# Probabilistic graphical models, Assignment 2

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## 3.5, Q1:

Monotonicity of Sample Complexity: Let  $\mathcal{H}$  be a hypothesis class for a binary classification task. Suppose that  $\mathcal{H}$  is PAC learnable and its sample complexity is given by  $m_{\mathcal{H}}(\cdot, \cdot)$ . Show that  $m_{\mathcal{H}}$  is monotonically increasing in both parameters. That is, show that:

- 1. for  $\delta \in (0,1)$  for  $0 \le \epsilon_1 \le \epsilon_2 \le 1$ , show that  $m_{\mathcal{H}}(\epsilon_1,\delta) \ge m_{\mathcal{H}}(\epsilon_2,\delta)$ .
- 2. for  $\epsilon \in (0,1)$  for  $0 \le \delta_1 \le \delta_2 \le 1$ , show that  $m_{\mathcal{H}}(\epsilon, \delta_1) \ge m_{\mathcal{H}}(\epsilon, \delta_2)$ .

#### Solution, fixed $\delta$

First recall what it means to be PAC-learnable:

For every  $\epsilon, \delta \in (0,1)$ , for every distribution  $\mathcal{D}$  over  $\mathcal{X}$ , for every labelling function  $f: \mathcal{X} \to \{0,1\}$ , if the realisability assumption holds with respect to  $(\mathcal{H}, \mathcal{D}, f)$ , then for all constants  $m \geq m_{\mathcal{H}}(\epsilon, \delta)$ , on running the learning algorithm on a random i.i.d sample  $S \in X^m \sim \mathcal{D}^m$ , the algorithm produces a hypothesis  $h_S$ , such that with probability  $(1 - \delta)$ , the learning error  $L_{(\mathcal{D}, f)}(h) \leq \epsilon$ .

 $\epsilon_1$  and  $\epsilon_2$  are the error rates that we wish to achieve. Intuitively, the smaller the error we want, the more samples we need. Hence  $\epsilon_1 \leq \epsilon_2 \implies m_{\mathcal{H}}(\epsilon_1, \delta) \geq m_{\mathcal{H}}(\epsilon_2, \delta)$ .

Formally, we shall provide a **proof by contradiction**. Let us assume that:

$$\epsilon_1 \leq \epsilon_2 \implies m_{\mathcal{H}}(\epsilon_1, \delta) \leq m_{\mathcal{H}}(\epsilon_2, \delta)$$
 (assumption for contradiction)

Unwrapping the definition, this means that:

- 1 Giving the learning algorithm  $m_1 = m_{\mathcal{H}}(\epsilon_1, \delta)$  samples, the best error rate we are able to achieve is  $\epsilon_1$ ,  $(1 \delta)$  of the time.
- 2 Similarly, on giving the learning algorithm learning algorithm  $m_2 = m_{\mathcal{H}}(\epsilon_1, \delta)$  samples, the best error rate we are able to achieve is  $\epsilon_2$ ,  $(1 \delta)$  of the time.

Since the fraction  $(1 - \delta)$  is the same in both cases, we can focus purely on the error rate. Recall that  $\epsilon_1 < \epsilon_2$  and  $(m_2 > m_1)$  from our assumption of contradiction.

Hence, in the  $m_2$  case, we can simply **discard samples** and provide the learning algorithm with  $m_1$  samples, thereby reducing our error rate to  $\epsilon_1$ . However, we had assumed that for  $m_2$ ,  $\epsilon_2$  was **the best error rate we could have achieved**.

this contradiction arose from our contradictory assumption. Hence, it must be the case that as  $\epsilon$  decreases, the quantity  $m_{\mathcal{H}}(\epsilon, \cdot)$  increses.  $\square$ .

#### Solution, fixed $\epsilon$

Once again, recall what it means to be PAC-learnable:

For every  $\epsilon, \delta \in (0,1)$ , for every distribution  $\mathcal{D}$  over  $\mathcal{X}$ , for every labelling function  $f: \mathcal{X} \to \{0,1\}$ , if the realisability assumption holds with respect to  $(\mathcal{H}, \mathcal{D}, f)$ , then for all constants  $m \geq m_{\mathcal{H}}(\epsilon, \delta)$ , on running the learning algorithm on a random i.i.d sample  $S \in X^m \sim \mathcal{D}^m$ , the algorithm produces a hypothesis  $h_S$ , such that with probability  $(1 - \delta)$ , the learning error  $L_{(\mathcal{D}, f)}(h) \leq \epsilon$ .

 $\delta_1$  and  $\delta_2$  are the rejection rates that we wish to achieve. Intuitively, the smaller the number of samples we wish to reject, the more general a solution we would need to construct, thus more samples need to be seen. Hence  $\delta_1 \leq \delta_2 \implies m_{\mathcal{H}}(\epsilon, \delta_1) \geq m_{\mathcal{H}}(\epsilon, \delta_2)$ .

