Bootstrapping and Learning PDFA in Data Streams

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Example Application: Web User Modeling



"Wish List"

- Process examples as fast as they arrive (10⁵ per sec. or more)
- Use small amount of memory (must fit into machine's main memory)
- Detect *changes* in customer behavior and *adapt* the model accordingly

Other Applications: Process Mining, Biological Models (DNA and aminoacid sequences)



Learning PDFA from Data Streams

Testing Similarity in Data Streams with the Bootstrap

Adapting to Changes in the Target

Outline

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The Data Streams Algorithmic Model

An algorithm receives an infinite stream $x_1, x_2, \ldots, x_t, \ldots$ from some domain X and must:

- Make only one pass over the data and process each item in time O(1)
- At every time t use sublinear memory (e.g. $O(\log t)$, $O(\sqrt{t})$)
- Adapt to possible "changes" in the data

It is a theoretically challenging model useful for applications:

- Originated in the algorithmics community
- Realistic for Data Mining and Machine Learning tasks in real-time
- Feasible way to deal with Big Data problems

When studying learning problems with streaming data:

- In the worst case setting it resembles Gold's model (with algorithmic constraints)
- But we consider a PAC-style scenario where:
 - x_t are all independent and generated from a distribution D_t
 - ▶ the sequence of distributions D₁, D₂,..., D_t,... either changes very slowly or presents only abrupt changes but very rarely

Hypothesis Class: PDFA

$\label{eq:probabilistic Deterministic Finite Automata = DFA + Probabilities$



Transition/Stop probabilities

\boldsymbol{q}	$p_q(a)$	$p_q(b)$	$p_q(\xi)$
1	0.3	0.7	0.0
2	0.5	0.5	0.0
3	0.8	0.0	0.2

Parameters

- n (states)
- $|\Sigma|$ (alphabet)
- L (expected length)
- μ (distinguishability, L_{∞})

 $\mu = \min_{q \neq q'} \max_{x \in \Sigma^*} |D_q(x) - D_{q'}(x)|$

State Merge/Split Algorithm

Usual approach to PDFA learning [Carrasco–Oncina '94, Ron et al. '98, Clark–Thollard '04, Palmer–Goldberg '05, Castro–Gavaldà '08, etc.]



Statistical tests

 $\begin{array}{c} S \not\approx a^{-1}S\\ S \approx b^{-1}a^{-1}S\\ S \not\approx b^{-1}S\\ a^{-1}S \not\approx b^{-1}S\\ b^{-1}S \approx a^{-1}a^{-1}S\\ b^{-1}S \approx b^{-1}b^{-1}S\end{array}$

Description of the Algorithm

System Architecture



Learner Module

initialize H with safe q_{λ} : foreach $\sigma \in \Sigma$ do add a candidate q_{σ} to H; schedule insignificance and similarity tests for q_{σ} ; **foreach** string x_t in the stream **do** foreach decomposition $x_t = wz$, with $w, z \in \Sigma^*$ do if q_w is defined then add z to \hat{S}_{w} : if q_w is a candidate and $|\hat{S}_w|$ is large enough then call SimilarityTest(q_w, δ): foreach candidate q_w do if it is time to test insignificance of q_w then if $|\hat{S}_w|$ is too small then declare q_w insignificant; else schedule another insignificance test for q_w ; if H has more than n safes or there are no candidates left then return H:

Sample Sketches for Similarity Testing

Note: Instead of keeping a sample S_w for each state q_w , the algorithm keeps a sketch \hat{S}_w of each sample

A sketch using memory $O(1/\mu)$ should be enough:

- Given samples S, S' from distributions D, D'
- \blacktriangleright Algorithm wants to test $\mathsf{L}_\infty(D,D')=0$ or $\mathsf{L}_\infty(D,D') \geqslant \mu$
- In the second case, if $|D(x) D'(x)| \ge \mu$ then either $D(x) \ge \mu$ or $D'(x) \ge \mu$
- It is enough to find all strings with D(x), $D'(x) = \Omega(\mu)$, of which there are $O(1/\mu)$

In our algorithm, each sketch uses a SpaceSaving data structure [Metwally et al. '05]:

- Uses memory $O(1/\mu)$
- \blacktriangleright Finds every string whose probability is $\Omega(\mu)$ (frequent strings)
- And approximates their probability with enough accuracy
- Easier to implement than sketches based on hash functions

Properties of the Algorithm

Streaming-specific features

- Adaptive test scheduling (decide as soon as possible)
- Similarity test based on Vapnik–Chervonenkis bound (slow similarity detection)
- Use bootstrapped confidence intervals in tests (faster convergence)

Complexity Bounds (with any reasonable test)

- Time per example O(L) (expected, amortized)
- The learner reads $O(n^2|\Sigma|^2/\varepsilon\mu^2)$ examples (in expectation)
- Memory usage is $O(n|\Sigma|L/\mu)$ (roughly $O(\sqrt{t})$)



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Testing Similarity between Probability Distributions

Goal: decide if $L_{\infty}(D, D') = 0$ or $L_{\infty}(D, D') \ge \mu$ from samples *S*, *S'*

Statistical Test Based on Empirical L_{∞} (the "default")

- Let $\mu_{\star} = L_{\infty}(D, D')$ and compute $\hat{\mu} = L_{\infty}(S, S')$
- Compute Δ_l, Δ_u such that $\hat{\mu} \Delta_l \leqslant \mu_\star \leqslant \hat{\mu} + \Delta_u$ holds w.h.p.
- If $\hat{\mu} \Delta_l > 0$ decide $D \neq D'$
- If $\hat{\mu} + \Delta_u < \mu$ decide D = D'
- Else, wait for more examples

