



# Model based prediction

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# Basic idea

1. Assume the data follow a probabilistic model
2. Use Bayes' theorem to identify optimal classifiers

## Pros:

- Can take advantage of structure of the data
- May be computationally convenient
- Are reasonably accurate on real problems

## Cons:

- Make additional assumptions about the data
- When the model is incorrect you may get reduced accuracy

# Model based approach

Wahrscheinlichkeit, dass outcome  $Y$   
zur Klasse  $k$  gehoert,  
gegeben bestimmte  
Prädiktorvariablen ( $x$ )

1. Our goal is to build parametric model for conditional distribution  $P(Y = k | X = x)$
2. A typical approach is to apply [Bayes theorem](#):

$$Pr(Y = k | X = x) = \frac{Pr(X = x | Y = k)Pr(Y = k)}{\sum_{\ell=1}^K Pr(X = x | Y = \ell)Pr(Y = \ell)}$$

total probability

$$Pr(Y = k | X = x) = \frac{f_k(x)\pi_k}{\sum_{\ell=1}^K f_{\ell}(x)\pi_{\ell}}$$

pi\_k: Prior

3. Typically prior probabilities  $\pi_k$  are set in advance.

4. A common choice for  $f_k(x) = \frac{1}{\sigma_k \sqrt{2\pi}} e^{-\frac{(x-\mu_k)^2}{\sigma_k^2}}$ , a Gaussian distribution

5. Estimate the parameters  $(\mu_k, \sigma_k^2)$  from the data.

6. Classify to the class with the highest value of  $P(Y = k | X = x)$

# Classifying using the model

A range of models use this approach

- Linear discriminant analysis assumes  $f_k(x)$  is multivariate Gaussian with same covariances
- Quadratic discriminant analysis assumes  $f_k(x)$  is multivariate Gaussian with different covariances
- [Model based prediction](#) assumes more complicated versions for the covariance matrix
- Naive Bayes assumes independence between features for model building

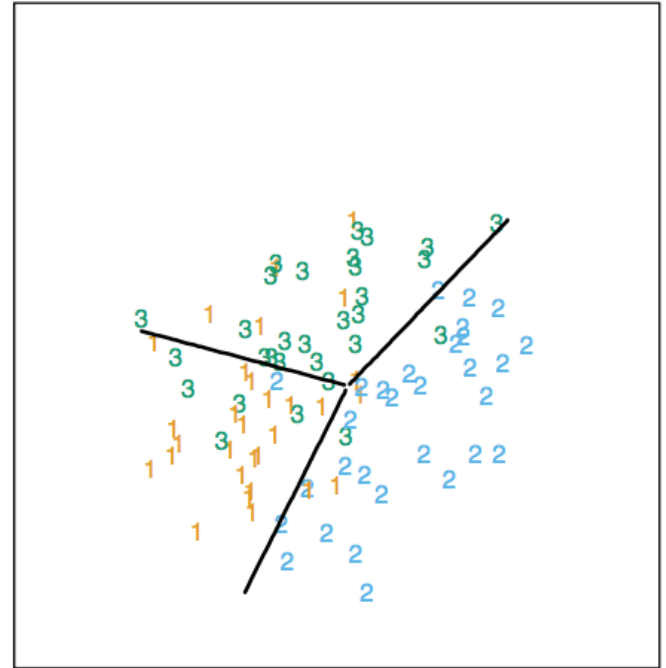
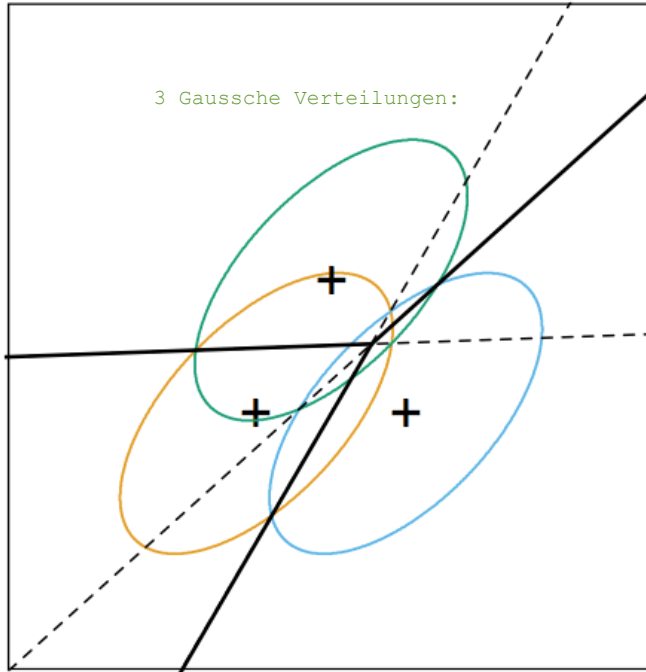
<http://statweb.stanford.edu/~tibs/ElemStatLearn/>

