Note: More Efficient Conversion of Equivalence-Query Algorithms to PAC Algorithms

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May 8th, 2008

Abstract

We present a method for transforming an Equivalence-query algorithm using Q queries into a PAC-algorithm using $\frac{Q}{\epsilon} + O(\frac{Q^{2/3}}{\epsilon} \log \frac{Q}{\delta})$ examples in expectation. The method is a variation of that by Schuurmans and Greiner which provides, for each $\gamma > 0$, an algorithm using $(1 + \gamma)\frac{Q}{\epsilon} + O(\frac{1}{\epsilon} \log \frac{Q}{\delta})$ examples in expectation. In other words, we show that the constant in front of the dominating term Q/ϵ can be made 1 + o(1).

1 Introduction

In her seminal paper on learning from queries, Angluin [Ang87] showed that algorithms using Equivalence queries can be rewritten as PAC algorithms. Her simulation uses a worst-case sample $O(\frac{Q^2}{\epsilon} \ln \frac{1}{\delta})$ to achieve (ϵ, δ) confidence from an algorithm using Q Equivalence queries, but it is not difficult to show that in her same simulation, sample size $O(\frac{Q}{\epsilon} \ln \frac{Q}{\delta})$ suffices.

^{*}Partially supported by the EU PASCAL2 Network of Excellence, and by the Spanish Ministry of Education through the MOISES-BAR project, TIN2005-08832-C03-03. gavalda@lsi.upc.edu, http://www.lsi.upc.edu/~gavalda.

It was shown later that, with a different algorithm, that the dependence on n can be made linear. Specifically, Littlestone [Lit89] showed that there is a simulation using a worst-case sample size $4\frac{Q}{\epsilon} + O(\frac{1}{\epsilon} \ln \frac{Q}{\delta})$) (his simulation was phrased in terms of on-line learning rather than Equivalence queries, but the distinction is irrelevant for our purpose). Schuurmans and Greiner [SG95, Sch96] showed how to build, for every constant $\gamma > 0$, a simulation that uses *expected* sample size $(1 + \gamma)\frac{Q}{\epsilon} + c(\gamma)\frac{1}{\epsilon} \ln \frac{Q}{\delta}$. Here $c(\gamma)$ is constant for each γ , but tends to infinity as γ tends to 0.

In this note we show that the leading constant in front of the Q/ϵ term can be made 1 + o(1), that is, arbitrarily close to 1 as Q grows. In fact, our algorithm is essentially the same as the Schuurmans-Greiner one, except that instead of using a fixed value for γ a priori, we let the value of γ decrease at a precisely controlled rate as the algorithm progresses.

2 The Algorithm

We view an Equivalence query algorithm as a particular case of a strategy for generating hypothesis from sequences of labelled examples. Given such an algorithm, we build a new algorithm S, given in Figure 1, which reads a sequence of example, uses the Equivalence-query strategy as a black box, and eventually outputs a hypothesis from those generated by the strategy. We will show that S is a PAC-learning algorithm.

Procedure **sprt** is Wald's Sequential Probability Ratio Test, discussed below, and also used in the Schuurmans-Greiner approach. The main difference with their method is that we do not fix a constant γ *a priori*, but rather use a different γ_i that varies with *i*. We will fix one particular setting for the sequence of γ_i to obtain our bound on the sample size used by **S**, but occasionally comment on the effect of using other values for γ_i .

We will argue that procedure **S** satisfies three conditions, which we formulate as theorems: Correctness, Completeness, and Efficiency.

Theorem 1 (Correctness) The probability that $S(\epsilon, \delta)$ outputs some $h \in H$ with $error(h) > \epsilon$ is less than δ .

The completeness condition can be stated in many ways, of which the following is but one example:

Theorem 2 (Completeness) If for some *i* we have that $error(h_i) = 0$ with probability 1, then $S(\epsilon, \delta)$ stops with probability 1.

Algorithm $S(\epsilon, \delta)$

Generate initial hypothesis h_1 ; 1 2 i := 1; t := 0;3 while TRUE 4 do 5t := t + 1;6get a training example $(x_t, c(x_t))$, labelled by the unknown target c; 7if $h_i(x_t) \neq c(x_t)$ (i.e., $(x_t, c(x_t))$) is a counterexample for h_i) 8 then 9 use (x_t, y_t) to generate h_{i+1} ; 10 start testing $error(h_i)$ on subsequent examples using $\operatorname{sprt}(\epsilon/(1+\gamma_i), \epsilon, \delta/(i(i+1)), 0);$ 11 12i := i + 1;if for some j < i, the sprt test for h_j rejects 1314then 15drop h_j from the list of hypothesis being tested if for some j < i, the sprt test for h_j accepts 1617then 18 output h_j and stop 19end while

Figure 1: Algorithm S

Putting both claims together, if the strategy used to generate hypothesis is an exact Equivalence-query algorithm learning with finitely many queries, with probability 1 the algorithm stops, and its output is, with probability $1 - \delta$, a hypothesis h having $error(h) < \epsilon$.

