# Active Learning and Optimized Information Gathering

Lecture 8 – Active Learning

CS 101.2 Andreas Krause

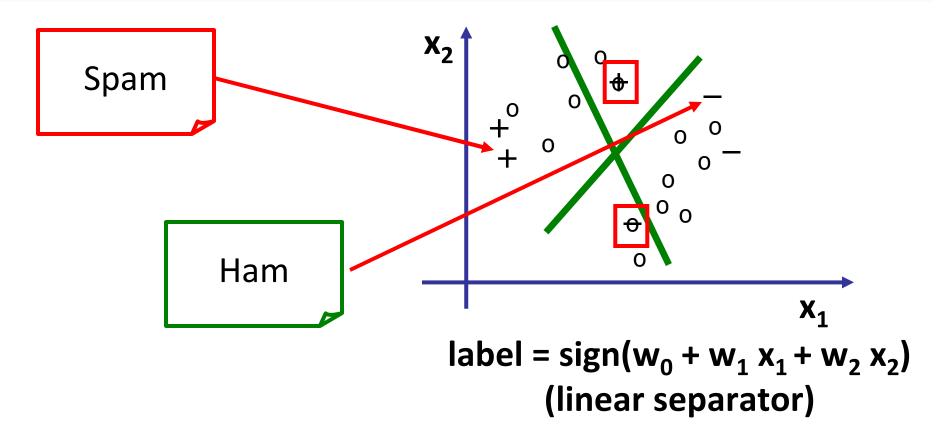
#### Announcements

- Homework 1: Due today
- Office hours
  - Come to office hours before your presentation!
  - Andreas: Monday 3pm-4:30pm, 260 Jorgensen
  - Ryan: Wednesday 4:00-6:00pm, 109 Moore

#### Outline

- Background in learning theory
- Sample complexity
- Key challenges
- Heuristics for active learning
- Principled algorithms for active learning

## Spam or Ham?



- Labels are expensive (need to ask expert)
- Which labels should we obtain to maximize classification accuracy?

## Recap: Concept learning

- Set X of instances, with distribution P<sub>X</sub>
- True concept c:  $X \rightarrow \{0,1\}$
- Data set D =  $\{(x_1, y_1), ..., (x_n, y_n)\}, x_i \sim P_X, y_i = c(x_i)$
- Hypothesis h: X  $\rightarrow$  {0,1} from H = {h<sub>1</sub>, ..., h<sub>n</sub>, ...}
- Assume  $c \in H$  (c also called "target hypothesis")
- $error_{true}(h) = E_X |c(x)-h(x)|$
- error<sub>train</sub>(h) =  $(1/n) \sum_{i} |c(x_i) h(x_i)|$

If n large enough, error<sub>true</sub>(h)  $\approx$  error<sub>train</sub>(h) for all h

#### Recap: PAC Bounds

How many samples **n** to we need to get error  $\leq \epsilon$  with probability 1- $\delta$  ?

```
No noise: n \ge 1/\epsilon (log |H| + log 1/\delta)
```

Noise: 
$$n \ge 1/\epsilon^2 (\log |H| + \log 1/\delta)$$

#### Requires that data is i.i.d.!

**Today: Mainly no-noise case (more next week)** 

#### Statistical passive/active learning protocol

Data source  $P_X$  (produces inputs  $x_i$ )



**Active learner assembles** 

data set 
$$D_n = \{(x_1, y_1), ..., (x_n, y_n)\}$$

by selectively obtaining labels



Learner outputs hypothesis h

$$\int$$

$$error_{true}(h) = E_{x^{\sim}P}[h(x) \neq c(x)]$$

Data set NOT sampled i.i.d.!!

## Example: Uncertainty sampling

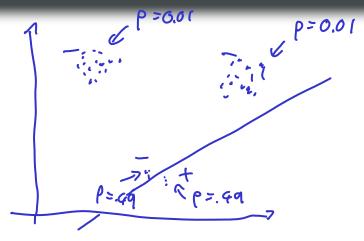
- Budget of m labels
- Draw n unlabeled examples
  \$\infty\$ = \frac{4}{2}\$
- Repeat until we've picked m labels
  - Assign each unlabeled data an "uncertainty score"
  - Greedily pick the most uncertain example

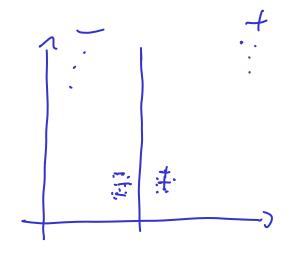
One of the most commonly used class of heuristics!

