6.842 Randomness and Computation

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Lecture 17

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The goal of today is to study boosting, which is a general technique that turns a distribution-free weak learner into a strong learner.

Definition 1 An algorithm A weakly PAC learns a concept class C (with confidence $\delta > 0$ and error $\gamma > 0$) if for all $f \in C$ and for all distribution D, with probability at least $1 - \delta$ and given samples from f, A outputs some c satisfying

 $\Pr_{\mathcal{D}}[f(x) \neq c(x)] \le \frac{1}{2} - \frac{\gamma}{2} .$

We emphasize here that the weak learning algorithm A does not know the distribution \mathcal{D} . It only sees samples from \mathcal{D} . The time and sampling complexity of A usually depends on δ and γ . The definition of weak learner is different from the normal but strong version of PAC learning in terms of the error guarantee: the strong version requires that for any $\varepsilon > 0$, the error $\Pr_{\mathcal{D}}[f(x) \neq c(x)]$ can be made as small as ε .

(Recall that we have learned from the previous lecture on how to obtain a weak leaner for monotone functions, but only when the distribution is uniform. If that weak leaner worked for all distributions (i.e., is distribution-free), it would imply that monotone functions can be PAC learned in the strong sense, contradicting to the impossibility result.)

1 Thought Experiments

We begin with some intuitive trials on how to turn a (distribution-free) weak leaner into a strong one.

The first (and very bad) trial is to run the weak leaner multiple times on the same distribution \mathcal{D} , and output for instance the majority of the predictions. This approach fails very miserably because typically in learning, repetition can only improve the confidence δ (see for instance Homework 7-3), but not the prediction accuracy. A simple example to illustrate this phenomenon is to consider some weak learner that always outputs the same prediction c no matter how many times it is run. In such a case, the majority of the same function c is still c itself, and therefore the prediction accuracy is not improved at all.

The second (and very promising) trial is to always run a weak learner on the samples where the previous learners fail to predict. More specifically, suppose that from distribution \mathcal{D} the weak learner A obtains some prediction $c_1(x)$ (from samples $(x_1, f(x_1)), (x_2, f(x_2)), \ldots, (x_m, f(x_m))$ if m samples are needed in A). Next, we query \mathcal{D} for more examples $(x_{m+1}, f(x_{m+1})), \ldots$, but only focus on those x_i such that $c_1(x_i) \neq f(x_i)$. In other words, we use $c_1(x)$ to filter out the successfully predicted samples, and only focus on the rest. This gives rise to a different distribution \mathcal{D}_1 , and we can pass it to the weak learner A to produce a second prediction $c_2(x)$. This process can go on for a number of stages.

The second trial looks very promising, because it uses the distribution-free feature of the weak leaner A, and always refines on the samples that are mistakenly predicted. However, a fundamental problem exists: how do we, in the end of the algorithm, combine the sequence of predictions $c_1(x), c_2(x), \ldots$? For instance, given some new sample x and suppose we want to predict the unknown label f(x), how do we choose from the answers of $c_1(x), c_2(x), \ldots$? The filtering techniques allows us to choose the $c_j(x)$ among all possible j only after seeing the ground-truth label f(x), so for new and to-be-predicted samples, the filtering no longer helps.

In this lecture, we are going to study the *boosting* algorithm that is essentially built from the second trial above. In fact, it chooses the majority of $c_1(x), c_2(x), \ldots$ as the final output (of the strong leaner), but changes the definition of filtering to ensure the correctness.

(One may ask if there are indeed example of concept classes where it is easy to design weaker learners but hard to design for strong ones. In fact, it is not clear if such examples exist so the final theorem of the boosting algorithm turns out to be a theoretical result. However, the strong learner of DNF was indeed introduced using boosting by Jackson: he uses weak learners that are not distribution-free, but work for a sufficiently large class of distributions that is sufficient for the boosting of DNF. Also, variants of boosting have found numerous applications in practical problems such as character recognitions.)

2 The Boosting Framework

Pick δ to be smaller than 1/T where T is the number of the phases, and suppose we are given samples from \mathcal{D} that are labelled according to f.

- Stage 0: initialization. $\mathcal{D}_0 \leftarrow \mathcal{D}$. Run the weak learner on \mathcal{D}_0 to generate $c_1(x)$ satisfying $\Pr_{\mathcal{D}_0}[f(x) = c_1(x)] \geq \frac{1}{2} + \frac{\gamma}{2}$.
- For stage $i \leftarrow 1$ to $T = O(1/\gamma^2 \varepsilon^2)$.
 - Stop the algorithm if $Maj(c_1, c_2, \ldots, c_i)$ is correct on 1ε fraction of the inputs with respect to \mathcal{D} , and output the function $Maj(x_1, \ldots, c_i)$ as the final prediction.

 (This step can be done by taking $O(1/\varepsilon)$ samples from \mathcal{D} and checking how many of them fail the test on $f(x) = Maj(c_1(x), \ldots, c_i(x))$.)
 - Construct \mathcal{D}_i via some "filtering procedure". (This will be introduced in the next section, but from a high level, \mathcal{D}_i is constructed from \mathcal{D} by favoring samples where the previous predictions are incorrect. It will be constructed in a probabilistic way based on how many previous predictions are correct.)
 - Run the weak leaner with distribution \mathcal{D}_i and get a new function $c_{i+1}(x)$ satisfying $\Pr_{\mathcal{D}_i}[f(x) = c_{i+1}(x)] \geq \frac{1}{2} + \frac{\gamma}{2}$.
- If the algorithm does not stop in T stages, output $C = Maj(c_1, ..., c_T)$. (Our theorem ensures that the algorithm will never reach here, at least with reasonable probability.)

