Day 3: Introduction to Machine Learning

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Data Mining and Statistical Learning

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How do we get "true" condition?

- In some domains: through more expensive or extensive tests
- In social sciences: typically by expert annotation or coding
- A scheme should be tested and reported for its reliability

Inter-rater reliability

Different types are distinguished by the way the reliability data is obtained.

Туре	Test Design	Causes of Disagreements	Strength
Stability	test-retest	intraobserver inconsistencies	weakest
Reproduc- ibility	test-test	intraobserver inconsistencies + interobserver disagreements	medium
Accuracy	test-standard	intraobserver inconsistencies + interobserver disagreements + deviations from a standard	strongest

Measures of agreement

- Percent agreement Very simple: (number of agreeing ratings) / (total ratings) * 100%
- Correlation
 - ▶ (usually) Pearson's *r*, aka product-moment correlation
 - ► Formula: $r_{AB} = \frac{1}{n-1} \sum_{i=1}^{n} \left(\frac{A_i \bar{A}}{s_A} \right) \left(\frac{B_i \bar{B}}{s_B} \right)$
 - May also be ordinal, such as Spearman's rho or Kendall's tau-b
 - Range is [0,1]
- Agreement measures
 - Take into account not only observed agreement, but also agreement that would have occured by chance
 - Cohen's κ is most common
 - Krippendorf's α is a generalization of Cohen's κ
 - Both range from [0,1]

Reliability data matrixes

Example here used binary data (from Krippendorff)

Article:	1	2	3	4	5	6	7	8	9	10	
Coder A	1	1	0	0	0	0	0	0	0	0	
Coder B	0	1	1	0	0	1	0	1	0	0	

- ▶ A and B agree on 60% of the articles: 60% agreement
- Correlation is (approximately) 0.10
- Observed disagreement: 4
- Expected *dis*agreement (by chance): 4.4211
- Krippendorff's $\alpha = 1 \frac{D_o}{D_e} = 1 \frac{4}{4.4211} = 0.095$
- Cohen's κ (nearly) identical

Naive Bayes classification

- The following examples refer to "words" and "documents" but can be thought of as generic "features" and "cases"
- We will being with a discrete case, and then cover continuous feature values
- Objective is typically MAP: identification of the maximum a posteriori class probability

Multinomial Bayes model of Class given a Word

Consider J word types distributed across I documents, each assigned one of K classes.

At the word level, Bayes Theorem tells us that:

$$P(c_k|w_j) = \frac{P(w_j|c_k)P(c_k)}{P(w_j)}$$

For two classes, this can be expressed as

$$=\frac{P(w_j|c_k)P(c_k)}{P(w_j|c_k)P(c_k)+P(w_j|c_{\neg k})P(c_{\neg k})}$$
(1)

Multinomial Bayes model of Class given a Word Class-conditional word likelihoods

$$P(c_k|w_j) = \frac{P(w_j|c_k)P(c_k)}{P(w_j|c_k)P(c_k) + P(w_j|c_{\neg k})P(c_{\neg k})}$$

- The word likelihood within class
- The maximum likelihood estimate is simply the proportion of times that word j occurs in class k, but it is more common to use Laplace smoothing by adding 1 to each observed count within class

Multinomial Bayes model of Class given a Word Word probabilities

$$P(c_k|w_j) = \frac{P(w_j|c_k)P(c_k)}{P(w_j)}$$

- This represents the word probability from the training corpus
- Usually uninteresting, since it is constant for the training data, but needed to compute posteriors on a probability scale

Multinomial Bayes model of Class given a Word Class prior probabilities

$$P(c_k|w_j) = \frac{P(w_j|c_k)P(c_k)}{P(w_j|c_k)P(c_k) + P(w_j|c_{\neg k})P(c_{\neg k})}$$

- This represents the class prior probability
- Machine learning typically takes this as the document frequency in the training set
- This approach is flawed for scaling, however, since we are scaling the latent class-ness of an unknown document, not predicting class – uniform priors are more appropriate

Multinomial Bayes model of Class given a Word Class posterior probabilities

$$P(c_k|w_j) = \frac{P(w_j|c_k)P(c_k)}{P(w_j|c_k)P(c_k) + P(w_j|c_{\neg k})P(c_{\neg k})}$$

