$$P(C|x_1, x_2) = \frac{P(x_1, x_2|C) \cdot P(C)}{P(x_1, x_2)}$$

$$= \frac{P(x_1|C) \cdot P(x_2|C) \cdot P(C)}{P(x_1, x_2)}$$

# Lecture 8: Naive Bayesian & LDA

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# AGENDA

01	Naive Bayesian Classifier
02	Linear Discriminant Analysis
03	R Exercise

# Naive Bayesian Classifier

#### Classification revisited





















Men

Vs.

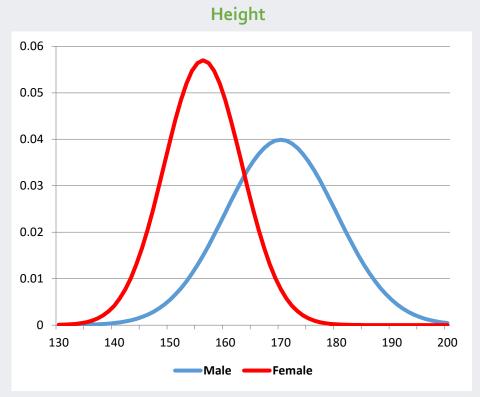
Women

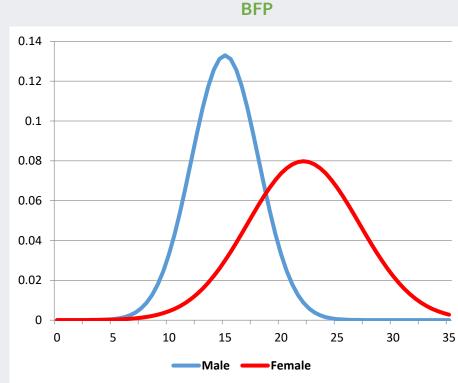




#### Assumption

- ✓ There are two attributes: height & body fat percentage (BFP)
- √ There are equal number of male and female
- ✓ Actual probability distributions for all attribute & gender pairs are known

