#### Naive Bayes and Gaussian Bayesian Inference Thomas Schwarz

Given two events A and B, we define the conditional probability as

$$P(A \mid B) = \frac{P(A \cap B)}{P(B)}$$

"probability of A given B"

• Write also as:

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

- Bayes' Theorem: An observation of extreme importance
  - Giving rise to a new way of statistics

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

- Expresses a probability conditioned on B in one conditioned on A
- Proof:

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

• Now solve for  $P(A \mid B)$ 

 We can express a probability for one event in terms of another event happening or not

 $P(A) = P(A \cap B) + P(A \cap \overline{B})$  $= P(A \mid B)P(B) + P(A \mid \overline{B})P(\overline{B})$  $\overbrace{A \cap \overline{B}_{A \cap B}}^{A}$ 



• We can expand Bayes by calculating P(B) as probabilities conditioned on A

$$P(A \mid B) = \frac{P(B \mid A) \cdot P(A)}{P(B)}$$
$$= \frac{P(B \mid A) \cdot P(A)}{P(B \cap A) + P(B \cap \overline{A})}$$
$$= \frac{P(B \mid A) \cdot P(A)}{P(B \mid A) P(A) + P(B \mid \overline{A}) P(\overline{A})}$$

- Example: Medical Tests
  - An HIV test is positive. What is the probability that you have HIV?
  - Need some data: The quality of the test
    - Type 1 error: Test is negative, but there is illness
    - Type 2 error: Test is positive, but there is no illness

- Abbreviate probabilities
  - T : Test is positive
  - H : Person infected with HIV
  - Interested in  $P(H \mid T)$ . The quality of the test is expressed in terms of the opposite conditional probability.
    - Type I error probability:  $P(\overline{T} | H)$
    - Type II error probability:  $P(T | \overline{H})$

• We calculate

$$P(H \mid T) = \frac{P(T \mid H)P(H)}{P(T \mid H)P(H) + P(T \mid \overline{H})P(\overline{H})}$$

 Assume test has 5% type I (false positive) error probability and 1% type II (false negative) error probability:

$$P(T \,|\, \overline{H}) = 0.95$$

 $P(T \mid H) = 0.99$ 

 The probability still depends on the prevalence of HIV in the population

 $P(H \mid T) = \frac{0.99P(H)}{0.99P(H) + 0.95(1 - P(H))}$ 

- Example: HIV rate in general population in the US is 13.3/100000 = 0.000,133
- After a positive test:
  - 0.000138599 (Almost no change!)
- Example 2: HIV in a high risk group in the US is 1,753.1/100000 = 0.017531
- After a positive test:
  - 0.0182557

 The type I and type II error rates are just too bad to use this.

- Bayes' theorem inverts conditional probabilities
- Can use this for classification based on observations
- Idea: Assume we have observations  $\overrightarrow{x}$ 
  - We have calculated the probabilities of seeing these observations given a certain classification
  - I.e.: for each category, we know  $P(\vec{x}, c_i)$ 
    - Probability to observe  $\overrightarrow{x}$  assuming that point lies in  $c_i$
  - We use Bayes formula in order to calculate  $P(c_i, \vec{x})$
  - And then select the category with highest probability

- Document classification:
  - Spam detection:
    - Is email spam or ham?
  - Sentiment analysis:
    - Is a review good or bad

- Bag of words method:
  - Model a document by only counting words
    - Restrict ourselves to non-structure = non-common words

"I love this movie! It's sweet, but with satirical humor. The dialogs are great and the adventure scenes are fun. It manages to be romantic and whimsical while laughing at the conventions of the fairy tale genre. I would recommend it to just about anyone. I have seen it several times and I'm always happy to see it again"

fun	1
great	2
happy	1
humor	1
love	1
recommend	1
satirical	1
sweet	1

- There is a whole theory about recognizing key-words automatically
  - Easy out:
    - Use all words that are not common

- Recognizing words
  - Actual documents have misspelling and grammatical forms
    - Grammatical forms less common in English but typical in other languages
      - Lemmatization: Recognize the form of the word
        - जाओ, जाओगे, ... -> जाना
        - went, goes -> to go
        - Usually difficult to automatize

- Recognizing words
  - Stemming
    - Several methods to automatically extract the stem
      - English: Porter stemmer (1980)
      - Other languages: Can use similar ideas
      - https://www.emerald.com/insight/content/doi/ 10.1108/00330330610681295/full/pdf?title=theporter-stemming-algorithm-then-and-now

- Need to calculate the probability to observe a set of keywords given a classification
  - This is too specific:
    - There are too many sets of keywords
- First reduction:
  - Only use existence of words.