Theorem 2 in fact follows from this more general statement:

Theorem 3 (Running time) Define $\gamma_i = i^{-1/3}$, and let the base Equivalencequery learner learn with at most Q queries. Then

$$E[running \ time \ of \ \mathbf{S}(\epsilon, \delta)] \leq \frac{Q}{\epsilon} + 7 \frac{Q^{2/3}}{\epsilon} \cdot (\ln \frac{Q(Q+1)}{\delta} + 2)$$

We do not describe here the **sprt** test. We quote, however, some relevant properties from [Sch96], appendix A:

Theorem 4 [Sch96] Let k > 1 and suppose $sprt(\epsilon/k, \epsilon, \delta_{acc}, \delta_{rej})$ is run on a sequence $X_1, X_2, \ldots, X_i, \ldots$ of i.i.d. boolean random variables. Then:

- 1. If $E[X_i] > \epsilon$, the probability that sprt accepts is at most δ_{acc} .
- 2. If $E[X_i] < \epsilon/k$, the probability that sprt rejects is at most δ_{rej} .
- 3. ([Sch96], Lemma A.4) If $\delta_{rej} = 0$, the expected running time of sprt is

$$\left(\frac{k}{k-1-\ln k}\right) \frac{1}{\epsilon} \left(\ln \frac{1}{\delta_{acc}} + 1\right).$$

3 Proof of Theorem 1

The proof is as in [SG95, Sch96], but we reproduce it for completeness. We say that a hypothesis $h \in H$ is ϵ -bad iff $error(h) \geq \epsilon$. Observe that the **sprt** instance associated to h_i is fed boolean variables whose expected value is precisely $error(h_i)$. Therefore, by Theorem 4, part (1), we have the following (where probabilities are taken over infinite sequences of independently generated examples).

$$\Pr[\mathbf{S}(\epsilon, \delta) \text{ outputs an } \epsilon\text{-bad hypothesis}] \\ \leq \sum_{i=1}^{\infty} \Pr[h_i \text{ is } \epsilon\text{-bad yet } \mathbf{S}(\epsilon, \delta) \text{ outputs } h_i]$$

$$\leq \sum_{i=1}^{\infty} \Pr[\operatorname{sprt}(\epsilon/(1+\gamma_i), \epsilon, \delta/(i(i+1)), 0) \text{ accepts } h_i \mid h_i \text{ is } \epsilon\text{-bad}]$$

$$\leq \sum_{i=1}^{\infty} \frac{\delta}{i(i+1)} = \delta.$$

4 Proof of Theorem 3

For every i, we define the following random variables and expected values:

- h_i is the random variable representing the *i*th generated hypothesis,
- ϵ_i is such that $1/\epsilon_i = E[1/error(h_i)],$
- T_i is the number of examples read from the moment in which h_i is generated until either h_{i+1} is generated (if h_{i+1} is ever generated; otherwise, $T_i = \infty$)
- R_i is the running time of the **sprt** test run on h_i , and
- T is the running time of the algorithm.

Proving Theorem 3 is thus bounding E[T]. Let *i* be the first index such that $\epsilon_i(1 + \gamma_i) < \epsilon$. Note that if the base Equivalence learner uses at most Q queries, we have $i \leq Q$. Observe also that

$$T \le \sum_{j < i} T_j + R_i \tag{1}$$

because, by definition of T_j and R_i , by this time h_i has been generated and the **sprt** test for h_i has stopped. Since the test is run with parameter δ_{rej} , it rejects h_i with probability 0, i.e., it accepts h_i . Therefore, by this time either **S** stops outputting h_i , unless it has stopped before due to another h_j .

Taking expectations in Equation (1), we have

$$E[T] \le \sum_{j < i} E[T_j] + E[R_i].$$

$$\tag{2}$$

We first bound $E[T_i]$; the proof of the lemma is given later.

Lemma 1 $E[T_j] = 1/\epsilon_j$.

Taking $k = (1 + \gamma_i)$ in Theorem 4, part (3), provides the following bound on $E[R_i]$:

$$E[R_i] \le \frac{1+\gamma_i}{\gamma_i - \ln(1+\gamma_i)} \frac{1}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 1\right).$$
(3)

As a detour, let us note how to get the result in [SG95, Sch96]. Since *i* is the first index such that $\epsilon_i(1 + \gamma_i) < \epsilon$, for j < i we have $\epsilon_j \ge \epsilon/(1 + \gamma_j)$, that is, $E[T_j] = 1/\epsilon_j \le (1 + \gamma_j)/\epsilon$. Fix $\gamma_i = \gamma$ for every *i*. Then from Equation (2) we get

$$E[T] \leq \sum_{j < i} \frac{1+\gamma}{\epsilon} + \frac{1+\gamma}{\gamma - \ln(1+\gamma)} \frac{1}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 1\right)$$
$$= (1+\gamma)\frac{i}{\epsilon} + c(\gamma) \frac{1}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 1\right).$$