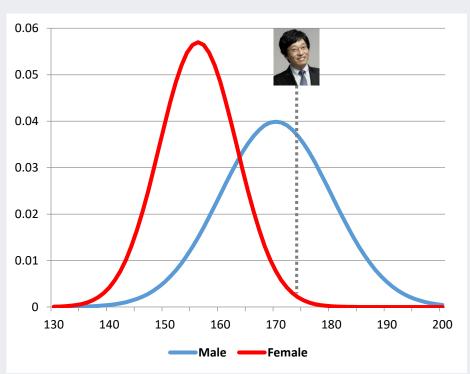


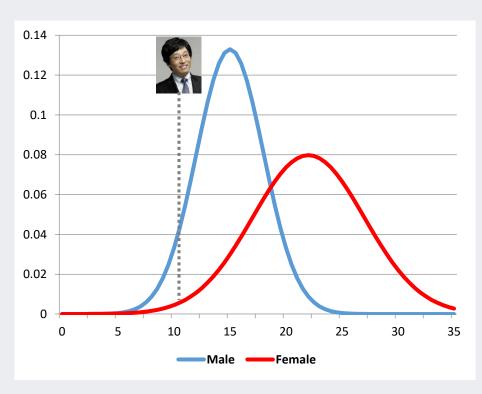


Let's classify



with the given information



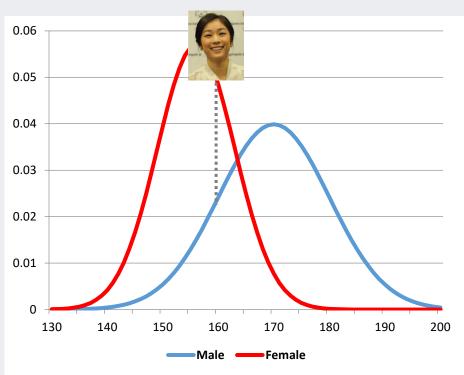


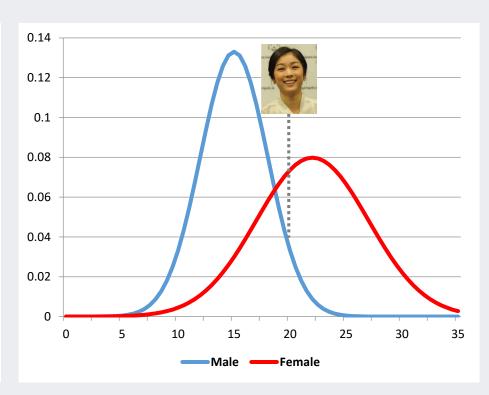
✓ Classify him as Male

Let's classify



with the given information



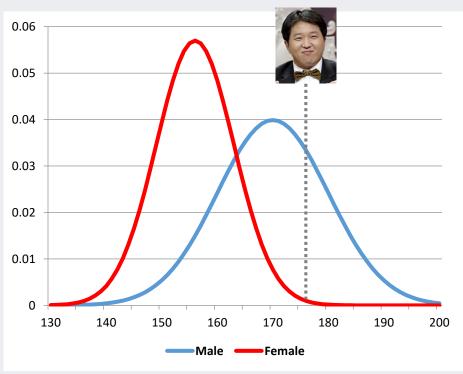


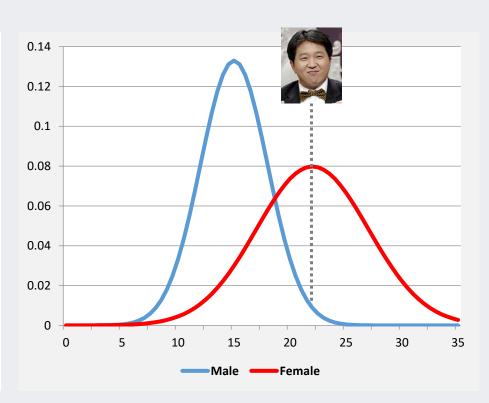
✓ Classify her as Female

What about



9





✓ Classify this person as Male or Female???

# Naïve Bayesian Classification: Theory

Bay's Rule (one of the most important rules in statistics)

$$P(C_i|x_1, x_2) = \frac{P(x_1, X_2|C_i) \cdot P(C_i)}{P(x_1, x_2)}$$

Naive: Let's assume that all variables are statistically independent to each other

$$= \frac{P(x_1|C_i) \cdot P(x_2|C_i) \cdot P(C_i)}{P(x_1, x_2)}$$

# Naïve Bayesian Classification: Theory

For the previous example, we should compare the following two probabilities

$$P(M|H, W, BFS) = \frac{P(H|M) \cdot P(W|M) \cdot P(BFS|M) \cdot P(M)}{P(H, W, BFS)}$$

$$P(\mathbf{F}|H, W, BFS) = \frac{P(H|\mathbf{F}) \cdot P(W|\mathbf{F}) \cdot P(BFS|\mathbf{F}) \cdot P(\mathbf{F})}{P(H, W, BFS)}$$

✓ Assign to the class with the highest posterior probability

### Naïve Bayesian Classification

Compute the posterior probabilities

$$P(H|M) \cdot P(W|M) \cdot P(BFS|M) = 0.035 \times 0.01 \times 0.5 = 0.000175$$
  
 $P(H|F) \cdot P(W|F) \cdot P(BFS|F) = 0.001 \times 0.08 \times 0.5 = 0.00004$ 

• Classify the person as Male

### Exact Bayesian Classifier

# Find all the other records whose variable values are exactly identical to the test entity



• Find all the other people with the same height and BFS.

Person	Height	BFS	Class
홍길동	178	11	M
김영희	178	11	F
김철수	178	11	М
김가네	178	11	М

Variables are not assumed to be statistically independent

$$P(C_i \mid x_1, x_2, ..., x_d) = \frac{P(x_1, x_2, ..., x_d \mid C_i)P(C_i)}{P(x)}$$

# Exact Bayesian Classifier

#### Find the prevalent class



Determine what classes they all belong to and which class is more prevalent.

