Classification via counting

Our goal in classification is to construct the conditional probability distribution p(y|x)

- given feature vector **x**, **y** is a vector of probabilities
- y_c is the probability that \mathbf{x} is in class c
 - Notation abuse alert: subscripts like c should be integers rather than class names
 - so technically should write $y_{m(c)}$ where m(c) is the integer index of class named c

This sounds difficult at first glance.

Let's start with something simpler: counting.

We will show to to construct this probability using nothing more than counting the features and targets of the training set!

From counting to probability

We introduce the topic by assuming all our variables (features and target) are discrete.

We will subsequently adapt this to continuous variables.

First, let's compute the distribution of target classes.

Let $\mathbf{X} = \{(\mathbf{x^{(i)}}, \mathbf{y^{(i)}}) | 1 \le i \le m\}$ be our m training examples.

Then

$$\mathbf{v} = \mathbf{y}' = \left| \{i \mid \mathbf{y}^{(i)} = \mathbf{y}'\} \right|$$

is the number of training examples with target y'.

We can easily convert this into an unconditional probability

$$p(\mathbf{y} = \mathbf{y}') = \frac{\text{cnt}\mathbf{y} = \mathbf{y}'}{m}$$

We can similarly compute the joint probability of any two features.

First we count the cooccurences of the two variables

$$\langle \text{cnt} \mathbf{x}_j = \mathbf{x}'_j, \mathbf{x}_k = \mathbf{x}'_k = |\{i | \mathbf{x}_j^{(i)} = \mathbf{x}'_j, \mathbf{x}_k^{(i)} = \mathbf{x}'_k\}|$$

And the the joint probability is

$$p(\mathbf{x}_j = \mathbf{x}'_j, \mathbf{x}_k = \mathbf{x}'_k) = \frac{\text{cnt}\mathbf{x}_j = \mathbf{x}'_j, \mathbf{x}_k = \mathbf{x}'_k}{m}$$

Our illustration is with two features but the notation generalizes for

- any number of variables
- any kind of variables: feature or target

Finally, we can define conditional probability

$$p(\mathbf{y} = \mathbf{y}' | \mathbf{x} = \mathbf{x}') = \frac{p(\mathbf{y} = \mathbf{y}', \mathbf{x} = \mathbf{x}')}{p(\mathbf{x} = \mathbf{x}')}$$

That is, the conditional probability

- is the joint probability
- ullet as a fraction, relative to the unconditional probability of ${f x}={f x}'$

Bayes theorem

The key for converting counts (really, associated probabilities) to predictions lies in Bayes Theorem.

Bayes Theorem relates conditional and unconditional probabilities.

Bayes Theorem

$$p(\mathbf{y} = \mathbf{y}' | \mathbf{x} = \mathbf{x}') = \frac{p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y}') * p(\mathbf{y} = \mathbf{y}')}{p(\mathbf{x} = \mathbf{x}')}$$

Let's think about Bayes Theorem in terms of our classification task:

- The left hand side is our prediction for the class probabilities, given the features
- The right hand side involves
 - the conditional probability of seeing examples with features \mathbf{x}' and target \mathbf{y}' .
 - the unconditional probability of seeing examples with label y
 - the unconditional probability of seeing examples with feature vector x.

All these elements can be obtained by counting (and filtering) the training set!

Hence, we can build an extremely simple classifier using nothing more than counting.

Posterior, Prior Probability, Evidence

Let's break down the parts of Bayes theorem and give them some names:

- p(y = y' | x = x): posterior probability
 - our prediction
 - this is the probability distribution of y conditional on the features being
 x
- p(y = y'): prior probability
 - this is the unconditional distribution of y
- $p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y})$: likelihood
 - given that y = y', what is the probability that x = x'?
 - this is the counting part: how often does the label y occur when the features are x?
- $p(\mathbf{x} = \mathbf{x}')$: evidence
 - how often do we see the features x?

