Approximate Statistical Discrepancy

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Problem Setup

Let X be a set of m points in \mathbb{R}^d .



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Let (X, A) be a range space, defining a family subsets A of X. Each data point can belong to an anomalous set

Find a range $A \in \mathcal{A}$ where the anomalous data is significantly denser than the baseline data using some function Φ



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Statistical Functions Φ

2 scalar values for each $x \in X$: b(x) and r(x). For any $A \cap X$, define $B = \sum_{x \in X} b(x)$ and $R = \sum_{x \in X} r(x)$ and

$$b(A) = \frac{1}{B} \sum_{x \in X \cap A} b(x)$$
 and $r(A) = \frac{1}{R} \sum_{x \in X \cap A} r(x)$

Define statistic $\Phi(A)$ as the log likelihood ratio

$$\Phi(A) = \log\left(rac{\Pr(\mathcal{H}_0|A,X)}{\Pr(\mathcal{H}_1|A,X)}
ight)$$

- ► H₀ : no anomaly, rate of measured points same inside as outside.
- \mathcal{H}_1 : anomaly, A has a different rate of measured points inside than outside.

► Formulate a model of the data and choose a corresponding scan statistic φ to score the likelihood of an anomaly in a region.

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► Assess whether the score φ(A*) indicates A* is significant. Requires typically 1000 "scans" for permutation tests Many existing papers on these algorithms:

- Classic discrepancy maximization [BDT16, DE93]
- Subroutine in algorithms ranging from computer graphics [DEM96] to association rules in data mining [FMMT96]
- Minimum disagreement problem in machine learning [LM96].

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 Scan Statistics [Kul97, Kul06, HKG07, NM04, APV06, AMP⁺06, KHPD06, TT05](many many more) Spatial Scan Statistics are heavily used to find spatial anomalies.

FiveThirtyEight

Politics Sports Science & Health Economics Culture

NOV. 16, 2016 AT 8:00 AM

How New York Hunts For Early Signs Of Disease Outbreaks

By Ian Evans

Filed under Public Health



On July 29, 2015, the New York City Department of Health and Mental Hygiene sent out an alert — 31 people in the South Bronx had contracted Legionnaires' disease, a lung infection from waterborne bacteria that kills about 1 out of every 10 people who get it. By the time officials found the source (a cooling tower) and contained the spread, 128 people had contracted Legionnaires' and 12 people had died. It was the largest outbreak of Legionnaires' disease in the city's history — an outbreak that was first detected by a computer program.

Find an approximate range $\hat{A} \in \mathcal{A}$ such that $\Phi(A^*) - \Phi(\hat{A}) < \varepsilon$.

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Existing approximation papers.

- [AMP⁺06] which introduced generic sampling bounds and a bound on approximating scan statistics with linear functions.
- [MSZ⁺16] which showed that a two-stage random sampling can provide some error guarantees.
- [Wal10] which showed approximation guarantees under the Bernoulli model.

Approximate Problem

Find an approximate range $\hat{A} \in \mathcal{A}$ such that $\Phi(A^*) - \Phi(\hat{A}) < \varepsilon$.

For m = |X|

	Known Exact	Known Approx [MSZ ⁺ 16]	New Runtime Bounds
General Range Space	$O(m^{ u+1})$		$O\left(m + \frac{1}{\varepsilon^{\nu+2}}\log^{\nu}\frac{1}{\varepsilon}\right)$
Halfspaces	<i>O</i> (<i>m</i> ^{<i>d</i>}) [DEM96]	-	$O\left(m + \frac{1}{\varepsilon^{d+1/3}}\log^{2/3}\frac{1}{\varepsilon}\right)$
Disks	O(m ³) [DEM96]	$O(m + \frac{1}{\varepsilon^4} \log^3 \frac{1}{\varepsilon})$	$O\left(m+\frac{1}{c^{3+1/3}}\log^{2/3}\frac{1}{\varepsilon}\right)$
Rectangles (Disc)	O(m ²) [BCNPL14]	$O(m + \frac{1}{\varepsilon^4} \log \frac{1}{\varepsilon})$ [APV06]	$O(m + \frac{1}{\varepsilon^2} \log \log \frac{1}{\varepsilon})$
Rectangles (SDF)	$O(m^4)$	$O(m + \frac{1}{\varepsilon^4} \log^4 \frac{1}{\varepsilon})$	$O\left(m + \frac{1}{\epsilon^{2.5}}\right)$
Rectangles (Gen)	$O(m^4)$	$O(m + \frac{1}{\varepsilon^4} \log^4 \frac{1}{\varepsilon}))$	$O\left(m+\frac{1}{\varepsilon^4}\right)$