Now, take take instead $\gamma_i = i^{-1/3}$. We have the following two lemmas, whose proofs are given later:

Lemma 2 For $\gamma_j = j^{-1/3}$,

$$\sum_{j < i} (1 + \gamma_j) \le i + \frac{3}{2} i^{2/3}.$$

Lemma 3 Define $c(\gamma) = (1 + \gamma)/(\gamma - \ln(1 + \gamma))$. Then $c(\gamma) \leq 7/\gamma^2$ for every $\gamma \in (0, 1]$, and $c(\gamma)$ tends to $2/\gamma^2$ as γ tends to 0.

From Equations (2) and (3) and Lemmas 2 and 3, and using again that for all j < i we have $E[T_j] = 1/\epsilon_j \leq (1 + \gamma_j)/\epsilon$, we obtain

$$E[T] \leq \sum_{j < i} \frac{1 + \gamma_j}{\epsilon} + \frac{7}{\gamma_i^2} \frac{1}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 1 \right)$$

$$\leq \frac{1}{\epsilon} \left(i + \frac{3}{2} i^{2/3} \right) + 7 \frac{i^{2/3}}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 1 \right)$$

$$\leq \frac{i}{\epsilon} + 7 \frac{i^{2/3}}{\epsilon} \left(\ln \frac{i(i+1)}{\delta} + 2 \right)$$

i.e., the statement of Theorem 3.

Proof of Lemma 1. Suppose that in a particular run of the algorithm the random variable h_j takes a particular value $h \in H$. Conditioned to $h_j = h$, the expected number of examples that have to be read to produce a counterexample for h_j is an exponential distribution with base error(h), and therefore,

$$E[T_j|h_j = h] = \sum_{\ell=1}^{\infty} (1 - error(h))^{\ell-1} \cdot error(h) \cdot \ell = 1/error(h).$$

So $E[T_j] = E[1/error(h_j)]$ (where the expectation is taken over h on the right-hand side), which is $1/\epsilon_j$ by definition of ϵ_j . \blacksquare (Lemma 1)

Proof of Lemma 2. We show by induction on i the following inequality, which implies the lemma:

$$\sum_{j \le i} (1+j^{-1/3}) \le \frac{i}{\epsilon} + \frac{3}{2} \frac{i^{2/3}}{\epsilon}.$$

For i = 1 it is obvious. Assume true for i, then

$$\sum_{j=1}^{i+1} j^{-1/3} \le \frac{3}{2} i^{2/3} + (i+1)^{-1/3}$$

and observe that

$$\frac{3}{2}i^{2/3} + (i+1)^{-1/3} \le \frac{3}{2}(i+1)^{2/3}$$

iff (multiplying on both sides by $(i+1)^{1/3}$)

$$\frac{3}{2} \left(i^2 (i+1) \right)^{1/3} + 1 \le \frac{3}{2} \left(i+1 \right)$$

iff (taking cubes on both sides)

$$\left(\frac{3}{2}\right)^3 (i^2(i+1)) \le \left(\frac{3}{2}(i+1) - 1\right)^3$$

which is verified to be true by simple algebra.

 $\blacksquare (Lemma \ 2)$

Proof of Lemma 3. We have $c(1)1^2 = 2/(1 - \ln(2)) < 7$, and studying the Taylor expansion of $c(\gamma)\gamma^2$ shows that it is strictly increasing with γ , so $c(\gamma)\gamma^2 < 7$ for all $\gamma < 1$. Also, for small enough γ we have $\ln(1+\gamma) \cong \gamma - \gamma^2/2$, from which $c(\gamma) \cong 2/\gamma^2$ follows. \blacksquare (Lemma 3)

5 Final Remarks

Observe that Theorem 3 does not strictly require that the algorithm produces an hypothesis with 0 error within the first Q queries. It is enough to assume that within the first Q queries it generates a hypothesis h_i with $\epsilon_i(1+\gamma_i) < \epsilon$.

Note also that a variety of bounds on the sample size are possible by taking other definitions for γ_i . In particular, with essentially the same proof, if we take $\gamma_i = 1/i^{\beta}$ for $\beta < 1$, we obtain (approximately)

$$E[T] \le \frac{Q}{\epsilon} + \frac{1}{1-\beta} \frac{Q^{1-\beta}}{\epsilon} + 7 \frac{Q^{2\beta}}{\epsilon} \ln \frac{Q(Q+1)}{\delta}.$$

We just chose $\beta = 1/3$ to make $1 - \beta = 2\beta$, but if the values of Q and δ are known in advance, other values of β may give better bounds.

Finally, as indicated by Lemma 3, the factor 7 in front of the second term is actually a decreasing function of Q that tends to 2 as Q grows.

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