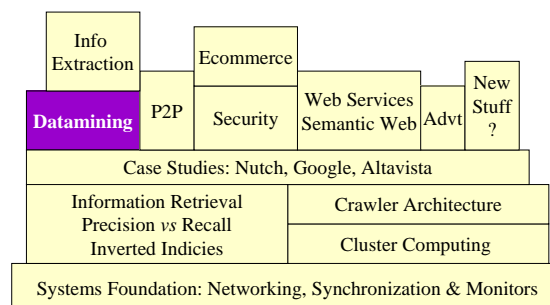


## Text Categorization (continued)

CSE 454

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## Course Overview



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## Immediate Organization

- **Tues 11/1**
  - Learning overview
  - Text categorization (Rocchio, nearest neighbor)
- **Thurs 11/3**
  - Text categorization (naïve Bayes); evaluation; topics
- **Tues 11/8**
  - Information extraction (HMMs)
- **Thurs 11/10**
  - KnowItAll (overview, rule learning, statistical model)
- **Tues 11/15**
  - KnowItAll (speedup, relational learning, opinion mining)

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## Review: Checkers as ML

- **Task T:**
  - *Playing checkers*
- **Performance Measure P:**
  - *Percent of games won against opponents*
- **Experience E:**
  - *Playing practice games against itself*
- **Target Function**
  - $V: \text{board} \rightarrow \mathcal{R}$
- **Representation of approx. of target function**

$$\hat{V}(b) = a + bx_1 + cx_2 + dx_3 + ex_4 + fx_5 + gx_6$$

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## Approximating the Target Function

- **Profound Formulation:**
  - *Can express any type of inductive learning as approximating a function*
- **E.g., Checkers**
  - $V: \text{boards} \rightarrow \text{evaluation}$
- **E.g., Handwriting recognition**
  - $V: \text{image} \rightarrow \text{word}$
- **E.g., Mushrooms**
  - $V: \text{mushroom-attributes} \rightarrow \{E, P\}$

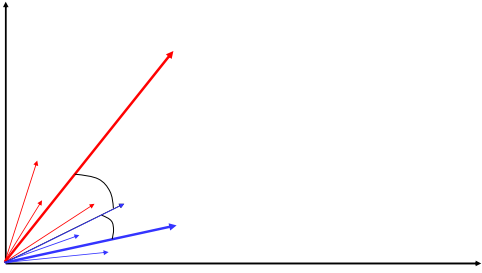
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## Supervised Learning

- **Inductive learning or “Prediction”:**
  - **Given** examples of a function  $(X, F(X))$
  - **Predict** function  $F(X)$  for new examples  $X$
- **Classification (“Categorization”)**
  - $F(X) = \text{Discrete}$
- **Regression**
  - $F(X) = \text{Continuous}$
- **Probability estimation**
  - $F(X) = \text{Probability}(X)$

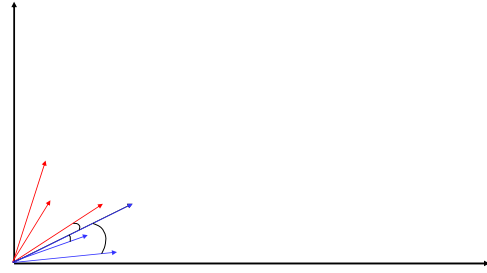
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### Illustration of Rocchio Text Categorization



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### Illustration of 3 Nearest Neighbor for Text



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### Learning ~ Prejudice meets Data

- The nice word for prejudice is “*bias*”.
- What kind of hypotheses will you consider?
  - What is allowable *range* of functions you use when approximating?
- What kind of hypotheses do you prefer?

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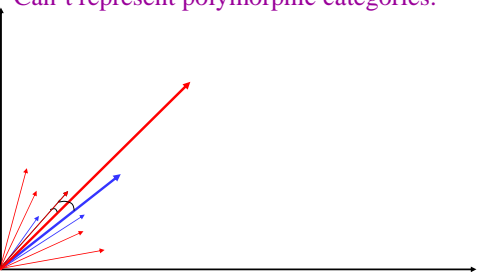
### Two Strategies for ML

- Restriction bias: use prior knowledge to specify a restricted hypothesis space.
  - Rocchio
  - Naïve Bayes
- Preference bias: use a broad hypothesis space, but impose an ordering on the hypotheses.
  - General Bayesian learning

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### Rocchio Anomaly

- Prototype models ~ very strong bias
- Can't represent polymorphic categories.



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### Bayesian Methods

- Learning and classification methods based on probability theory.
  - Bayes theorem plays a critical role in probabilistic learning and classification.
  - Uses *prior* probability of each category given no information about an item.
- Categorization produces a *posterior* probability distribution over the possible categories given a description of an item.