Person	Height	BFS	Class
홍길동	178	11	М
김영희	178	11	F
김철수	178	11	М
김가네	178	11	М

• 3 males and I female.

### Exact Bayesian Classifier

#### Assign the prevalent to the new record



Person	Height	BFS	Class
홍길동	178	11	М
김영희	178	11	F
김철수	178	11	М
김가네	178	11	М

- 3 males and I female.
- He is classified as male.

Difficult to find the exact same records when the there are many attributes(features) with small number of training data.

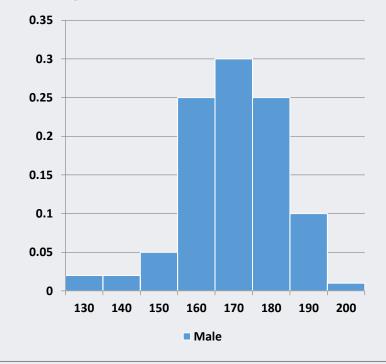
#### Prepare the training data

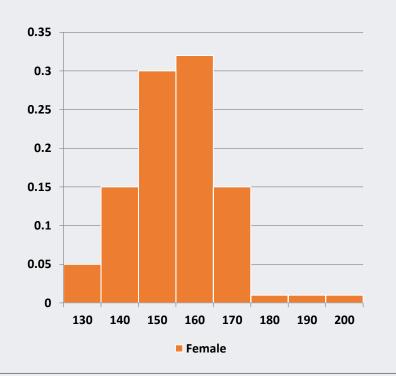
- Define attributes and collect training data
  - ✓ Total training data: 200 (100 males, 100 females)
  - ✓ Height & BFS

Record	Height	BFS	Class
1	187	15	M
2	165	25	F
3	174	14	М
4	156	29	F
N	168	12	М

#### Estimate the probability distribution

- Estimate the probability distribution of the attributes for each class.
- Height

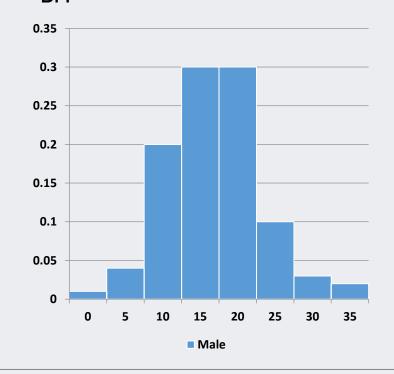


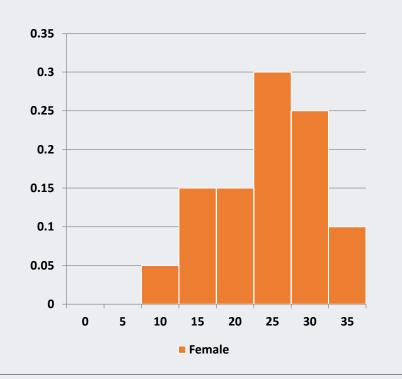


#### Estimate the probability distribution

■ Estimate the probability distribution of the attributes for each class.