Algorithm times for (ε -approximately) maximizing different range spaces. Here dimension d, VC-dimension ν , and probability of failure are all constants. For Rectangles (Disc) we show it takes $\Omega(m + 1/\varepsilon^2)$ time, assuming hardness of APSP.

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$$\phi(r,b)=C_1r+C_2b$$

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- New result: only need $O(1/\sqrt{\varepsilon})$ linear functions
- ► In practice only need 3 to 4 linear functions.

Approximating ϕ (cont)

Approximate ϕ with $O(1/\sqrt{\varepsilon})$ linear functions.

• If ϕ is a convex function then max lies on convex hull of:

$$V_{\mathcal{A},m,b} = \{(r(A), b(A)) \mid A \in \mathcal{A}\}$$

► Approximate convex hull of V_{A,r,b} by picking linear functions in an iterative way.



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► Size bound by Dudley's approximation (ε-kernel)

Approximate Rectangle Scanning

• Grid G over X so that each row and column has $\approx \varepsilon |X|$ points.



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- Grid G over X so that each row and column has $\approx \varepsilon |X|$ points.
- ► Each rectangle R ∈ (X, R₂) is approximated by a subgrid R_G ∈ G.



Approximate Rectangle Scanning

- Grid G over X so that each row and column has $\approx \varepsilon |X|$ points.
- ► Each rectangle $R \in (X, \mathcal{R}_2)$ is approximated by a subgrid $R_G \in G$.
- Can enumerate all subgrids in O(¹/_{ε⁴}) time and compute Φ on each.(O(m + ¹/_{ε⁴}))



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Approximate Rectangle Scanning for Linear Function

- Grid G over X so that each row and column has $\approx \varepsilon |X|$ points.
- Fix $1/\varepsilon$ upper end-points and sweep $1/\varepsilon$ lower end-points.



Approximate Rectangle Scanning for Linear Function

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- ▶ Fix $1/\varepsilon$ upper end-points and sweep $1/\varepsilon$ lower end-points.
- ► Run Kadane's algorithm for each upper and lower endpoint to compute maximum horizontal subgrid in time O(¹/_ε).
- Max subgrid in $O(\frac{1}{\varepsilon^3})$ time (max likelihood function in $O(1/\varepsilon^{3.5})$ time.)



Can be made faster:

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Most subgrids are very similar.

Can be made faster:

- Previously scanning was exact and approximation was from restricting to a grid.
- Most subgrids are very similar.
- Make scanning approximate so that only small updates occur to similar subgrids.

• Consider computing the max subgrid spanning a slab *M*.



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- Consider computing the max subgrid spanning a slab *M*.
- ► Divide upper subgrid into subgrids T_T and T_B and lower subgrids into B_T and B_B.
- Decompose into 4 seperate problems.



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- Divide upper subgrid into subgrids T_T and T_B and lower subgrids into B_T and B_B.
- Decompose into 4 seperate problems.
- Idea inspired by [BCNPL14, Tak02, DEM96]



Have to merge T_B and/or merge B_T into M.

- 1. Merge neither into M.
- 2. Merge T_B into M.
- 3. Merge B_T into M.
- 4. Merge T_B and B_T into M.



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Merging can be done in time proportional to non-zeros columns (see paper for details).