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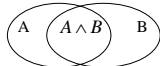
## Axioms of Probability Theory

- All probabilities between 0 and 1  
 $0 \leq P(A) \leq 1$
- True proposition has probability 1, false has probability 0.

$$P(\text{true}) = 1 \quad P(\text{false}) = 0.$$

- The probability of disjunction is:

$$P(A \vee B) = P(A) + P(B) - P(A \wedge B)$$

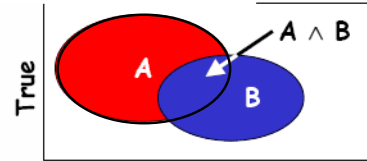


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## Probability: Simple & Logical

The definitions imply that certain logically related events must have related probabilities

$$\text{E.g. } P(A \vee B) = P(A) + P(B) - P(A \wedge B)$$



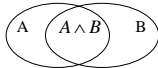
de Finetti (1931): an agent who bets according to probabilities that violate these axioms can be forced to bet so as to lose money regardless of outcome.

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## Conditional Probability

- $P(A | B)$  is the probability of  $A$  given  $B$
- Assumes:
  - $B$  is all and only information known.
- Defined by:

$$P(A | B) = \frac{P(A \wedge B)}{P(B)}$$



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## Independence

- $A$  and  $B$  are *independent* iff:

$$P(A | B) = P(A) \quad \text{These two constraints are logically equivalent}$$

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

- Therefore, if  $A$  and  $B$  are independent:

$$P(A | B) = \frac{P(A \wedge B)}{P(B)} = P(A)$$

$$P(A \wedge B) = P(A)P(B)$$

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## Bayes Theorem

$$P(H | E) = \frac{P(E | H)P(H)}{P(E)}$$

Simple proof from definition of conditional probability:

$$P(H | E) = \frac{P(H \wedge E)}{P(E)} \quad (\text{Def. cond. prob.})$$

$$P(E | H) = \frac{P(H \wedge E)}{P(H)} \quad (\text{Def. cond. prob.})$$

$$P(H \wedge E) = P(E | H)P(H) \quad (\text{Mult both sides of 2 by } P(H).)$$

$$\text{QED: } P(H | E) = \frac{P(E | H)P(H)}{P(E)} \quad (\text{Substitute 3 in 1.})$$

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## Bayesian Categorization

- Let set of categories be  $\{c_1, c_2, \dots, c_n\}$
- Let  $E$  be description of an instance.
- Determine category of  $E$  by determining for each  $c_i$

$$P(c_i | E) = \frac{P(c_i)P(E | c_i)}{P(E)}$$

- $P(E)$  can be determined since categories are complete and disjoint.

$$\sum_{i=1}^n P(c_i | E) = \sum_{i=1}^n \frac{P(c_i)P(E | c_i)}{P(E)} = 1$$

$$P(E) = \sum_{i=1}^n P(c_i)P(E | c_i)$$

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## Bayesian Categorization (cont.)

- Need to know:
  - Priors:  $P(c_i)$
  - Conditionals:  $P(E | c_i)$
- $P(c_i)$  are easily estimated from data.
  - If  $n_i$  of the examples in  $D$  are in  $c_i$ , then  $P(c_i) = n_i / |D|$
- Assume instance is a conjunction of binary features:
 
$$E = e_1 \wedge e_2 \wedge \dots \wedge e_m$$
- Too many possible instances (exponential in  $m$ ) to estimate all  $P(E | c_i)$

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## Naïve Bayesian Motivation

- Too many possible instances (exponential in  $m$ ) to estimate all  $P(E | c_i)$
- If we assume features of an instance are independent given the category ( $c_i$ ) (*conditionally independent*).

$$P(E | c_i) = P(e_1 \wedge e_2 \wedge \dots \wedge e_m | c_i) = \prod_{j=1}^m P(e_j | c_i)$$

- Therefore, we then only need to know  $P(e_j | c_i)$  for each feature and category.

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## Naïve Bayesian Categorization

- If we assume features of an instance are independent given the category ( $c_i$ ) (*conditionally independent*).

$$P(E | c_i) = P(e_1 \wedge e_2 \wedge \dots \wedge e_m | c_i) = \prod_{j=1}^m P(e_j | c_i)$$

- Therefore, we then only need to know  $P(e_j | c_i)$  for each feature and category.

