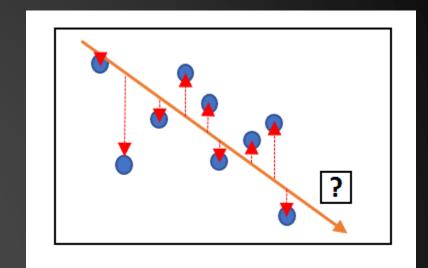


5.6. Dimension Reduction

Principal Component Analysis

Variance: Measure of the spread of data with its mean

$$\sigma^2 = \text{var}(X) = \frac{\sum_{i=1}^{n} (X_i - \bar{X})^2}{n-1}$$



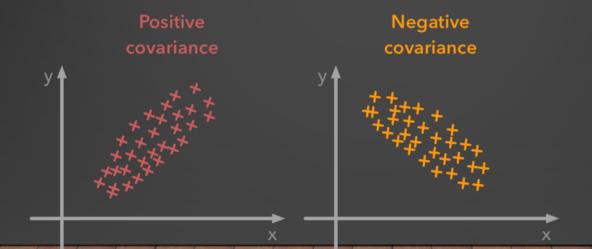


5.6. Dimension Reduction

Principal Component Analysis

Covariance: Measure how much each of the dimensions varies from the mean with respect to each other

$$cov(X,Y) = \frac{\sum_{i=1}^{n} (X_i - \bar{X})(Y_i - \bar{Y})}{n-1}$$







5.6. Dimension Reduction

Principal Component Analysis

Covariance Matrix: A square matrix for 3 or more dimension data

$$M = \begin{bmatrix} Cov(X,X) & Cov(X,Y) & Cov(X,Z) \\ Cov(Y,X) & Cov(Y,Y) & Cov(Y,Z) \\ Cov(Z,X) & Cov(Z,Y) & Cov(Z,Z) \end{bmatrix}$$

- Diagonal: variances of variables
- Cov(X,Y) = Cov(Y,X)
- m dimensional data results in mxm matrix



5.6. Dimension Reduction

Principal Component Analysis

Eigenvalue problem:

$$M. v = \lambda. v$$

$$\begin{bmatrix} 2 & 3 \\ 2 & 1 \end{bmatrix} \times \begin{bmatrix} 3 \\ 2 \end{bmatrix} = \begin{bmatrix} 12 \\ 8 \end{bmatrix} = 4 \times \begin{bmatrix} 3 \\ 2 \end{bmatrix}$$

M: m x m matrix

v: m x 1 non-zero vector

λ: scalar

$$\lambda$$
 = 4: eigenvalues

$$v = \begin{bmatrix} 3 \\ 2 \end{bmatrix}$$
: eigenvectors

Eigenvector: direction

Eigenvalue: how much variance there is in the eigenvector



5.6. Dimension Reduction

Principal Component Analysis

Some characteristics of eigenvalues/eigenvectors

- Given mxm matrix, we can find m eigenvectors and m eigenvalues
- Eigenvectors can only be found for square matrix.
- Not every square matrix has eigenvectors
- A square matrix A and its transpose have the same eigenvalues but different eigenvectors
- The eigenvalues of a diagonal or triangular matrix are its diagonal elements.
- Eigenvectors of a matrix **A** with distinct eigenvalues are linearly independent.





5.6. Dimension Reduction

Principal Component Analysis

Covariance Matrix: A square matrix for 3 or more dimension data

$$M = \begin{bmatrix} Cov(X,X) & Cov(X,Y) & Cov(X,Z) \\ Cov(Y,X) & Cov(Y,Y) & Cov(Y,Z) \\ Cov(Z,X) & Cov(Z,Y) & Cov(Z,Z) \end{bmatrix}$$

Find m eigenvectors and m eigenvalues of the Matrix M



5.6. Dimension Reduction

Principal Component Analysis

Eigenvector with the largest eigenvalue forms the first principal component of the data set ... and so on ...