Problem: asymmetry — deciding *dissimilarity* is easier that deciding *similarity*

- When $D \neq D'$ will decide correctly w.h.p. when |S|, $|S'| \approx 1/\mu_{\star}^2$
- When D = D' will decide correctly w.h.p. when |S|, $|S'| \approx 1/\mu^2$

In the later we are always competing against the worst case $\mathsf{L}_\infty(D,D')=\mu$

Enter the Bootstrap

- ▶ In the test I just described there is another *worst case* assumption the confidence interval $\mu_{\star} \leq \hat{\mu} + \Delta_u$ must hold *for any D and D'*
- ▶ But it may be the case that for some D, certifying that S, S' ~ D come from the same distribution is easier
- The *bootstrap* is widely used in statistics for computing *distribution dependent* confidence intervals (among many other things)

Basic Idea

- Suppose we have *r* different samples $S_{(1)}, \ldots, S_{(r)} \sim D$
- Compute distances $\hat{\mu}_i = L_{\infty}(S_{(i)}, S'_{(i)})$
- Use them to compute a histogram of the distribution of $\hat{\mu}$



Enter the Bootstrap

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Bootstrapped Confidence Intervals

- Given a sample S, obtain other samples $\tilde{S}_{(i)}$ by sampling from S uniformly with replacement
- Sort estimates increasingly $\tilde{\mu}_1 \leqslant \ldots \leqslant \tilde{\mu}_r$
- \blacktriangleright Say that $\mu_\star \leqslant \tilde{\mu}_{[(1-\delta)r]}$ with prob. $\geqslant 1-\delta$

Bootstrapped Confidence Intervals in Data Streams

Question: Do you need to store the full sample to do bootstrap resampling?

Answer: No, if you can test from sketched data

The Bootstrap Sketch

- Keep r copies of the sketch you use for testing (e.g. SpaceSaving)
- For each item x_t in the stream, randomly insert r copies of x_t into the r sketches
- Comparing each pair $\tilde{S}_{(i)}$, $\tilde{S}'_{(j)}$ can obtain r^2 approximations $\tilde{\mu}_{i,j}$
- Choosing r involves a trade-off between accuracy and memory



In theory can prove bound (asymptotically) comparable to Vapnik–Chervonenkis In practice assuming $\mu_{\star} \leqslant \tilde{\mu}_{[(1-\delta)r^2]}$ gives accurate and statisically efficient similarity test

Experimental Results for Learner

- \blacktriangleright Prototype written in C++ and Boost, run in this laptop
- Evaluated with Reber Grammar (typical Grammatical Inference benchmark)

• $|\Sigma| = 5, n = 6, \mu = 0.2, L \approx 8$

• Compared VC and Bootstrap (r = 10) based tests

	Examples	Memory (MiB)	Time/item (ms)
Hoeffding	57617	6.1	0.05
Bootstrap	23844	53.7	1.2

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What if n and μ are unknown (or change)?

Want to design strategy for fast and accurate parameter estimation

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Parameter Search Algorithm
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 $n \leftarrow 2, \ \mu \leftarrow 1/8;$ while true do

> $H \leftarrow \text{Learner}(n, \mu);$ if |H| < n then $\mu \leftarrow \mu/8;$ else $n \leftarrow 2n;$ if $n > (1/\mu)^{1/3}$ then $\mu \leftarrow \mu/8;$

Complexity Bounds

- Needs only $O(\log(n_\star/\mu_\star^{1/3}))$ calls to Learner
- In expectation will read $O(n_\star^6|\Sigma|^2/\epsilon\mu_\star^2)$ elements
- Memory usage grows like $O(t^{2/3})$

Note: can tweak parameters to trade-off convergence speed and memory usage

Adapting the Hypothesis to Changes

Adapter block — Once the structure is known...

- Estimating probabilities is easy
- Estimations can be adapted to changes (e.g. moving average)



Transition/Stop probabilities

 $S = \{abb, baab, bbaabb\}$

q	$p_q(a)$	$p_q(b)$	$p_q(\xi)$
1	2/6	4/6	0/6
2	2/6	4/6	0/6
3	1/4	0/4	3/4

But, sometimes the current structure is not good anymore

Detecting Structural Changes

Idea: "change" is *difficult to define* in general, focus on changes explained in terms of *structure*



- Given a PDFA, compute the expected number of times each state is visited when generating a string
- Given a sample, compute the average number of times strings hit any state
- If there is a significant difference, conclude the structure has changed

$$S = \{abb, baab, bbaabb\}$$

$$\begin{array}{c|cc} h_1 & h_2 & h_3 \\ \hline 6/3 & 6/3 & 4/3 \end{array}$$

- Restart structure learning when a change is detected
- Adapting probabilities may be enough, but re-learning does no damage

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Conclusion

Summary of Contributions

- Adaptation of state-merging paradigm to streaming data
- Fast convergence achieved by:
 - adaptive test scheduling
 - better similarity testing
 - efficient parameter search
- Use of sketching algorithms for implementing the bootstrap and reducing memory usage Future Work
 - Deploy real system and exploit parallelization oportunities
 - Develop further similarity tests based on the bootstrap
 - Adapt other GI algorithms to the data streams framework

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