3 The Specific Choice of Filtering Procedure

At stage i of the algorithm, given predictions c_1, \ldots, c_i from stage 0 through stage i-1, we construct \mathcal{D}_i as follows. Given a sample (x, f(x)) from \mathcal{D} , we compare $Maj(c_1(x), \ldots, c_i(x))$ and f(x).

- If $Maj(c_1(x), \ldots, c_i(x)) = f(x)$, we keep this sample.
- Otherwise, letting #right be the number of correct predictions among $c_1(x), \ldots, c_i(x)$, and #wrong be the number of incorrect predictions (so we have #right + #wrong = i).

 If a large majority is correct —that is, $\#right \#wrong > \frac{1}{\gamma\varepsilon}$ (which is equivalent to $\#wrong \le \frac{i}{2} \frac{1}{2\gamma\varepsilon}$)—we discard this sample.
- If $\#right \#wrong = \frac{\alpha}{\gamma \varepsilon} < \frac{1}{\gamma \varepsilon}$ for some $\alpha \in [0, 1]$, we keep this sample with probability 1α .

This above random procedure (of generating samples) gives the definition of \mathcal{D}_i . Note that when i is small, the threshold $\frac{i}{2} - \frac{1}{2\gamma\varepsilon}$ is negative so nearly all samples from \mathcal{D} are kept, and in other words, $\mathcal{D}_i \approx \mathcal{D}$ for small i.

As another remark, in principle, we need to make sure that the weak learner receives enough samples in each stage i; or in other words, the sampling complexity does not blow up from weak to strong learner. This is true because, at stage i, the non-stopping criterion $\Pr_{\mathcal{D}}[Maj(c_1(x),\ldots,c_i(x))=f(x)]<1-\varepsilon$ ensures that we only need $\leq \frac{1}{\varepsilon}$ samples from \mathcal{D} in order to generate a sample from \mathcal{D}_i .

4 Sketched Proof of the Correctness

We mostly only introduce some notations here, and the full proof will be given in the next lecture. Let $R_c(x)$ capture the correctness of a prediction c at input x:

$$R_c(x) := \begin{cases} +1, & \text{if } f(x) = c(x); \\ -1, & \text{otherwise.} \end{cases}$$

Let $N_i(x)$ indicate "#correct - #wrong" for the first i predictions:

$$N_i(x) := \sum_{1 \le j \le i} R_{c_j}(x) .$$

Let

$$M_i(x) := \begin{cases} +1, & \text{if } N_i(x) \leq 0; \\ 0, & \text{if } N_i(x) \geq \frac{1}{\gamma \varepsilon}; \\ 1 - \varepsilon \gamma \cdot N_i(x), & \text{otherwise.} \end{cases}$$

so that the distribution $\mathcal{D}_{M_i}(x) := \frac{M_i(x)}{\sum_x M_i(x)}$ coincides with our distribution \mathcal{D}_i . We emphasize again that this distribution \mathcal{D} includes all incorrectly predicted x plus some others.

We define also the advantage of a prediction c(x) over M_i as

$$Adv_c(M_i) := \sum_{x} R_c(x) \cdot M_i(x)$$

It is clear from the definition that $\Pr_{x \in \mathcal{D}_{M_i}}[c(x) = f(x)] = \frac{1}{2} + \frac{Adv_c(M_i)}{2\sum_x M_i(x)}$. Note that, by the definition of the weak learner, we have that the special choice of $c = c_{i+1}$ makes sure that $\Pr_{x \in \mathcal{D}_{M_i}}[c(x) = f(x)] \ge \frac{1}{2} + \frac{\gamma}{2}$, and therefore $\frac{Adv_c(M_i)}{\sum_x M_i(x)} \ge \gamma$ for $c = c_{i+1}$.

(We do not have enough time to continue the proof today, so the following texts provide a sketch of the proof.)

We should always have $\sum_x M_i(x) \ge \varepsilon 2^n$ if the algorithm does not stop. This gives a lower bound on $\sum_x R_{c_{i+1}}(x) \cdot M_i(x) = Adv_{c_{i+1}}(M_i) \ge \gamma \varepsilon 2^n$.

On the other hand, for any fixed input x, letting $A_i(x) := \sum_{0 \le j \le i-1} R_{c_{j+1}}(x) \cdot M_j(x)$, we should have $A_i(x) < \frac{1}{x_i} + \frac{\varepsilon \gamma}{2} \cdot i$.

have $A_i(x) \leq \frac{1}{\varepsilon\gamma} + \frac{\varepsilon\gamma}{2} \cdot i$. At last, we combine the upper and lower bounds, deduce a contradiction and therefore proving that the algorithm must terminate in T stages.

¹This is because, if $R_{c_{i+1}}(x)$ keeps being large for a few iterations j, then $M_i(x)$ drops very quickly by the definition.