We can re-state Bayes Theorem as $posterior = \frac{prior*likelihood}{evidence}$

That is:

- ullet starting from an uninformed *prior* distribution of $oldsymbol{y}$
- derive a conditional posterior distribution (i.e., informed by evidence x) by updating via the likelihood of seeing x, y together.

Proof of Bayes Theorem
$$p(\mathbf{y} = \mathbf{y}' | \mathbf{x} = \mathbf{x}') = \frac{p(\mathbf{y} = \mathbf{y}', \mathbf{x} = \mathbf{x}')}{p(\mathbf{x} = \mathbf{x}')} \qquad \text{(def. of conditional probability)}$$

$$= \frac{p(\mathbf{y} = \mathbf{y}', \mathbf{x} = \mathbf{x}')}{p(\mathbf{x} = \mathbf{x}')} * \frac{\frac{1}{p(\mathbf{y} = \mathbf{y}')}}{\frac{1}{p(\mathbf{y} = \mathbf{y}')}} \qquad \text{(multiply by identity)}$$

$$= \frac{p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y}')}{p(\mathbf{x} = \mathbf{x}')} * \frac{1}{\frac{1}{p(\mathbf{y} = \mathbf{y}')}} \qquad \text{(def. of conditional probability)}$$

$$= \frac{p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y}')}{p(\mathbf{x} = \mathbf{x}')} * p(\mathbf{y} = \mathbf{y}')$$

Length of x is n

Remember that \mathbf{x} is a vector, so that $p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y}')$ is a *joint* probability of n terms $p(\mathbf{x}_1 = \mathbf{x}'_1, \mathbf{x}_2 = \mathbf{x}'_2, \dots, \mathbf{x}_n = \mathbf{x}'_n | \mathbf{y} = \mathbf{y}')$

We an obtain this by counting (as described above)

- Let $\langle \operatorname{cnt} \mathbf{x}_j \rangle$ denote the number of distinct values for the j^{th} feature
- We need to count

$$\prod_{1 \le j \le n} \operatorname{lcnt} x_j$$

combinations

That's a lot of counting. More importantly, it's a lot of parameters to remember (i.e, size of Θ is big).

We need a short-cut.

The Naive part of Naive Bayes

We will assume that each feature is *conditionally* independent of one another

$$p(\mathbf{x}_j = \mathbf{x}'_j, \mathbf{x}_k = \mathbf{x}'_k, |\mathbf{y} = \mathbf{y}') = p(\mathbf{x}_j = \mathbf{x}'_j | \mathbf{y} = \mathbf{y}') * p(\mathbf{x}_k = \mathbf{x}'_k | \mathbf{y} = \mathbf{y}')$$

That is

- x_i an x_k are not independent
- ullet they are independent *conditional* on ${f y}={f y}'$

Think of x_j and x_k being correlated through their individual relationships with y.

Excluding that mutual dependence, they may be uncorrelated.

if we make the "naive" assumption of conditional independence

$$p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = \mathbf{y}') = \prod_{i=1}^{n} p(\mathbf{x}_i = \mathbf{x}_i' | \mathbf{y} = \mathbf{y}')$$

That is, the joint conditional probability of the vector of length n is **assumed** to be the product of the individual conditional probabilities of each element of the vector.

This is probably not true, but

- is very easy to compute
- ues few parameters
- may be close enough

What about computing the unconditional $p(\mathbf{x} = \mathbf{x}')$?

We can obtain this from conditional probabilities as well

$$p(\mathbf{x} = \mathbf{x}') = \sum_{c \in C} p(\mathbf{x} = \mathbf{x}' | \mathbf{y} = c) * p(\mathbf{y} = c)$$

That is, the unconditional probability follows from the

- ullet conditional probability given $oldsymbol{y}$
- weighted by the probability p(y) for each possible value of y

This follows from the definition of conditional probability.