- Want:  $P(w_1, w_2, w_3, ..., w_n | c_i)$ 
  - The probability to find a certain word in documents of a certain category depends on the existence of other words.
    - E.g.: "Malicious Compliance"
  - We make now a big assumptions:
    - The probabilities of a keyword showing up are independent of each other
    - That's why this method is called "<u>Naïve</u> Bayes"

#### • Want:

 $P(w_1, w_2, w_3, \dots, w_n | c_i) = P(w_1 | c_i) \times P(w_2 | c_i) \times P(w_3 | c_i) \times \dots P(w_n | c_i)$ 

- Can estimate this from a training set:
  - E.g. a set of movie reviews classified with the sentiment
  - Algorithm: for document in set: sentiment = document.sentiment for word in document: count[word]+=1 if sentiment=='positive': countPos[word]+=1 else: countNeg[word]+=1 return countPos/count, countNeg/count

- This algorithm has a problem:
  - It can return a probability as zero
    - Because we use multiplication in our estimator:

 $P(w_1, w_2, w_3, \dots, w_n | c_i) = P(w_1 | c_i) \times P(w_2 | c_i) \times P(w_3 | c_i) \times \dots P(w_n | c_i)$ 

- Would create zero probabilities
- Solution: start all counts at 1
  - No more zero probabilities

• Result: Simple classifier

- Example: Use NLTK, a natural language processor
  - NLTK has several corpus (which you might have to download separately)

```
import nltk
from nltk.corpus import movie_reviews
import random
```

• First step: Get the documents

- Second step: Get all "features" (important words)
- Strategy: Get a list of all words, then order it, then select the frequent ones with exception of the most frequent ones.

```
all_words = nltk.FreqDist(w.lower() for w in movie_reviews.words())
word_features = list(all_words)[200:2000]
```

#### • Here is all\_words:

- FreqDist({',': 77717, 'the': 76529, '.': 65876, 'a': 38106, 'and': 35576, 'of': 34123, 'to': 31937, "'": 30585, 'is': 25195, 'in': 21822, ...})
- Therefore, just drop the first ones.

Create a bag of words for each document

```
def document_features(document):
    document_words = set(document)
    features = {}
    for word in word_features:
        features['contains({})'.format(word)] = (word in
    document_words)
    return features
```

```
featuresets = [(document_features(d), c) for (d,c) in documents]
train_set, test_set = featuresets[500:], featuresets[:500]
```

• Use NLTK Naive Bayes Classifier

classifier = nltk.NaiveBayesClassifier.train(train\_set)

print(nltk.classify.accuracy(classifier, test\_set))

- Results: 80.2% sentiments classified correctly
- Can see how the classifier works

>>> classifier.show\_most\_informative\_features(5)
Most Informative Features

contains(seagal)	= True	neg : pos	=	11.3 : 1.0
contains(outstanding)	= True	pos : neg	=	8.6 : 1.0
contains(wasted)	= True	neg : pos	=	7.3 : 1.0
contains(mulan)	= True	pos : neg	=	7.2 : 1.0
contains(wonderfully)	= True	pos : neg	=	6.3 : 1.0

• And already can see improvements

- Continuous features
  - Assumption: Features are distributed normally
  - Example: Look again at Iris set
    - All features are look normally distributed



- Possibility one: Disregard correlation —> Naïve
  - For each feature:
    - Calculate sample mean  $\mu$  and sample standard deviation  $\sigma$
    - Use these as estimators of the population mean and deviation
  - For a given feature value *x*, calculate the probability density assuming that *x* is in a category *c* 
    - $P(x \mid c) \sim \mathcal{N}(\mu_c, \sigma_c)$