## Uncertainty sampling for linear separators



## Active learning bias





Uncertainty sampling with m= 3/2

Fren if

n > 00

m = n2

I hypothesis h

consistent with labels

ne've seen

error(h) ≥ 0.01

#### Active learning bias

If we can pick at most m = n/2 labels, with overwhelmingly high probability, US pick points such that there remains a hypothesis with error > 1!!!

• With standard passive learning, error  $\rightarrow$  0 as n $\rightarrow \infty$ 

## Wish list for active learning

- Minimum requirement
  - Consistency: Generalization error should go to 0 asymptotically
- We'd like more than that:
  - Fallback guarantee: Convergence rate of error of active learning "at least as good" as passive learning
- What we're really after
  - Rate improvement: Error of active learning decreases much faster than for passive learning

#### From passive to active

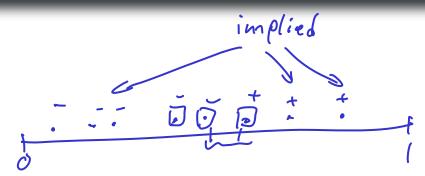
#### Passive PAC learning

- 1. Collect data set D of n  $\geq$  1/ $\epsilon$  (log |H| + log 1/ $\delta$ ) data points and their labels i.i.d. from P<sub>x</sub>
- 2. Output consistent hypothesis h
- 3. With probability at least 1- $\delta$ , error<sub>true</sub>(h)  $\leq \epsilon$

#### Key idea

- Sample **n unlabeled** data points  $D_X = \{x_1, ..., x_n\}$  i.i.d.
- Actively query labels until all hypotheses consistent with these labels agree on the labels of all unlabeled data

# Why might this work?



#### Formalization: "Relevant" hypothesis

- Data set D =  $\{(x_1,y_1),...,(x_n,y_n)\}$ , Hypothesis space H
- Input data:  $D_X = \{x_1, ..., x_n\}$
- Relevant hypothesis
  H'(D<sub>x</sub>) = H' = Restriction of H on D<sub>x</sub>
- Formally:

$$H' = \{h': D_x \rightarrow \{0,1\} \exists h \in H \text{ s.t. } \forall x \in D_x: h'(x) = h(x)\}$$

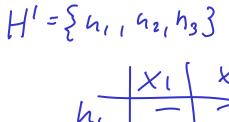
## Example: Threshold functions



$$H = \{h(x) = [x \ge t] \text{ for some } t \in [0,1]\}$$

$$h(x) > 1 \text{ if } x \ge t$$

$$0 \text{ otw.}$$



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49	1	†
hg	+	+

#### Version space

- Input data  $D_X = \{x_1, ..., x_n\}$
- Partially labeled: Have L =  $\{(x_{i_1}, y_{i_1}), ..., (x_{i_m}, y_{i_m})\}$
- The (relevant) version space is the set of all relevant hypotheses consistent with the labels L
- Formally:

- Why useful?
  - Partial labels L imply all remaining labels for  $D_x \Leftrightarrow |V|=1$

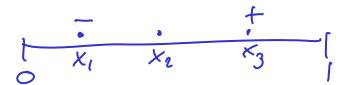
#### Version space

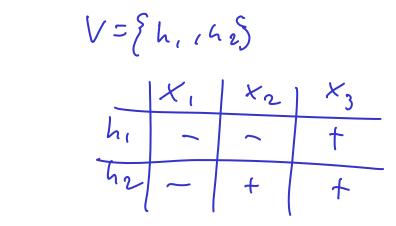
- Input data  $D_X = \{x_1, ..., x_n\}$
- Partially labeled: Have L =  $\{(x_{i_1}, y_{i_1}), ..., (x_{i_m}, y_{i_m})\}$
- The (relevant) version space is the set of all relevant hypotheses consistent with the labels L
- Formally:

$$V(D_X,L) = V = \{h' \in H'(D_X): h'(x_{i_j}) = y_{i_j} \text{ for } 1 \leq j \leq m\}$$