What this means is that the only parameters we need to remember are

- the unconditional probabilities p(y)
 - depends on number of classes |C|
- probabilities conditional on \mathbf{y} : $p(\mathbf{x}|\mathbf{y})$
 - depends on length of x: n

Example

Here is a hypothetical trading example for equities

- there are two attributes
 - Valuation: Rich/Cheap
 - is the current stock price expensive (Rich) or inexpensive (Cheap) ?
 - Yield: High/Low
 - is the divident yield of the stock desirable (High) or undesirable (Low)?
- target: Long/Short/Neutral
 - What should our position be ?

We are given a number of examples and want a classifier to tell us

- for a new equity
 - given it's Valuation and Yield: what should our position be ?



```
In [4]: d_df = pd.read_csv("valuation_yield_action.csv")
    target_name = "Action"
    d_df
```

Out[4]:

	Valuation	Yield	Action
0	Cheap	High	Long
1	Cheap	High	Long
2	Cheap	High	Long
3	Cheap	High	Neutral
4	Rich	Low	Short
5	Rich	Low	Short
6	Rich	Low	Short
7	Rich	Low	Short
8	Rich	Low	Neutral
9	Cheap	Low	Neutral
10	Cheap	Low	Long
11	Cheap	Low	Short
12	Cheap	Low	Neutral
13	Rich	High	Long
14	Rich	High	Short
15	Rich	High	Neutral
16	Fair	Low	Long
17	Fair	Low	Short
18	Fair	Low	Neutral
19	Fair	High	Long
20	Fair	High	Short
21	Fair	High	Neutral
22	Cheap	Low	Long
23	Fair	Low	Short
24	Fair	High	Long



```
In [5]:
        grouped by target = d df.groupby(target name)
        for gp in grouped_by_target.groups.keys():
             print(gp, "\n")
            print(grouped by target.get group(gp).head())
        Long
           Valuation Yield Action
        0
               Cheap High
                             Long
        1
               Cheap High
                             Long
               Cheap High
                             Long
        10
               Cheap
                      Low
                             Long
        13
                Rich
                      High
                             Long
        Neutral
           Valuation Yield
                            Action
                            Neutral
        3
               Cheap High
        8
                Rich
                       Low
                            Neutral
        9
                            Neutral
               Cheap
                       Low
        12
               Cheap
                       Low
                            Neutral
        15
                            Neutral
                Rich
                      High
        Short
           Valuation Yield Action
```

Low Short

Low Short

Low Short

Low Short

Short

Low

4

5

6

7

11

Rich

Rich

Rich

Rich

Cheap

Looks like we

- go Long if the stock is Cheap (valuation) and High (Yield)
- go short if the stock is Rich (expensive valuation) and Low (Yield)



Out[6]:

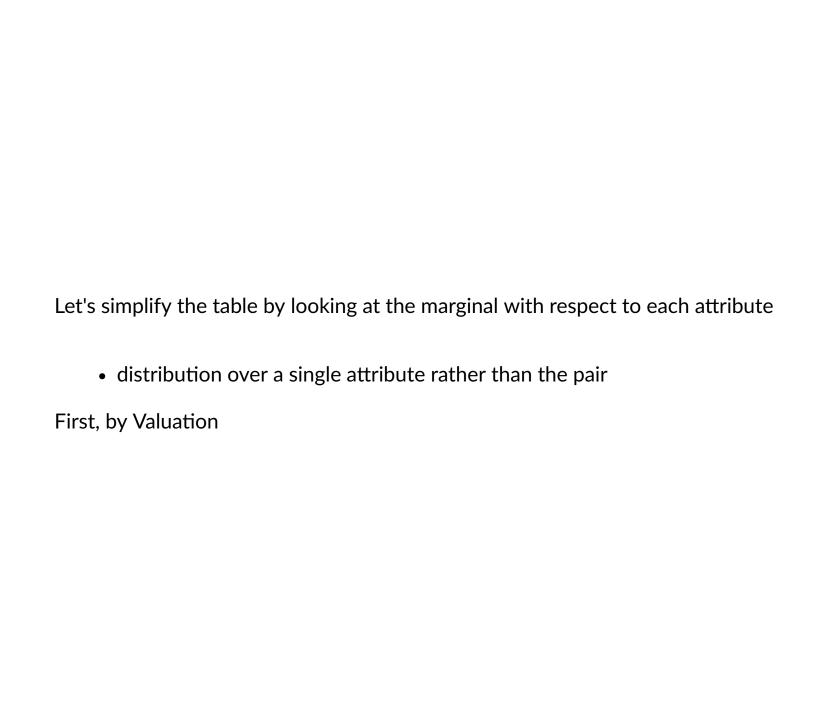
	count						
Valuation	Cheap)	Fair		Rich		All
Yield	High	Low	High	Low	High	Low	
Action							
Long	3.0	2.0	2.0	1.0	1.0	NaN	9
Neutral	1.0	2.0	1.0	1.0	1.0	1.0	7
Short	NaN	1.0	1.0	2.0	1.0	4.0	9
All	4.0	5.0	4.0	4.0	3.0	5.0	25