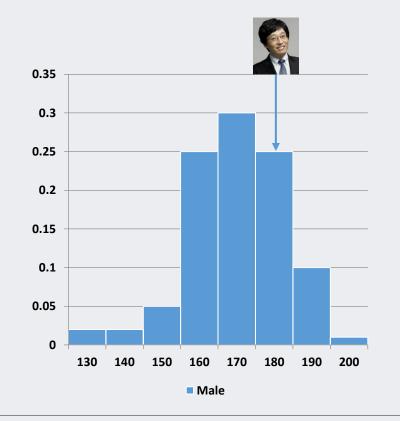
#### BFP

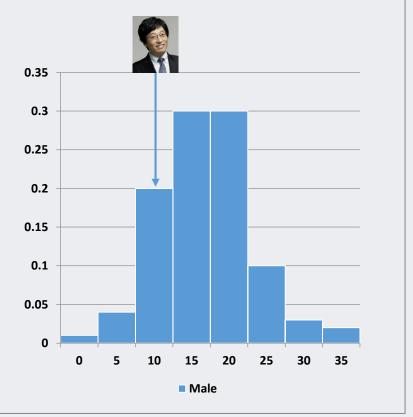




#### Compute the conditional probability for each attribute

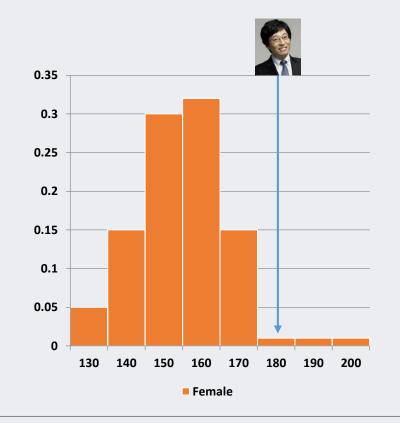
P(Height = 178 | Male) = 0.25, P(BFP = 11 | Male) = 0.2

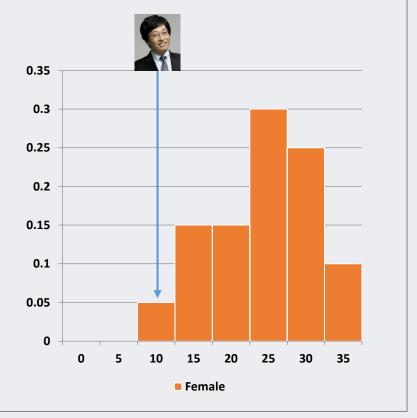




#### Compute the conditional probability for each attribute

■ P(Height = 178 | Female) = 0.01, P(BFP = 11 | Female) = 0.05





#### Compute the posterior probability

Compute the posterior probability for each class

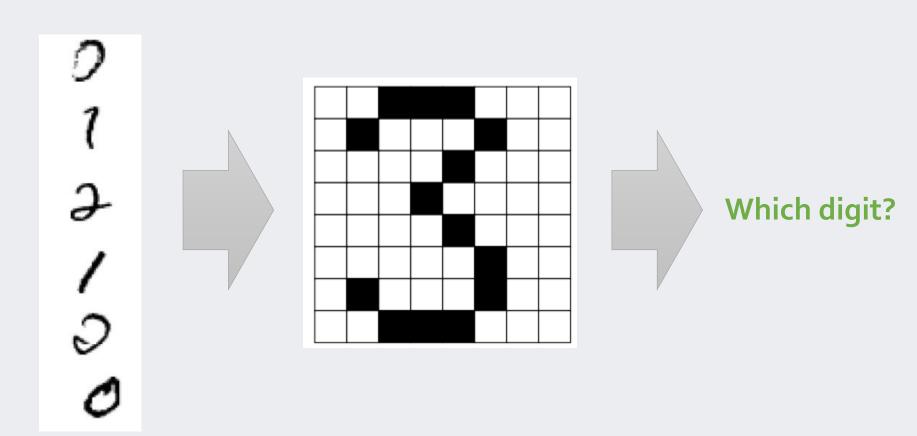
```
    ✓ P(Height = 178, BFP = 11 | Male)*P(Male)
    = P(Height = 178 | Male)* P(BFP = 11 | Male) *P(Male)
    = 0.25*0.2*0.5 = 0.025
    ✓ P(Height = 178, BFP = 11 | Female) *P(Female)
    = P(Height = 178 | Female)* P(BFS = 11 | Female) *P(Female)
    = 0.01*0.05*0.5 = 0.00025
```

#### Make a decision

- P(Height=178, BFS=11 | Male)P(Male) > P(Height=178, BFS=11 | Female)P(Female)
  - ✓ Classify him as male
- What if there are 400 males and 100 females in the training data?
  - ✓ Consider the prior probability P(Male) & P(Female)
  - $\checkmark$  P(Height=178, BFS=11 | Male)\*P(Male) = 0.05\*0.8 = 0.04
  - ✓ P(Height=178, BFS=11 | Female)\*P(Female) = 0.0005\*0.2 = 0.0001
  - ✓ Classify him as male