- Why useful?
  - Partial labels L imply all remaining labels for  $D_X \Leftrightarrow |V|=1$

## Example: Binary thresholds





#### Pool-based active learning with fallback

- 1. Collect n  $\geq$  1/ $\epsilon$  ( log |H| + log 1/ $\delta$  ) unlabeled data points D<sub>X</sub> from P<sub>X</sub>
- Actively request labels L until there remains a single hypothesis h'∈ H' that's consistent with these labels (i.e., |V(H',L)| = 1)
- 3. Output any hypothesis  $h \in H$  consistent with the obtained labels. With probability  $\geq 1-\delta$  error<sub>true</sub>(h) $\leq \epsilon$

Get PAC guarantees for active learning Bounds on #labels for fixed error  $\varepsilon$  carry over from passive to active

→ Fallback guarantee

## Wish list for active learning

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#### Pool-based active learning with fallback

- 1. Collect  $n \ge 1/\epsilon$  (log |H| + log  $1/\delta$ ) unlabeled data points  $D_X$  from  $P_X$
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## Example: Threshold functions

#### Generalizing binary search [Dasgupta '04]

- Want to shrink the version space (number of consistent hypotheses) as quickly as possible.
- General (greedy) approach:
  - For each unlabeled instance  $x_i$  compute  $v_{i,1} = V(t)'$ ,  $L \cup \xi(x_i, l)\beta)$  e "hall using the "label  $|V_{i,0}| = V(t)'$ ,  $L \cup \xi(x_i, l)\beta$ ) or  $v_{i,0} = V(t)'$
  - $v_i = \min \{v_{i,1}, v_{i,0}\}$
  - Obtain label y<sub>i</sub> for x<sub>i</sub> where i = argmax<sub>i</sub> {v<sub>i</sub>}

#### Ideal case

$$L_{o} = \frac{23}{3} CL_{i} CL_{2} C... CL_{m} | L_{mn}| = m$$

$$| \text{deal case: } \forall i \exists x : |V(L_{i} \cup \frac{2}{3}(x_{i}, 1))| = |V(L_{i} \cup \frac{2}{3}(x_{i}, 0))|$$

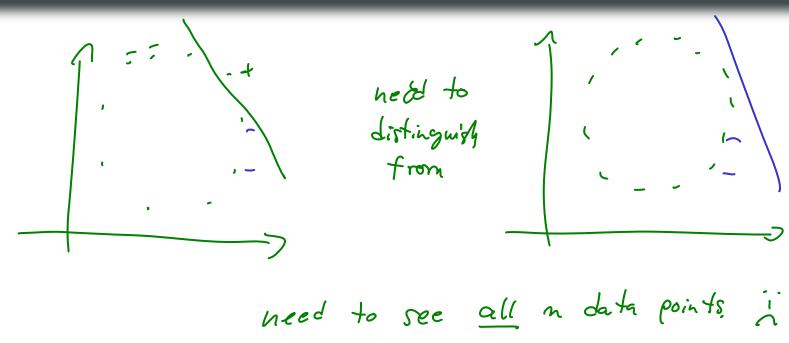
$$= \frac{1}{3} |V(L_{i+1})| \leq \frac{1}{2} |V(L_{i})|$$

$$= \frac{1}{3} |V(L_{m})| \leq \frac{1}{2} |V(L_{i})|$$

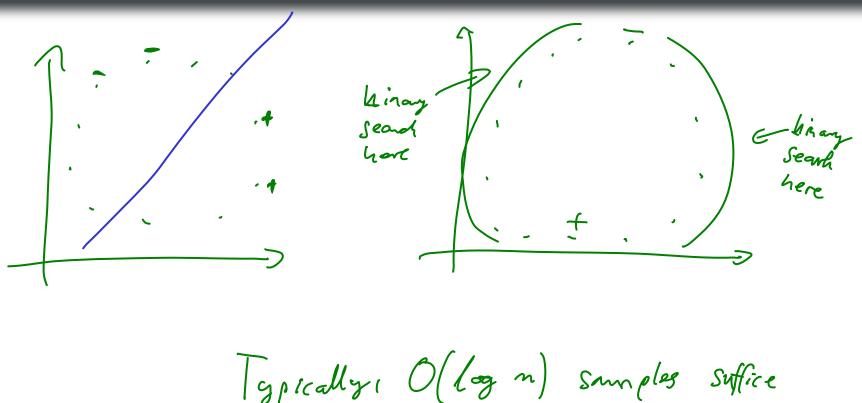
$$= \frac{1}{3} |V(L_{m})| \leq \frac{1}{3} |V(L_{i})|$$