This gives us everything we need for the Naive Bayes algorithm

- these are counts
- we can easily turn the counts into unconditional probabilities by dividing by total number of examples
- will leave them as counts for now

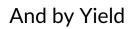
Let's parse this table:

- Columns: \cnty|x
 - a column (defined by concrete values for each of the two atttributes)
 - defines a distribution over the target (Action)
- Column Sum: $\c x = \sum_{a \in Action} \c x$
 - total number of examples with attribute pair x
- Rows: \cntx | y
 - a row (defined by a concrete value for the Action)
 - defines a distribution over the attributes pairs for which this action is taken
- Row sums: $\c = \sum_{\mathbf{x}} \c |a|$
 - total number of examples with Action a



Out[7]:

	count			
	Yield			
Valuation	Cheap	Fair	Rich	All
Action				
Long	5	3	1	9
Neutral	3	2	2	7
Short	1	3	5	9
All	9	8	8	25



Out[8]:

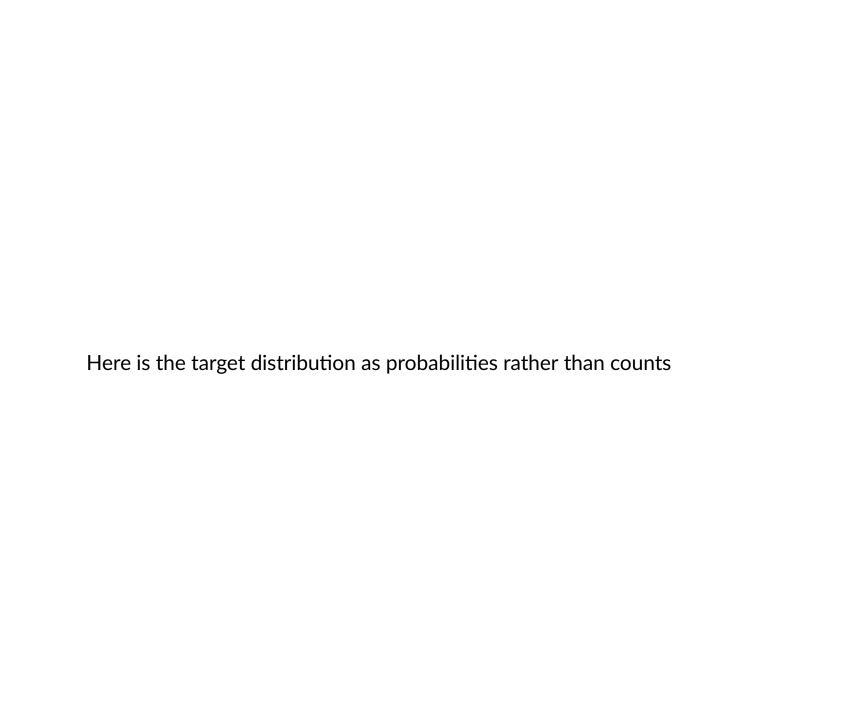
count			
	Valuation		
Yield	High	Low	All
Action			
Long	6	3	9
Neutral	3	4	7
Short	2	7	9
All	11	14	25



```
In [9]: t.loc[:, idx["count","All",:]]
```