- This represents the posterior probability of membership in class k for word j
- Under certain conditions, this is identical to what LBG (2003) called P_{wr}
- ► Under those conditions, the LBG "wordscore" is the linear difference between P(c_k|w_j) and P(c_{¬k}|w_j)

"Certain conditions"

- The LBG approach required the identification not only of texts for each training class, but also "reference" scores attached to each training class
- Consider two "reference" scores s₁ and s₂ attached to two classes k = 1 and k = 2. Taking P₁ as the posterior P(k = 1|w = j) and P₂ as P(k = 2|w = j), A generalised score s_i^{*} for the word j is then

$$s_{j}^{*} = s_{1}P_{1} + s_{2}P_{2}$$

= $s_{1}P_{1} + s_{2}(1 - P_{1})$
= $s_{1}P_{1} + s_{2} - s_{2}P_{1}$
= $P_{1}(s_{1} - s_{2}) + s_{2}$

Moving to the document level

The "Naive" Bayes model of a joint document-level class posterior assumes conditional independence, to multiply the word likelihoods from a "test" document, to produce:

$$P(c|d) = P(c) \prod_{j} \frac{P(w_j|c)}{P(w_j)}$$

- ► This is why we call it "naive": because it (wrongly) assumes:
 - conditional independence of word counts
 - positional independence of word counts

Naive Bayes Classification Example

(From Manning, Raghavan and Schütze, *Introduction to Information Retrieval*)

► Table 13.1	Data for parameter estimation examples.				
	docID	words in document	in $c = China$?		
training set	1	Chinese Beijing Chinese	yes		
	2	Chinese Chinese Shanghai	yes		
	3	Chinese Macao	yes		
	4	Tokyo Japan Chinese	no		
test set	5	Chinese Chinese Chinese Tokyo Japan	?		

Naive Bayes Classification Example

Example 13.1: For the example in Table 13.1, the multinomial parameters we need to classify the test document are the priors $\hat{P}(c) = 3/4$ and $\hat{P}(\overline{c}) = 1/4$ and the following conditional probabilities:

$$\begin{array}{rcl} \hat{P}({\sf Chinese}|c) &=& (5+1)/(8+6) = 6/14 = 3/7\\ \hat{P}({\sf Tokyo}|c) = \hat{P}({\sf Japan}|c) &=& (0+1)/(8+6) = 1/14\\ \hat{P}({\sf Chinese}|\overline{c}) &=& (1+1)/(3+6) = 2/9\\ \hat{P}({\sf Tokyo}|\overline{c}) = \hat{P}({\sf Japan}|\overline{c}) &=& (1+1)/(3+6) = 2/9 \end{array}$$

The denominators are (8 + 6) and (3 + 6) because the lengths of $text_c$ and $text_{\overline{c}}$ are 8 and 3, respectively, and because the constant *B* in Equation (13.7) is 6 as the vocabulary consists of six terms.

We then get:

$$\begin{split} \hat{P}(c|d_5) &\propto 3/4 \cdot (3/7)^3 \cdot 1/14 \cdot 1/14 \approx 0.0003. \\ \hat{P}(\overline{c}|d_5) &\propto 1/4 \cdot (2/9)^3 \cdot 2/9 \cdot 2/9 \approx 0.0001. \end{split}$$

Thus, the classifier assigns the test document to c = China. The reason for this classification decision is that the three occurrences of the positive indicator Chinese in d_5 outweigh the occurrences of the two negative indicators Japan and Tokyo.

Naive Bayes with continuous covariates

library(e1071) # has a normal distribution Naive Bayes

```
# Congressional Voting Records of 1984 (abstentions treated as missing)
data(HouseVotes84, package="mlbench")
model <- naiveBayes(Class ~ ., data = HouseVotes84)</pre>
```

##		Predicted	Actual	postPr.democrat	postPr.republican
##	1	republican	republican	1.029209e-07	9.999999e-01
##	2	republican	republican	5.820415e-08	9.999999e-01
##	3	republican	democrat	5.684937e-03	9.943151e-01
##	4	democrat	democrat	9.985798e-01	1.420152e-03
##	5	democrat	democrat	9.666720e-01	3.332802e-02
##	6	democrat	democrat	8.121430e-01	1.878570e-01
##	7	republican	democrat	1.751512e-04	9.998248e-01
##	8	republican	republican	8.300100e-06	9.999917e-01
##	9	republican	republican	8.277705e-08	9.999999e-01
##	10	democrat	democrat	1.000000e+00	5.029425e-11