Formally, we once again proceed with contradiction. We assume:

$$\delta_1 \leq \delta_2 \implies m_{\mathcal{H}}(\epsilon, \delta_1) \leq m_{\mathcal{H}}(\epsilon, \delta_2)$$
 (assumption for contradiction)

Unwrapping the definition, this means that:

- 1 Giving the learning algorithm  $m_1 = m_{\mathcal{H}}(\epsilon, \delta_1)$  samples,  $(1 \delta)$  of the time, the learning error will be less than  $\epsilon$ .
- 2 Similarly, on giving the learning algorithm learning algorithm  $m_2 = m_{\mathcal{H}}(\epsilon_1, \delta)$  samples,  $(1 \delta_2)$  of the time, the learning error will be less that  $\epsilon$ .

### 3.5, Q2:

Let  $\mathcal{X}$  be a discrete domain, and let  $\mathcal{H}_{singleton} \equiv \{[z] : z \in \mathcal{X}\} \cup \{h\}$ , where

$$[z]: \mathcal{X} \to \{0, 1\}; \quad [z](x) \equiv \begin{cases} 1 & x = z \\ 0 & \text{otherwise} \end{cases}$$
  
 $h: \mathcal{X} \to \{0, 1\}; \quad h^-(\underline{\ }) \equiv 0$ 

The realizability assumption here implies that the true hypothesis f labels negatively all examples in the domain, perhaps except one.

- 1. Describe an algorithm that implements the ERM rule for learning  $\mathcal{H}_{singleton}$  in the realizable setup.
- 2. Show that  $\mathcal{H}_{singleton}$  is PAC learnable. Prove an upper bound on the sample complexity.

#### Solution, part (a)

Let  $\mathcal{X}$  be the domain, let  $f: \mathcal{X} \to \{0,1\}$  be the underlying target function f that we are trying to approximate using  $\mathcal{H}$ .

We define the sample loss  $L_S(h)$  as the number of elements in S that are mis-classified by h. More formally,  $L_S(h) \equiv |\{(x,y) \in \S : h(x) \neq y\}|$ .

The ERM algorithm must, given a particular sample set  $S \in \mathcal{X}^n \sim \mathcal{D}^n$ , provides a function  $h_0 \in \mathcal{H} = ERM(S)$  which has minimum sample loss  $L_S(h_0)$  across all functions in  $\mathcal{H}$ .

We can check over the classification of all the samples  $s \in S$ .

- If all samples  $s \in S$  are classified as 0: we return  $h^-$  this will always return 0.
- If some sample  $s_1 \in S$  is classified as 1: notice that our hypothesis space  $\mathcal{H}$  can only allow us to set at most one sample to 1. So, we can pick any sample  $s_1$  to create our hypothesis function  $h = [s_1]$ , since that is the best we can do.

```
def hminus(_): return 0 # h-: sends all samples to 0
def indicator(z): return lambda x: 1 if x == z else 0 #indicator of z
def erm_sample(S):
    # all samples which have label 1
    one_samples = [y for (x, y) in S if y == 1]
    if len(one_samples) == 0: return hminus # send all samples to 0!
    else: # we will have at least on element in one_samples
        return indicator(ones_samples[0])
```

#### Solution, part (b)

The definition of a hypothesis class  $\mathcal{H}$  to be PAC-learnable is that there exists a function  $m_{\mathcal{H}}$ :  $(0,1)^2 \to \mathbb{N}$  and a learning algorithm such that:

For every  $\epsilon, \delta \in (0,1)$ , for every distribution  $\mathcal{D}$  over  $\mathcal{X}$ , for every labelling function  $f: \mathcal{X} \to \{0,1\}$ , if the realisability assumption holds with respect to  $(\mathcal{H}, \mathcal{D}, f)$ , then for all constants  $m \geq m_{\mathcal{H}}(\epsilon, \delta)$ , on running the learning algorithm on a random i.i.d sample  $S \in X^m \sim \mathcal{D}^m$ , the algorithm produces a hypothesis  $h_S$ , such that with probability  $(1 - \delta)$ , the learning error  $L_{(\mathcal{D}, f)}(h) \leq \epsilon$ .