Out[9]:

	count
Valuation	All
Yield	
Action	
Long	9
Neutral	7
Short	9
All	25



```
In [10]: num_examples = t.loc["All", idx["count","All",:]][0]
    print("There are {e:d} training examples".format(e=int(num_examples)))
# Class probabilities
    t.loc[:, idx["count","All",:]]/t.loc["All", idx["count","All",:]]
```

There are 25 training examples

Out[10]:

	count
Valuation	All
Yield	
Action	
Long	0.36
Neutral	0.28
Short	0.36
All	1.00

Why not just use the empirical distribution?

At this point, it's fair to ask:

- given a test example $\mathbf{x} = \mathbf{x}'$
- we can read p(y = y' | x = x') directly from the table

Why do we need Naive Bayes?

Answer: Because the table can be big!

one entry for every possible combination of features

The conditional independence assumption allows us

- to have a single vector for each feature \mathbf{x}_1 , \mathbf{x}_2 individually: total $|\mathbf{x}_1| + |\mathbf{x}_2|$
- rather than *combinations* of features \mathbf{x}_1 , \mathbf{x}_2 : total $|\mathbf{x}_1| * |\mathbf{x}_2|$
 - in this case n=2 but in general: total $\prod_{j=0}^{n} |\mathbf{x}_j|$

This is usually much smaller.

Drawbacks

The zero frequency problem

In order for Naive Bayes to work we must have

$$p(\mathbf{x}_j = \mathbf{x}_j' | \mathbf{y} = \mathbf{y}')$$

for all possible values of \mathbf{x}'_{j} that we will encounter during inference (test) time.

There is no guarantee that we will see each of these values in the training set.

especially when the training set is small

If we don't, the probability is 0, which is not only probably wrong but can cause problems.

Additive smoothing

additive smoothing (https://en.wikipedia.org/wiki/Additive_smoothing)

There is a simple solution to the zero frequency problem

• artificially inflate all counts by some parameter α .

This eliminates zero counts at the cost of biasing all counts.

Note that when converting counts to probabilities

- we have increased the count of each of the |C| classes by α
- so the total count for the denominator is $m + |C| * \alpha$

Replace empirical distributions by functional forms

Another way to address the zero frequency problem is to avoid the empirical distribution of training data (the counts)

- assume the features come from a paramaterized distribution
 - Bernoulli distribution for binary variables
 - Multinomial distribution for variables with more than two classes

This also has the advantage of fewer parameters: one parameter per feature.

Assumption of conditional independence

This is a questionable assumption.

In its defense:

- if *n* (the number of features) is very large
 - the conditional independence assumption is more likely to hold.

Advantages

- Very simple: just counting!
 - Easy and powerful Baseline Model to use in your Recipe for Machine Learning

Continuous variables for features

The above discussion was limited to features that could take on discrete values.

We now discuss how to include features that are continuous variables.

Discretizing continous variables

The simplest way to deal with a continuous feature \mathbf{x}_j is to turn it into one or more discrete variables.

- define a threshhold t_i and replace the continuous \mathbf{x}_i with a binary variable
 - $Is_{\mathbf{X}_j < t_j}$
- ullet define multiple intervals on the range of ${f x}_j$ and create a binary variable per interval
 - $\operatorname{Is}_{t_{j,l-1} \leq x_j \leq t_{j,l}}$
- the threshholds are a hyper-parameter: can search for optimal

Unfortunately the ordering relationship between continous values is lost

we have made them categorical

We see this same technique used in Decision Trees, so it's worth mentioning.

Replacing empirical distributions by functional forms for continous variables

Another technique for continous variables

- Replace the discrete empirical distribution by a functional form
 - Gaussian

This has the advantage that many distributions are characterized by a small number of parameters

• Gaussian: 2 per feature -- a mean and standar deviation

This also deals with the zero frequency problem by eliminating the empirical distribution.

```
In [11]: print("Done")
```

Done