• If T_b or B_b has k rows then can construct sparse grid with $O(k \log \frac{1}{\varepsilon})$ non zero columns that misplaces $\varepsilon / \log \frac{1}{\varepsilon} |X|$ points with respect to vertical intervals.



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- ► Have to construct a tree of sparse *subgrids* first.
- Error adds over log ¹/_ε levels in the recurrence leading to ε|X| misplaced points.
- Total run time is $O(m + \frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon})$
 - $O(m + \frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon})$ time to build grid.
 - Takes $O(\frac{1}{\epsilon^2})$ time to construct tree of sparse subgrids.
 - ► Takes O(¹/_{ε²} log ¹/_ε) to compute max slab spanning subgrid approximately.

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- $\log \frac{1}{\varepsilon}$ can be made $\log \log \frac{1}{\varepsilon}$ with some work.
- New lower bound conditional on APSP of $\Omega(|X| + \frac{1}{\epsilon^2})$.

Faster in Practice

Significant improvement in convergence

- ► SatScan is exact algorithm run on sample.
- gridScan is simple scanning over a grid.
- ▶ gridScan_linear is Kadane based algorithm.
- Do not have an implementation of fastest algorithm.



Other range spaces

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Why are exact algorithms slow on a Sample?



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Problem: Far too many combinatorial regions.



Idea: Use smaller sample of size $O_d(\frac{1}{\varepsilon})$ to induce regions.



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Compute ϕ using dense sample.



Repeat procedure log $\frac{1}{\delta}$ times and take median to amplify probability of success.



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Procedure works on many different range spaces.



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Fast Scanning of Halfspaces

Dobkin and Eppstein [DE93] maximize (X, \mathcal{H}) of *n* points in \mathbb{R}^d in $O(n^d)$ time.





Dual

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Fast Scanning of Halfspaces

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But need to count points in S.

Can annotate dual arrangement in O(n) time in \mathbb{R}^2 for each $x \in S$.

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S is usually of size $O(\frac{1}{\varepsilon^2})$ for constant d

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• Halfspaces
$$\mathcal{A} = \mathcal{H}$$
 in \mathbb{R}^2 then $|S| = s = O((1/\varepsilon)^{4/3})$

• Balls
$$\mathcal{A} = \mathcal{B}$$
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Provides further speedup.

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Method exists for computing Halfspaces samples [MP18]:

•
$$|S| = O((1/\varepsilon)^{2d/(d+1)} \log^{d/(d+1)}(1/\varepsilon))$$

• Computable in $O(n + \frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon})$ time.

Provides further speedup.

For m = |X|

	Known Exact	Known Approx [MSZ ⁺ 16]	New Runtime Bounds
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Halfspaces	<i>O</i> (<i>m^d</i>) [DEM96]	-	$O\left(m + \frac{1}{\varepsilon^{d+1/3}}\log^{2/3}\frac{1}{\varepsilon}\right)$
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Any Questions?

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Evaluating Discrepancy

$$b(A) = \frac{1}{B} \sum_{x \in X \cap A} b(x) \text{ and } r(A) = \frac{1}{R} \sum_{x \in X \cap A} r(x)$$

Let $b(x) = -1$ and $m(x) = \{0, +2\}$
 $\phi(b, r) = |B(m(A) + b(A))|$

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Evaluating Discrepancy

$$b(A) = \frac{1}{B} \sum_{x \in X \cap A} b(x) \text{ and } r(A) = \frac{1}{R} \sum_{x \in X \cap A} r(x)$$

Let $b(x) = -1$ and $m(x) = \{0, +2\}$

$$\phi(b,r) = |B(m(A) + b(A))|$$

Equivalent to *discrepancy evaluation* for a range space (X, A) and a coloring $\chi : X \to \{-1, +1\}$: Find

$$\mathsf{disc}_\chi(X,\mathcal{A}) = rg\max_{A\in\mathcal{A}} \left|\sum_{x\in X\cap A} \chi(x)
ight|.$$

If $\chi(x) = +1$ then r(x) = 2 otherwise if $\chi(x) = -1$ then r(x) = 0.

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