#### **TODO**

### 4.5, Q1:

Prove that the following two statements are equivalent for any learning algorithm A, any probability distribution  $\mathcal{D}$ , and any loss function whose loss is in the range [0,1]:

$$(1): \quad \forall \epsilon, \delta > 0, \exists M \equiv m(\epsilon, \delta), \forall m \ge M: \underset{S \sim D^m}{\mathbb{P}} [L_{\mathcal{D}}(A(S)) > \epsilon] < \delta.$$

$$(2): \quad \lim_{m \to \infty} \underset{S \sim D^m}{\mathbb{E}} [L_{\mathcal{D}}(A(S)) > \epsilon] = 0.$$

Solution:  $(2) \implies (1)$ 

We start with (2):

(2): 
$$\lim_{m \to \infty} \mathbb{E}_{S \sim \mathcal{D}^m} [L_{\mathcal{D}}(A(S)) > \epsilon] = 0.$$

We then look at the definition of the limit abstractly, for a function  $f: \mathbb{N} \to \mathbb{R}$  (In our case, the function takes  $m \in \mathbb{N}$  as input and produces  $\mathbb{E}[\dots]$  as output):

$$\lim_{m \to \infty} f(m) = y^* \equiv$$

$$\forall p > 0, \ \exists M \in \mathbb{N}, \forall m \in \mathbb{N}, m \ge M \implies |f(m) - y^*| < p$$

Plugging in to our case, we recieve:

$$\forall p > 0, \ \exists M \in \mathbb{N}, \forall m \in \mathbb{N}, m \ge M \implies \left| \underset{S \sim \mathcal{D}^m}{\mathbb{E}} [L_{\mathcal{D}}(A(S)) > \epsilon] - 0 \right| < p$$

Recall that our loss function  $L_{\mathcal{D}}$  is non-negative, hence we can remove the  $|\cdot|$  and -0 completely, giving:

$$\forall p > 0, \ \exists M \in \mathbb{N}, \forall m \in \mathbb{N}, m \geq M \implies \underset{S \in \mathcal{D}^m}{\mathbb{E}} [L_{\mathcal{D}}(A(S)) > \epsilon] < p$$

However, note that  $L_{\mathcal{D}}(A(S)) > \epsilon$  is a binary random variable, and hence:

$$\mathbb{E}_{S \sim \mathcal{D}^m}[L_{\mathcal{D}}(A(S)) > \epsilon] = 1 \cdot \mathbb{P}_{S \sim \mathcal{D}^m}[L_{\mathcal{D}}(A(S)) > \epsilon] + 0 \cdot \mathbb{P}_{S \sim \mathcal{D}^m}[L_{\mathcal{D}}(A(S)) \not> \epsilon] = \mathbb{P}_{S \sim \mathcal{D}^m}[L_{\mathcal{D}}(A(S)) > \epsilon]$$

Plugging this in, we get:

$$\forall p > 0, \ \exists M \in \mathbb{N}, \forall m \in \mathbb{N}, m \geq M \implies \underset{S \sim \mathcal{D}^m}{\mathbb{P}} [L_{\mathcal{D}}(A(S)) > \epsilon] < p$$

If we replace p with  $\delta$  in the above, we recover (1).

Hence, we show that  $(2) \implies (1)$ .  $\square$ 

Solution:  $(1) \implies (2)$ 

To show that  $(2) \implies (1)$  we only ever argued with equalities. We can run the proof backwards to derive  $(1) \implies (2)$ .  $\square$ 

### 6.8, Q1:

For two hypothesis classes  $\mathcal{H}, \mathcal{H}'$ , if  $\mathcal{H}' \subseteq \mathcal{H}$  then  $VCdim(H') \leq VCdim(H)$ .

#### Solution

Recall that the VC dimension of a given set family  $\mathcal{H}$  is the size of the largest set C such that H shatters C. That is, the intersection of C with every element is H is equal to the powerset of C:

$$\operatorname{VCdim}(\mathcal{H}) \equiv \max_{C} \{h \cap C : h \in \mathcal{H}\} = 2^{C}$$
 We denote powerset of C by  $2^{C}$ 

Now, if a set family  $\mathcal{H}'$  is a subset of another set family  $\mathcal{H}$ , and if  $\mathcal{H}'$  shatters C, then:

$$\mathcal{H}'$$
 shatters  $C \equiv \{h \cap C : h \in \mathcal{H}'\} = 2^C$  Given, (1)  
 $\{h \cap C : h \in \mathcal{H}'\} \subseteq \{h \cap C : h \in \mathcal{H}\}$  Since  $H' \subseteq H$   
 $2^C \subseteq \{h \cap C : h \in \mathcal{H}\}$  From (1)

Hence, any set that can be shattered by H' can be shattered by H if  $H' \subseteq H \implies \operatorname{VCdim}(H') \leq \operatorname{VCdim}(H)$ .

On the other hand, clearly if H is larger than H', then H can shatter more. For example, let  $H' = \emptyset \subsetneq H$ . Then H' can only shatter the empty set, while H can in general shatter sets larger than the empty set. Hence, we have have strict inequality:  $\operatorname{VCdim}(\emptyset) < \operatorname{VCdim}(H)$  for example.