### Hand digit recognition

- ✓ Input: pixel grids
- ✓ Classes: a digit 0-9



#### Feature definition

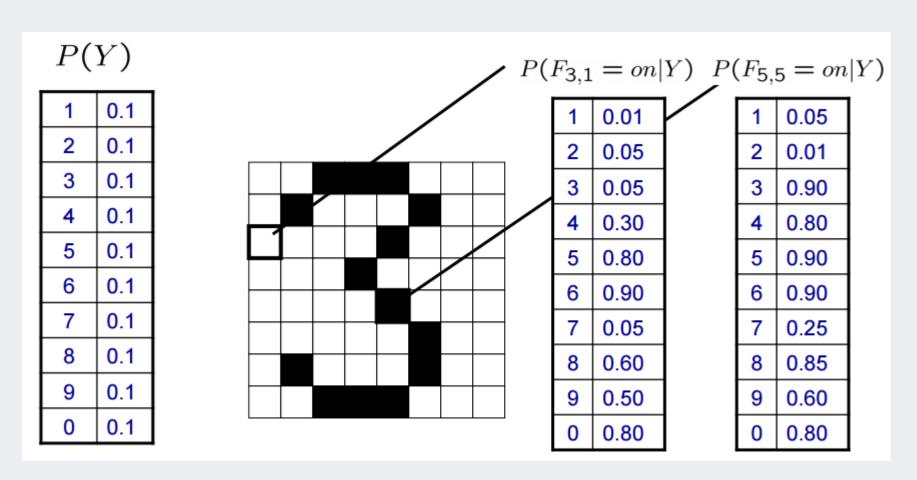
- ✓ One feature  $f_{ij}$  for each grid position <i, j>
- ✓ Possible feature values are on/off, based on whether intensity is more or less than 0.5 in underlying image.
- ✓ Each input maps to a feature vector, e.g.

$$Arr VF_{0,0} = 0 \quad F_{0,1} = 0 \quad F_{0,2} = 1 \quad F_{0,3} = 1 \quad F_{0,4} = 0 \quad \dots F_{15,15} = 0$$

#### Naïve Bayesian Model

$$P(Y|F_{0,0}...F_{15,15}) \propto P(Y) \prod_{i,j} P(F_{i,j}|Y)$$

What has to be learned?



#### Training

√ Count the target class ratio for each grid

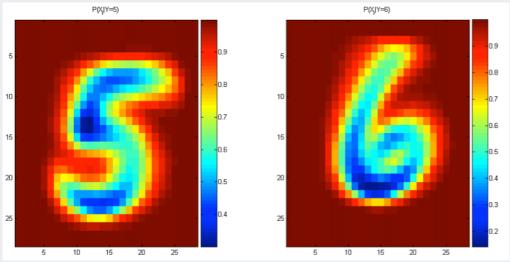
- Prior:

$$P(Y = y) = \frac{Count(Y = y)}{\sum_{y'} Count(Y = y')}$$

– Observation distribution:

$$P(X_i = x | Y = y) = \frac{Count(X_i = x, Y = y)}{\sum_{x'} Count(X_i = x', Y = y)}$$