$$= \frac{1}{3} |V(L_{m})| \leq \frac{1}{3} |V(L_{i})|$$

#### Is it always possible to half the version space?



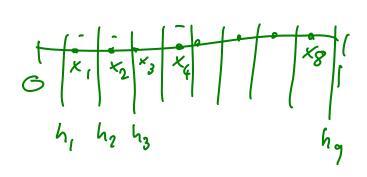
# Typical case much more benign

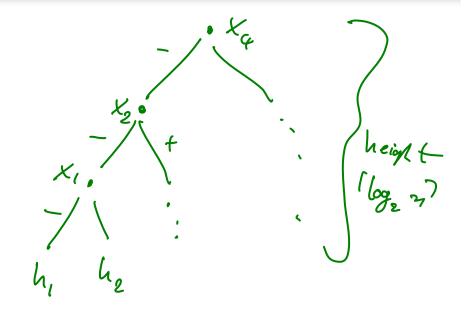


#### Query trees

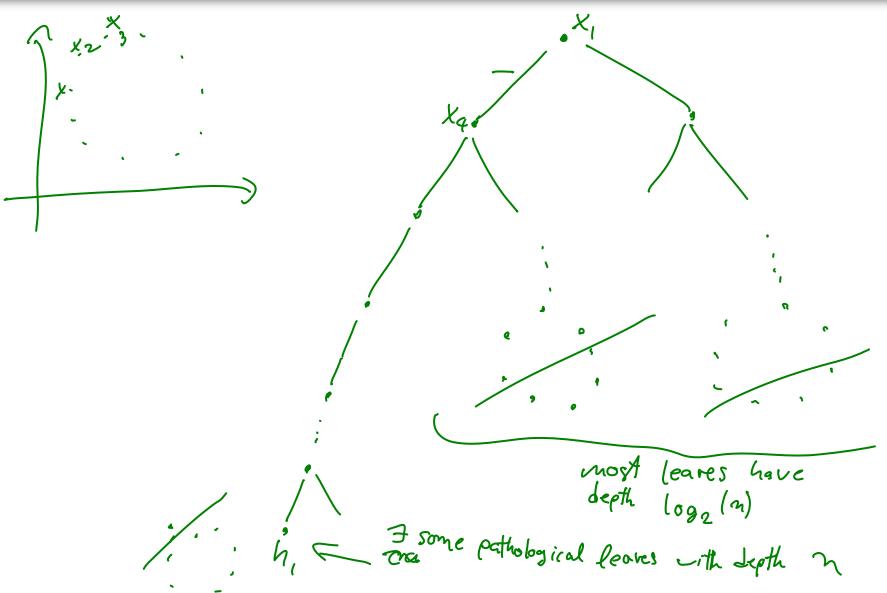
- A query tree is a rooted, labeled tree on the relevant hypothesis H'
- Each node is labeled with an input  $x \in D_X$
- Each edge is labeled with {0,1}
- Each path from root to hypothesis h'∈ H' is a labeling L such that V(D<sub>x</sub>,L) = {h'}
- Want query trees of minimum height

# Example: Threshold functions





# Example: linear separators (2D)



#### Number of labels needed to identify hypothesis

- Depends on target hypothesis!
- Binary thresholds (on n inputs D\_X)
  - Optimal query tree needs O(log n) labels! ©
- For linear separators in 2D (on n inputs D\_X)
  - For some hypotheses, even optimal tree needs n labels ⊗
  - On average, optimal query tree needs O(log n) labels!
- → Average-case analysis of active learning