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## Naïve Bayes Example

- $C = \{\text{allergy, cold, well}\}$
- $e_1 = \text{sneeze}; e_2 = \text{cough}; e_3 = \text{fever}$
- $E = \{\text{sneeze, cough, -fever}\}$

Prob	Well	Cold	Allergy
$P(c_i)$	0.9	0.05	0.05
$P(\text{sneeze} c_i)$	0.1	0.9	0.9
$P(\text{cough} c_i)$	0.1	0.8	0.7
$P(\text{fever} c_i)$	0.01	0.7	0.4

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## Naïve Bayes Example (cont.)

Probability	Well	Cold	Allergy
$P(c_i)$	0.9	0.05	0.05
$P(\text{sneeze}   c_i)$	0.1	0.9	0.9
$P(\text{cough}   c_i)$	0.1	0.8	0.7
$P(\text{fever}   c_i)$	0.01	0.7	0.4

$E = \{\text{sneeze, cough, -fever}\}$

$$P(\text{well} | E) = (0.9)(0.1)(0.1)(0.99)/P(E) = 0.0089/P(E)$$

$$P(\text{cold} | E) = (0.05)(0.9)(0.8)(0.3)/P(E) = 0.01/P(E)$$

$$P(\text{allergy} | E) = (0.05)(0.9)(0.7)(0.6)/P(E) = 0.019/P(E)$$

Most probable category: allergy  
 $P(E) = 0.089 + 0.01 + 0.019 = 0.0379$   
 $P(\text{well} | E) = 0.23$   
 $P(\text{cold} | E) = 0.26$   
 $P(\text{allergy} | E) = 0.50$

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## Evidence is Easy?

$$P(c_i | E) = \frac{\# \text{ [category]} \text{ in } E}{\# \text{ [category]} + \# \text{ [other categories]}}$$

- Or.... Are their problems?

Assume evidence is words in document

## Smooth with a Prior

$$P(c_i | E) = \frac{\# \text{ 🍌} + mp}{\# \text{ 🍌} + \# \text{ 🍌} + m}$$

$p$  = prior probability  
 $m$  = weight

Note that if  $m = 10$ , it means “I’ve seen 10 samples that make me believe  $P(X_i | S) = p$ ”

Hence,  $m$  is referred to as the **equivalent sample size**

## Estimating Probabilities

- Normally, probabilities are estimated based on observed frequencies in the training data.
- If  $D$  contains  $n_i$  examples in category  $c_i$ , and  $n_{ij}$  of these  $n_i$  examples contains feature  $e_j$ , then:

$$P(e_j | c_i) = \frac{n_{ij}}{n_i}$$

- However, estimating such probabilities from small training sets is error-prone.
- If due only to chance, a rare feature,  $e_k$ , is always false in the training data,  $\forall c_i: P(e_k | c_i) = 0$ .
- If  $e_k$  then occurs in a test example,  $E$ , the result is that  $\forall c_i: P(E | c_i) = 0$  and  $\forall c_i: P(c_i | E) = 0$

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## Smoothing

- To account for estimation from small samples, probability estimates are adjusted or *smoothed*.
- Laplace smoothing using an  $m$ -estimate assumes that each feature is given a prior probability,  $p$ , that is assumed to have been previously observed in a “virtual” sample of size  $m$ .

$$P(e_j | c_i) = \frac{n_{ij} + mp}{n_i + m} = (n_{ij} + 1) / (n_i + 2)$$

- For binary features,  $p$  is simply assumed to be 0.5.

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## Naïve Bayes for Text

- Modeled as generating a bag of words for a document in a given category by repeatedly sampling with replacement from a vocabulary  $V = \{w_1, w_2, \dots, w_m\}$  based on the probabilities  $P(w_j | c_i)$ .
- Smooth probability estimates with Laplace  $m$ -estimates assuming a uniform distribution over all words ( $p = 1/|V|$ ) and  $m = |V|$ 
  - Equivalent to a virtual sample of seeing each word in each category exactly once.

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## Text Naïve Bayes Algorithm (Train)

Let  $V$  be the vocabulary of all words in the documents in  $D$   
 For each category  $c_i \in C$   
 Let  $D_i$  be the subset of documents in  $D$  in category  $c_i$   
 $P(c_i) = |D_i| / |D|$   
 Let  $T_i$  be the concatenation of all the documents in  $D_i$   
 Let  $n_i$  be the total number of word occurrences in  $T_i$   
 For each word  $w_j \in V$   
 Let  $n_{ij}$  be the number of occurrences of  $w_j$  in  $T_i$   
 Let  $P(w_j | c_i) = (n_{ij} + 1) / (n_i + |V|)$

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## Text Naïve Bayes Algorithm (Test)

Given a test document  $X$   
 Let  $n$  be the number of word occurrences in  $X$   
 Return the category:  

$$\operatorname{argmax}_{c_i \in C} P(c_i) \prod_{i=1}^n P(a_i | c_i)$$
 where  $a_i$  is the word occurring the  $i$ th position in  $X$