Overall prediction performance

```
# now all of them: this is the resubstitution error
(mytable <- table(predict(model, HouseVotes84[,-1]), HouseVotes84$Class))</pre>
##
##
                democrat republican
                     238
                                13
##
     democrat
##
     republican
                29
                                155
prop.table(mytable, margin=1)
##
##
                  democrat republican
     democrat 0.94820717 0.05179283
##
##
     republican 0.15760870 0.84239130
```

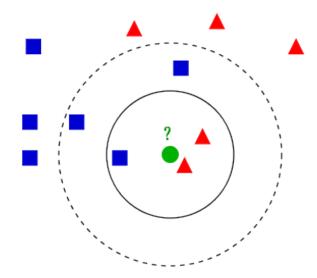
With Laplace smoothing

```
model <- naiveBayes(Class ~ ., data = HouseVotes84, laplace = 3)</pre>
(mytable <- table(predict(model, HouseVotes84[,-1]), HouseVotes84$Class))</pre>
##
##
                democrat republican
##
    democrat
                     237
                                12
    republican
               30 156
##
prop.table(mytable, margin=1)
##
##
                  democrat republican
    democrat 0.95180723 0.04819277
##
    republican 0.16129032 0.83870968
##
```

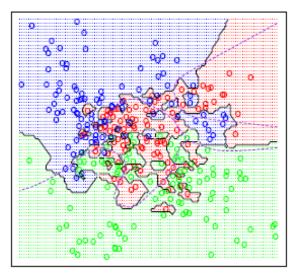
k-nearest neighbour

- A non-parametric method for classifying objects based on the training examples taht are *closest* in the feature space
- A type of instance-based learning, or "lazy learning" where the function is only approximated locally and all computation is deferred until classification
- An object is classified by a majority vote of its neighbors, with the object being assigned to the class most common amongst its k nearest neighbors (where k is a positive integer, usually small)
- Extremely *simple*: the only parameter that adjusts is k (number of neighbors to be used) - increasing k smooths the decision boundary

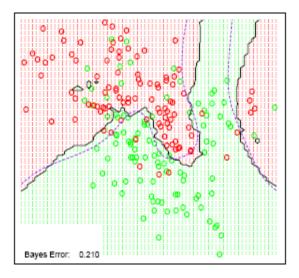
k-NN Example: Red or Blue?



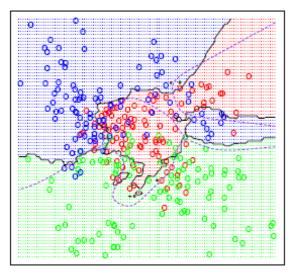
k = 1



k = 7



k = 15



```
## kNN classification
require(class)
## Loading required package: class
require(quantedaData)
## Loading required package:
                              auantedaData
## Loading required package:
                              quanteda
data(amicusCorpus)
# create a matrix of documents and features
amicusDfm <- dfm(amicusCorpus, ignoredFeatures=stopwords("english"),</pre>
                 stem=TRUE, verbose=FALSE)
## note: using english builtin stopwords, but beware that one size may not fit
# threshold-based feature selection
amicusDfm <- trim(amicusDfm, minCount=10, minDoc=20)</pre>
## Features occurring less than 10 times: 9920
```

```
## Features occurring in fewer than 20 documents: 11381
```

```
# tf-idf weighting
amicusDfm <- weight(amicusDfm, "tfidf")
# partition the training and test sets
train <- amicusDfm[!is.na(docvars(amicusCorpus, "trainclass")),]
test <- amicusDfm[!is.na(docvars(amicusCorpus, "testclass")),]
trainclass <- docvars(amicusCorpus, "trainclass")[1:4]</pre>
```

```
# classifier with k=1
classified <- knn(train, test, trainclass, k=1)
table(classified, docvars(amicusCorpus, "testclass")[-c(1:4)])</pre>
```

classified AP AR ## P 13 6 ## R 6 73

```
# classifier with k=2
classified <- knn(train, test, trainclass, k=2)
table(classified, docvars(amicusCorpus, "testclass")[-c(1:4)])</pre>
```