#### Average case query tree learning

- Query tree T
- Cost(T) =  $1/|H'| \sum_{h' \in H'} depth(h',T)$

Want 
$$T^* = \operatorname{argmin}_T \operatorname{Cost}(T)$$

- Superexponential number of query trees < < </p>
- Finding the optimal one is hard

#### Greedy construction of query trees [Dasgupta '04]

```
Algorithm GreedyTree(D_X, L)
V' = H'(D_X)
 If V'={h}
                                                              return Leaf(h)
  Else
                                                              For each unlabeled instance x<sub>i</sub> compute
                                                             v_{i,1} = |V'(H',L \cup \{(x_i,1)\}| \text{ and } v_{i,0} = |V'(H',L \cup \{(x_i,0)\}|)
                                                        \begin{aligned} & v_i = \min \left\{ v_{i,1}, v_{i,0} \right\} & \text{max.} & \text{"Jisagreemon}A\text{"} \\ & \text{Let i = argmax}_j \left\{ v_j \right\} & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 1) \right\}) & \text{RightSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree} &= \text{GreedyTree}(D_X, L \cup \left\{ (x_i, 0) \right\}) & \text{LeftSubTree}(D_X, L \cup \left\{ (x_
                                                             return Node x<sub>i</sub> with children LeftSubTree (1) and
```

RightSubTree(0)

#### Near-optimality of greedy tree [Dasgupta '04]

**Theorem**: Let  $T^* = \operatorname{argmin}_T \operatorname{Cost}(T)$ 

Then GreedyTree constructs a query tree T such that

Cost(T) = O(log |H'|) Cost(T\*)

## Limitations of this algorithm

- Often computationally intractable
  - Finding "most-disagreeing" hypothesis is difficult
- No-noise assumption

 Will see how we can relax these assumptions in the talks next week.

#### Bayesian or not Bayesian?

- Greedy querying needs at most O(log |H'|) queries more than optimal query tree on average
- Assumes prior distribution (uniform) on hypotheses
- If our assumption is wrong, generalization bound still holds! (but might need more labels)

#### Can also do a pure Bayesian analysis:

- Query by Committee algorithm [Freund et al '97]
- Assumes that Nature draws hypotheses from known prior distribution

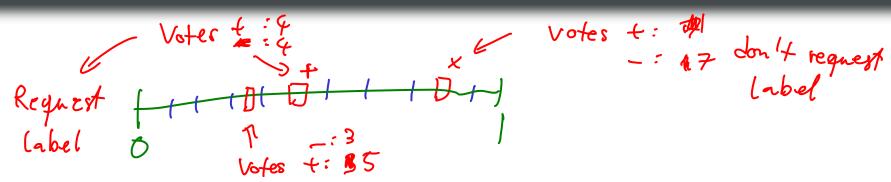
## Query by Committee

- Assume prior distribution on hypotheses
- Sample a "committee" of 2k hypotheses drawn from the prior distribution
- Search for an input such that k "members" assign label 1, and k "members" assign 0, and query that label ("maximal disagreement")

#### **Theorem** [Freund et al '97]

For linear separators in R<sup>d</sup> where both the coefficients w and the data X are drawn uniformly from the unit sphere, QBC requires exponentially fewer labels than passive learning to achieve same error

# Example: Threshold functions



## Wish list for active learning

- Minimum requirement
  - Consistency: Generalization error should go to 0 asymptotically
- We'd like more than that:
  - Fallback guarantee: Convergence rate of error of active learning "at least as good" as passive learning

#### What we're **really** after

 Rate improvement: Error of active learning decreases much faster than for passive learning

generalized binory search is "competitive" with optimal querying on average if hypotheses equally likely

## Beyond pool-based analysis

- Pool-based active learning just one convenient analysis technique
  - Gets around active learning bias by generalizing from a pool drawn i.i.d. at random
  - In pool-based analysis, there are examples where active learning does not outperform passive learning
- Exciting recent theoretical results show that using a more involved analysis, active learning always helps (asymptotically) [Balcan, Hanneke, Wortman COLT '08]
- Also other active learning paradigms
  - E.g.: Active querying (constructing rather than selecting inputs)

## What you need to know

- Uncertainty sampling
- Active learning bias
- Pool-based active learning scheme
- Relevant hypotheses and version spaces
- Generalized binary search algorithm