#### Trained examples

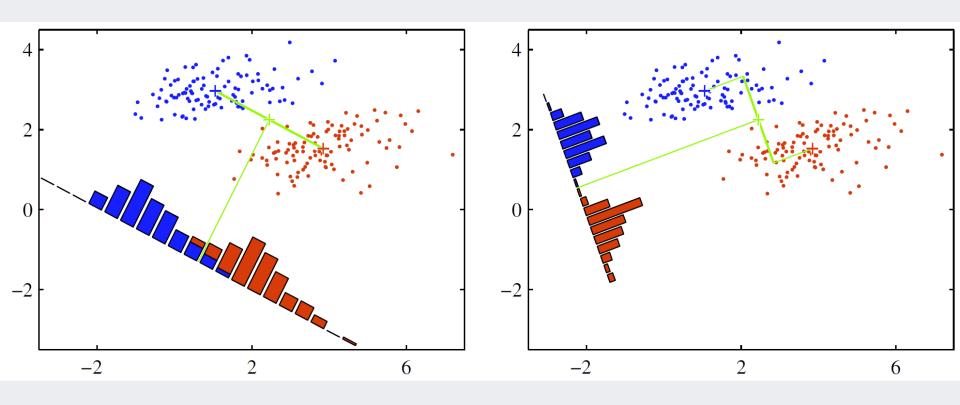


# AGENDA

01	Naive Bayesian Classifier
02	Linear Discriminant Analysis
03	R Exercise

#### • Fisher's LDA

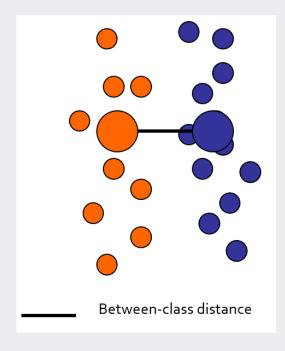
√ Which line is better to discriminate two classes after projection?

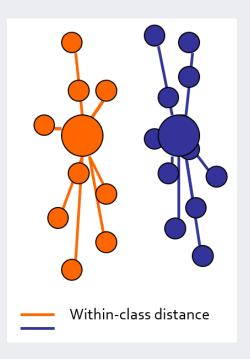


√ Find the most distinguishable vector!

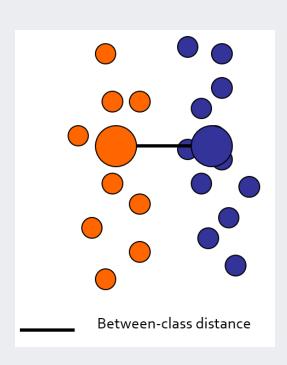
(Source: Bishop (2006))

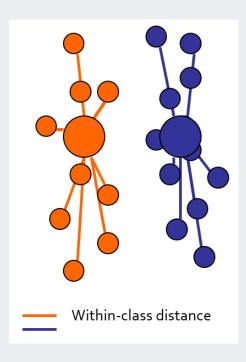
- Two type of class distances
  - √ Between-class distance
    - Distance between the centroids of different classes.
  - √ Within-class distance
    - Accumulated distance of an instance to the centroid of its class

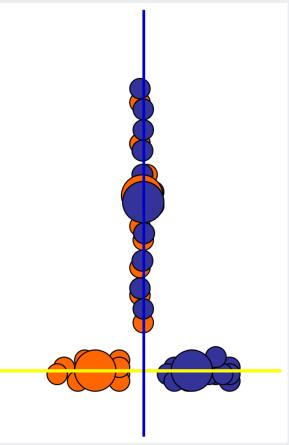




- (Fisher's) Linear Discriminant Analysis
  - ✓ Find most discriminant projection by maximizing between-class distance (variance) and minimizing within-class distance (variance)







#### • Fisher's LDA (cont')

 $\checkmark$  Take the D-dimensional input vector  $\mathbf{x}$  and project it down to one dim.

$$y = \mathbf{w}^T \mathbf{x}$$

✓ Consider a two-class problem in which there are  $N_1$  &  $N_2$  observations in  $C_1$  and  $C_2$ , respectively.

$$\mathbf{m}_1 = \frac{1}{N_1} \sum_{n \in C_1} \mathbf{x}_n \qquad \mathbf{m}_2 = \frac{1}{N_2} \sum_{n \in C_2} \mathbf{x}_n$$

#### Fisher's LDA (cont')

✓ Objective I: Choose w to maximize the separation of the projected class means (between class variance)

$$m_2 - m_1 = \mathbf{w}^T (\mathbf{m}_2 - \mathbf{m}_1)$$
  $m_k = \mathbf{w}^T \mathbf{m}_k$ 