# Why linear discriminant analysis?

$$\begin{aligned} \log \frac{\Pr(Y = k | X = x)}{\Pr(Y = j | X = x)} & \quad \begin{array}{l} \text{Prob(Y gehoert zu Klasse k, gegeben Predikt x)} \\ \text{Prob(Y gehoert zu Klasse j, gegeben Predikt x)} \end{array} \\ &= \log \frac{f_k(x)}{f_j(x)} + \log \frac{\pi_k}{\pi_j} \\ &= \log \frac{\pi_k}{\pi_j} - \frac{1}{2} (\mu_k + \mu_j)^T \Sigma^{-1} (\mu_k + \mu_j) \\ & \quad + x^T \Sigma^{-1} (\mu_k - \mu_j) \end{aligned}$$

<http://statweb.stanford.edu/~tibs/ElemStatLearn/>

# Decision boundaries



# Discriminant function

$$\delta_k(x) = x^T \Sigma^{-1} \mu_k - \frac{1}{2} \mu_k^T \Sigma^{-1} \mu_k + \log(\mu_k)$$

`mu_k`: mean of class k for all features  
`sigma`: Kovarianzmatrix for class k

- Decide on class based on  $\hat{Y}(x) = \operatorname{argmax}_k \delta_k(x)$
- We usually estimate parameters with maximum likelihood

# Naive Bayes

Suppose we have many predictors, we would want to model:  $P(Y = k | X_1, \dots, X_m)$

We could use Bayes Theorem to get:

$$P(Y = k | X_1, \dots, X_m) = \frac{\pi_k P(X_1, \dots, X_m | Y = k)}{\sum_{\ell=1}^K P(X_1, \dots, X_m | Y = \ell) \pi_{\ell}}$$

$$\propto \pi_k P(X_1, \dots, X_m | Y = k)$$

proportional, da Nenner eine Konstante ist

This can be written:

$$P(X_1, \dots, X_m, Y = k) = \pi_k P(X_1 | Y = k) P(X_2, \dots, X_m | X_1, Y = k)$$

$$= \pi_k P(X_1 | Y = k) P(X_2 | X_1, Y = k) P(X_3, \dots, X_m | X_1, X_2, Y = k)$$

$$= \pi_k P(X_1 | Y = k) P(X_2 | X_1, Y = k) \dots P(X_m | X_1, \dots, X_{m-1}, Y = k)$$

We could make an assumption to write this:

naive Annahme:  
dass alle Prediktoren unabhaengig sind.

$$\approx \pi_k P(X_1 | Y = k) P(X_2 | Y = k) \dots P(X_m | Y = k)$$

Speziell nuetzlich bei ganz vielen  
Prediktoren die binaer oder  
kategorial sind.



# Example: Iris Data

```
data(iris); library(ggplot2)
names(iris)
```

```
[1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Width"  "Species"
```

```
table(iris$Species)
```

| setosa | versicolor | virginica |
|--------|------------|-----------|
| 50     | 50         | 50        |

# Create training and test sets

```
inTrain <- createDataPartition(y=iris$Species,  
                                p=0.7, list=FALSE)  
  
training <- iris[inTrain,]  
testing <- iris[-inTrain,]  
dim(training); dim(testing)
```

```
[1] 45 5
```

# Build predictions

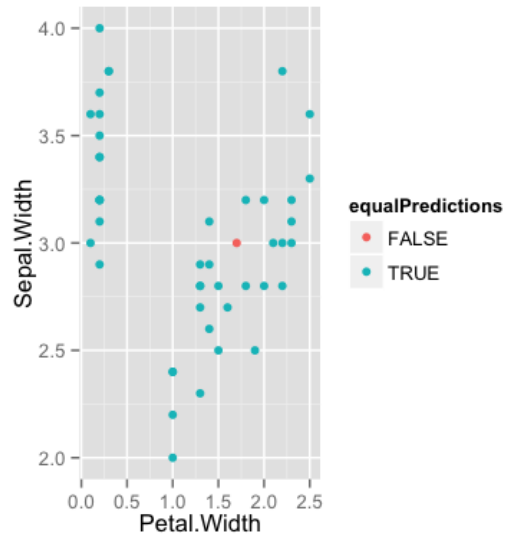
```
modlda = train(Species ~ ., data=training, method="lda") l_inear d_iscriminant a_nalYSIS
modnb = train(Species ~ ., data=training, method="nb") n_aive b_ayes
plda = predict(modlda, testing); pnb = predict(modnb, testing)
table(plda, pnb)
```

|            | pnb    |            |           |
|------------|--------|------------|-----------|
| plda       | setosa | versicolor | virginica |
| setosa     | 15     | 0          | 0         |
| versicolor | 0      | 13         | 1         |
| virginica  | 0      | 0          | 16        |

Bis auf einen Fall uebereinstimmende Kategorisierung von beiden Klassifizierungsmethoden.

# Comparison of results

```
equalPredictions = (plda==pnb)  
ggplot(Petal.Width,Sepal.Width,colour=equalPredictions,data=testing)
```



# Notes and further reading

- [Introduction to statistical learning](#)
- [Elements of Statistical Learning](#)
- [Model based clustering](#)
- [Linear Discriminant Analysis](#)
- [Quadratic Discriminant Analysis](#)