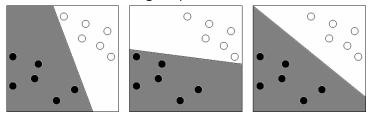
##
classified AP AR
P 9 33
R 10 46

k-nearest neighbour issues: Dimensionality

- Distance usually relates to all the attributes and assumes all of them have the same effects on distance
- Misclassification may results from attributes not confirming to this assumption (sometimes called the "curse of dimensionality") – solution is to reduce the dimensions
- There are (many!) different metrics of distance

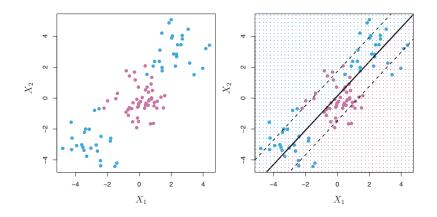
(Very) General overview to SVMs

- Generalization of maximal margin classifier
- The idea is to find the classification boundary that maximizes the distance to the marginal points



Unfortunately MMC does not apply to cases with non-linear decision boundaries

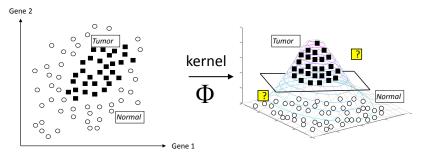
No solution to this using support vector classifier



One way to solve this problem

- Basic idea: If a problem is non-linear, don't fit a linear model
- Instead, map the problem from the *input space* to a new (higher-dimensional) *feature space*
- Mapping is done through a non-linear transformation using suitably chosen basis functions
 - the "kernel trick": using kernel functions to enable operations in the high-dimensional feature space without computing coordinates of that space, through computing inner products of all pairs of data in the feature space
 - different kernel choices will produce different results (polynomial, linear, radial basis, etc.)
- Makes it possible to then use a linear model in the feature space

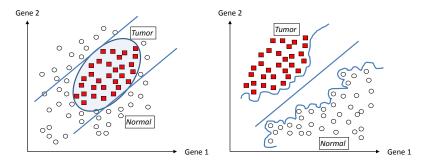
SVMs represent the data in a *higher* dimensional projection using a kernel, and bisect this using a hyperplane



Data is not linearly separable in the input space

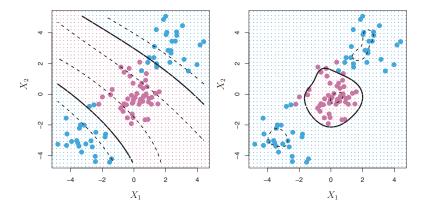
Data is linearly separable in the <u>feature space</u> obtained by a kernel

This is only needed when no linear separation plane exists - so not needed in second of these



Different "kernels" can represent different decision boundaries

- This has to do with different projections of the data into higher-dimensional space
- The mathematics of this are complicated but solveable as forms of optimization problems - but the kernel choice is a user decision



Precision and recall

Illustration framework

		True condition				
		Positive	Negative			
	Positive	True Positive	False Positive (Type I error)			
Prediction	Negative	False Negative (Type II error)	True Negative			

Precision and recall and related statistics

Precision: true positives true positives + false positives

Recall: true positives true positives + false negatives

Accuracy: Correctly classified Total number of cases

Example: Computing precision/recall

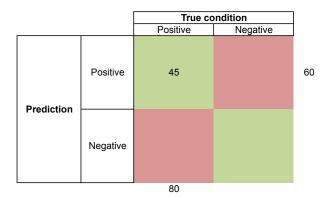
Assume:

- We have a sample in which 80 outcomes are really positive (as opposed to negative, as in sentiment)
- Our method declares that 60 are positive
- ▶ Of the 60 declared positive, 45 are actually positive

Solution:

$$\begin{aligned} \text{Precision} &= (45/(45+15)) = 45/60 = 0.75\\ \text{Recall} &= (45/(45+35)) = 45/80 = 0.56 \end{aligned}$$

Accuracy?



add in the cells we can compute

		True condition				
		Positive	Negative			
Prediction	Positive	45	15	60		
Prediction	Negative	35				

80

Receiver Operating Characteristic (ROC) plot