✓ Objective 2: Choose w to minimize the variance in each class after projection (within class variance)

$$s_k^2 = \sum_{n \in C_k} (y_k - m_k)^2$$

- Fisher's LDA (cont')
  - √ Fisher's criterion
    - The ratio of the between-class variance to the within-class variance

$$J(\mathbf{w}) = \frac{(m_2 - m_1)^2}{s_1^2 + s_2^2} = \frac{\mathbf{w}^T \mathbf{S}_B \mathbf{w}}{\mathbf{w}^T \mathbf{S}_W \mathbf{w}}$$

$$\mathbf{S}_B = (\mathbf{m}_2 - \mathbf{m}_1)(\mathbf{m}_2 - \mathbf{m}_1)^T$$

$$\mathbf{S}_W = \sum_{n \in C_1} (\mathbf{x}_n - \mathbf{m}_1)(\mathbf{x}_n - \mathbf{m}_1)^T + \sum_{n \in C_2} (\mathbf{x}_n - \mathbf{m}_2)(\mathbf{x}_n - \mathbf{m}_2)^T$$

• Fisher's LDA (cont')

• Differentiating the Fisher's criterion w.r.t. w, then J(w) is maximized when

$$(\mathbf{w}^T \mathbf{S}_B \mathbf{w}) \mathbf{S}_W \mathbf{w} = (\mathbf{w}^T \mathbf{S}_W \mathbf{w}) \mathbf{S}_B \mathbf{w}$$

- $S_B w$  is always in the direction of  $(m_2 m_1)$
- Can drop the scalar factor  $(\mathbf{w}^{\mathrm{T}}\mathbf{S}_{\mathrm{B}}\mathbf{w})$  and  $(\mathbf{w}^{\mathrm{T}}\mathbf{S}_{\mathrm{W}}\mathbf{w})$
- Then, obtain Fisher's linear discriminant

$$\mathbf{w} \propto \mathbf{S}_W^{-1}(\mathbf{m}_2 - \mathbf{m}_1)$$

# AGENDA

01	Naive Bayesian Classifier
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#### Target Data: Wisconsin Breast Cancer

- √ Predicting whether a patient is malignant or benign
- √ The real-valued features for the following information are computed for each cell nucleus:
  - a) radius (mean of distances from center to points on the perimeter)
  - b) texture (standard deviation of gray-scale values)
  - c) perimeter
  - d) area
  - e) smoothness (local variation in radius lengths)
  - f) compactness (perimeter^2 / area 1.0)
  - g) concavity (severity of concave portions of the contour)
  - h) concave points (number of concave portions of the contour)
  - i) symmetry
  - j) fractal dimension ("coastline approximation" 1)
- ✓ Mean, standard error, and worst values are used as input variables.

• Write a performance evaluation function and initialize the result summary table

```
# Performance Evaluation Function -----
perf eval <- function(cm){</pre>
 # True positive rate: TPR (Recall)
  TPR \leftarrow cm[2,2]/sum(cm[2,])
  # Precision
  PRE <- cm[2,2]/sum(cm[,2])
  # True negative rate: TNR
  TNR \langle -cm[1,1]/sum(cm[1,])
  # Simple Accuracy
  ACC \leftarrow (cm[1,1]+cm[2,2])/sum(cm)
  # Balanced Correction Rate
  BCR <- sqrt(TPR*TNR)
 # F1-Measure
  F1 <- 2*TPR*PRE/(TPR+PRE)
  return(c(TPR, PRE, TNR, ACC, BCR, F1))
# Result summary
Perf.Table <- matrix(0, nrow = 2, ncol = 6)
rownames(Perf.Table) <- c("Naive Bayes", "LDA")</pre>
colnames(Perf.Table) <- c("TPR", "Precision", "TNR", "Accuracy", "BCR", "F1-Measure")</pre>
```

• Load the data and divide the dataset into training (70%) and test (30%)

```
# Load the wdbc data
Wdbc.Data <- read.csv("wdbc.csv", header = FALSE)
# Divide the dataset into the training (70%) and Test (30%) datasets
trn.idx <- sample(1:nrow(Wdbc.Data), round(0.7*nrow(Wdbc.Data)))</pre>
```