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## Naïve Bayes Time Complexity

- **Training Time:**  $O(|D|L_d + |C||V|)$   
where  $L_d$  is the average length of a document in  $D$ .
  - Assumes  $V$  and all  $D_i$ ,  $n_i$ , and  $n_{ij}$  pre-computed in  $O(|D|L_d)$  time during one pass through all of the data.
  - Generally just  $O(|D|L_d)$  since usually  $|C||V| < |D|L_d$
- **Test Time:**  $O(|C|/L_t)$   
where  $L_t$  is the average length of a test document.
- Very efficient overall, linearly proportional to the time needed to just read in all the data.
- Similar to Rocchio time complexity.

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## Easy to Implement

- But...
- If you do... it probably won't work...

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## Probabilities: Important Detail!

- $P(\text{spam} | E_1 \dots E_n) = \prod_1 P(\text{spam} | E_i)$   
**Any more potential problems here?**
- We are multiplying lots of small numbers  
**Danger of underflow!**
  - $0.5^{57} = 7 \text{ E } -18$
- **Solution? Use logs and add!**
  - $p_1 * p_2 = e^{\log(p_1) + \log(p_2)}$
  - Always keep in log form

## Underflow Prevention

- Multiplying lots of probabilities, which are between 0 and 1 by definition, can result in floating-point underflow.
- Since  $\log(xy) = \log(x) + \log(y)$ , it is better to perform all computations by summing logs of probabilities rather than multiplying probabilities.
- Class with highest final un-normalized log probability score is still the most probable.

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## Naïve Bayes Posterior Probabilities

- **Classification results of naïve Bayes**
  - I.e. the class with maximum posterior probability...
  - Usually fairly accurate (?!?!?)
- **However, due to the inadequacy of the conditional independence assumption...**
  - Actual posterior-probability estimates *not* accurate.
  - Output probabilities generally very close to 0 or 1.

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## Evaluating Categorization

- Evaluation must be done on test data that are independent of the training data (usually a disjoint set of instances).
- **Classification accuracy:**  $c/n$  where
  - $n$  is the **total** number of test instances.
  - $c$  is the number of **correctly classified** test instances.
- Results can vary based on sampling error due to different training and test sets.
  - Bummer... what should we do?
- Average results over multiple training and test sets (splits of the overall data) for the best results.
  - Bummer... that means we need **lots** of labeled data...

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## N-Fold Cross-Validation

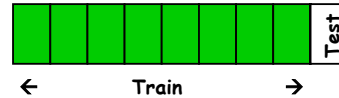
- Ideally: test, training sets are independent on each trial.
  - But this would require too much labeled data.
- Cool idea:
  - Partition data into  $N$  equal-sized disjoint segments.
  - Run  $N$  trials, each time hold back a different segment for testing
  - Train on the remaining  $N-1$  segments.
- This way, at least test-sets are independent.
- Report average classification accuracy over the  $N$  trials.
- Typically,  $N = 10$ .

Also nice to report standard deviation of averages

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## Cross validation

- Partition examples into  $k$  disjoint equiv classes
- Now create  $k$  training sets
  - Each set is union of all equiv classes *except one*
  - So each set has  $(k-1)/k$  of the original training data



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## Cross Validation

- Partition examples into  $k$  disjoint equiv classes
- Now create  $k$  training sets
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## Cross Validation

- Partition examples into  $k$  disjoint equiv classes
- Now create  $k$  training sets
  - Each set is union of all equiv classes *except one*
  - So each set has  $(k-1)/k$  of the original training data



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## Learning Curves

- In practice, labeled data is usually rare and expensive.
- Would like to know how performance varies with the number of training instances.
- *Learning curves* plot classification accuracy on independent test data ( $Y$  axis) versus number of training examples ( $X$  axis).

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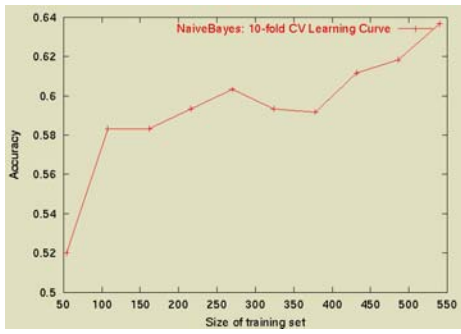
## N-Fold Learning Curves

- Want learning curves averaged over multiple trials.
- Use  $N$ -fold cross validation to generate  $N$  full training and test sets.
- For each trial,
  - train on increasing fractions of the training set
  - measure accuracy on the test data
    - for each point on the desired learning curve.

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## Sample Learning Curve (Yahoo Science Data)

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