• Estimate the probability for observation  $(x_1, x_2, ..., x_n)$  as the product of the densities

 $P((x_1, ..., x_n) | c_j) \sim \mathcal{N}(x_1, \sigma_{1,c_j}, \mu_{1,c_j}) \cdot ... \cdot \mathcal{N}(x_1, \sigma_{n,c_j}, \mu_{1,c_j})$ 

- Then use Bayes formula to invert the conditional probabilities
  - This means estimating the prevalence of the categories

• 
$$P(c_j | (x_1, ..., x_n)) = \frac{P((x_1, ..., x_n) | c_j)P(c_j)}{P((x_1, ..., x_n))}$$

- The denominator does not depend on the category  $c_i$
- So, we just leave it out:
  - $P(c_j | (x_1, ..., x_n)) \sim P((x_1, ..., x_n) | c_j) P(c_j)$
- We calculate  $P((x_1, ..., x_n) | c_j) P(c_j)$ 
  - And select the highest value

- Implemented in sklearn.naive\_bayes
  - Example with Iris data-set

from sklearn import datasets
from sklearn.naive\_bayes import GaussianNB

```
iris = datasets.load_iris()
model = GaussianNB()
model.fit(iris.data, iris.target)
print('means', model.theta_)
print('stds', model.sigma )
```

for x,t, p in zip(iris.data, iris.target, model.predict(iris.data)):
 print(x, t, p)

means [[5.006 3.428 1.462 0.246] [5.936 2.77 4.26 1.326]  $[6.588 \ 2.974 \ 5.552 \ 2.026]]$ stds [[0.121764 0.140816 0.029556 0.010884]  $[0.261104 \ 0.0965 \ 0.2164 \ 0.038324]$  $[0.396256 \ 0.101924 \ 0.298496 \ 0.073924]]$ [5.1 3.5 1.4 0.2] 0 [4.9 3. 1.4 0.2] 0 [4.7 3.2 1.3 0.2] 0 [4.6 3.1 1.5 0.2] 0 [5. 3.6 1.4 0.2] 0[5.4 3.9 1.7 0.4] 0

• There are a few errors:



 Caution: We did not divide the data set into a training and verification set.

- We did not use correlation between features
  - If we do, use the multi-variate probability density
  - Need to estimate correlation coefficients:

$$\sigma_{k,l} = \frac{1}{|C_j|} \sum_{\mathbf{x} \in C_j} (x_k - \mu_k)(x_l - \mu_l)$$

• Then use the multi-variate normal probability density  $\operatorname{norm}_{\mu,\Sigma}(x) = \frac{1}{(\sqrt{2\pi})^d \sqrt{|\Sigma|}} \exp\left(-\frac{(x-\mu)^T \Sigma^{-1} (x-\mu)}{2}\right)$ 

• Luckily, implemented in scipy.stats

from scipy.stats import multivariate\_normal

- Estimate means and correlations
- Similarly to before, estimate category by looking at the multi-variate normal density for each category and updating

```
def diagnose(tupla):
```

```
return np.argmax(
[multivariate_normal.pdf(tupla,mean=Gl.mu_setosa, cov=Gl.sigma_setosa),
multivariate_normal.pdf(tupla,mean=Gl.mu_ver, cov=Gl.sigma_ver),
multivariate_normal.pdf(tupla,mean=Gl.mu_vgc, cov=Gl.sigma_vgc)])
```

- This works slightly better: three mis-classifications
  - Example:
    - Virginica features:

>>> get\_probs((6.3, 2.8, 5.1, 1.5))
setosa 6.551299963143457e-116
versicolor 0.3895029363227387
virginica 0.25720254045708846

• Versicolor and virginica probs are similar

- This works slightly better: three mis-classifications
  - Example:
    - Versicolor features:

>>> get\_probs((6.0, 2.7, 5.1, 1.6))
setosa 3.4601607892612445e-119
versicolor 0.09776449471242309
virginica 0.56568607797792

Versicolor and virginica probs are somewhat similar