Train the Naive Bayesian Classifier

```
# Classifier 1: Naive Bayesian Classifier
# e1071 package install
install.packages("e1071")
library(e1071)

# Use the dataset without normalization
NB.Trn.Data <- Wdbc.Data[trn.idx,]
colnames(NB.Trn.Data)[31] <- "Target"

NB.Tst.Data <- Wdbc.Data[-trn.idx,]
colnames(NB.Tst.Data)[31] <- "Target"

# Training the Naive Bayesian Classifier
NB.model <- naiveBayes(Target ~ ., data = NB.Trn.Data)
NB.model$apriori
NB.model$tables</pre>
```

#### Check the trained parameters

```
> NB.model$apriori
Y
B M
150 248
```

```
> NB.model$tables
$V1
  V1
       [,1]
                [,2]
 B 17.44373 3.259898
 M 12.12940 1.780143
$V2
   V2
       [,1]
                [,2]
  B 21.67427 3.723244
 M 17.78270 3.960919
$V3
   V3
        [,1]
              [,2]
  B 115.16073 22.17670
 M 77.91379 11.74798
$V4
   V4
       [,1]
             [,2]
 B 976.8833 374.8957
 M 461.6625 133.4494
$V5
   V5
         [,1]
                    [,2]
  B 0.10332747 0.01313281
 M 0.09271343 0.01372253
$V6
   V6
          [,1]
                    [,2]
  B 0.14425960 0.05483254
  M 0.07890472 0.03337068
```

Evaluate the classification performance

```
# Predict the new input data based on Naive Bayesian Classifier
NB.posterior = predict(NB.model, NB.Tst.Data, type = "raw")
NB.prey = predict(NB.model, NB.Tst.Data, type = "class")

NB.cfm <- table(NB.Tst.Data[,31], NB.prey)
NB.cfm

Perf.Table[1,] <- perf_eval(NB.cfm)
Perf.Table</pre>
```

#### • Train LDA

```
# Classifier 2: Linear Discriminant Analysis ----
install.packages("MASS")
library(MASS)

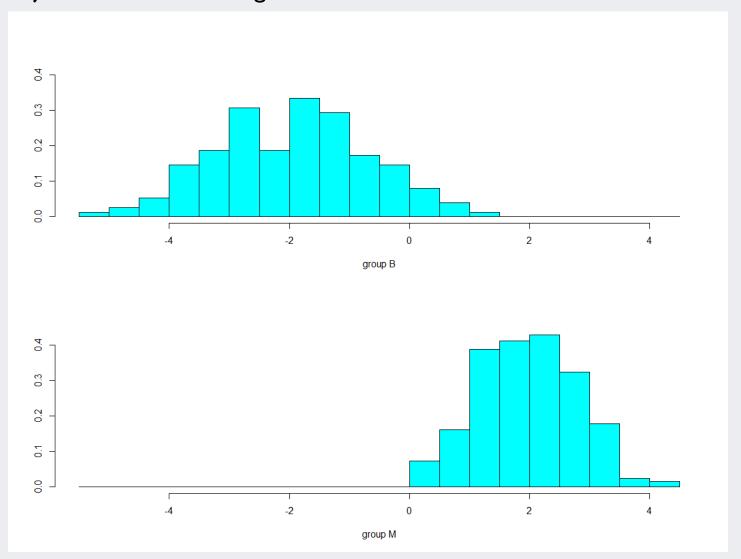
# Use the dataset without normalization
LDA.Trn.Data <- Wdbc.Data[trn.idx,]
colnames(LDA.Trn.Data)[31] <- "Target"

LDA.Tst.Data <- Wdbc.Data[-trn.idx,]
colnames(LDA.Tst.Data)[31] <- "Target"

# Training LDA
LDA.model <- lda(Target ~ ., data = LDA.Trn.Data)

# Training result of LDA
plot(LDA.model)</pre>
```

• Best projection and the histogram of the two classes



#### Evaluate the performance

```
# Predict the unknown observations based on the LDA
LDA.Predict <- predict(LDA.model, LDA.Tst.Data)

names(LDA.Predict)
LDA.Predict$class
LDA.Predict$posterior
LDA.Predict$x

LDA.cfm <- table(LDA.Tst.Data$Target, LDA.Predict$class)
LDA.cfm

Perf.Table[2,] <- perf_eval(LDA.cfm)
Perf.Table</pre>
```

