A Polynomial Number of Random Points does not Determine the Volume of a Convex Body

Ronen Eldan, Tel Aviv University

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- We are looking for polynomial time algorithms which approximate the volume up to a (multiplicative) constant C with probability ≈ 1 .

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 - Since the early 90's many improvements were found, and the running time was improved from O(n²⁴) to O(n⁴), given by L. Lovász, S. Vempala in 2004. Additional works by B.Bollobás, N. Goyal, A.Kalai, L.Rademacher, M.Simonovits and other people whose names are also very hard to pronounce.

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- ► The volume cannot be approximated even if every point is given in an infinitely good accuracy.
- ▶ A related result of N.Goyal and L.Rademacher shows that in order to learn a convex body one needs $2^{\Omega(\sqrt{n})}$ random points.

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- ▶ The reason: Consider the event that two points are picked from the same cell. If we condition on this event, *C* does not change anything.
- ▶ But since the number of cells is exponential, this event is negligible.

A very simple toy model - continued

A constant C induces a probability measure on the set of sequences of points, Ω. The **total variation** distance between two such measures is small.

$$d_{TV}(X_1, X_2) = \sup_{A \subset \Omega} |Prob(X_1 \in A) - Prob(X_2 \in A)|$$

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- Maybe we can try some other random deletion process?

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- ▶ Suppose K_1 and K_2 convex-body-valued random variables with the following properties:
 - ▶ With high probability, $Vol(K_1) \approx C_1$ and $Vol(K_2) \approx C_2$, and $\frac{C_1}{C_2}$ is very large.

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 - ▶ The random convex body K_i induces a probability measure on Ω in the following simple way: first generate a body K and then generate a sequence of uniformly distributed random points from K.
 - ► These corresponding measures have a small total variation distance.



▶ A small total variation distance between the random variables X_1, X_2 representing sequences of random points implies that for every algorithm $F: \Omega \to \mathbb{R}$, we would have a small total variation distance between $F(X_1)$ and $F(X_2)$.

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- ▶ To summarize: we are looking for two different random constructions of convex bodies whose volumes are very different, yet whose output of random points is very similar, so that no algorithm can distiguish between them.

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At this point, let us imagine that $T(\theta) = T_0(\theta)$. Let $\zeta = (\theta_1, ..., \theta_{m'})$ be a rotation invariant Poisson process on S^{n-1} with intensity m. Define

$$K = K_{T,m} = \bigcap_{1 \le i \le m'} T(\theta_i)$$

Let $x_1, x_2 \in D_n$. Let us try to compare $\mathbb{P}(x_1 \in K \& x_2 \in K)$ with $\mathbb{P}(x_1 \in K)\mathbb{P}(x_2 \in K)$.

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$$\frac{\mathbb{P}(x_1 \in K \& x_2 \in K)}{\mathbb{P}(x_1 \in K)\mathbb{P}(x_2 \in K)} = \frac{e^{-(m(S_1) + m(S_2) + m(S_b))}}{e^{-(m(S_1) + m(S_b) + m(S_2) + m(S_b))}} = e^{m(S_b)}$$

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- ► Therefore, even if the chances of a single point to be removed are rather high, the chances for both to be removed in the same deletion are very small.
- ► This means that we can cut a large portion of mass, still having

$$\mathbb{P}(x_1 \in K \& x_2 \in K)(1 + O(e^{-n^{\epsilon}})) = \mathbb{P}(x_1 \in K)\mathbb{P}(x_2 \in K)$$

for some $\epsilon > 0$.



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▶ The above "almost independence" implies,

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► Hence, the volumes are very concentrated around their expectation.

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we get

$$P(A) \approx \frac{1}{\mathbb{E}(Vol(K^N))} \int_A \prod_i \mathbb{P}(\forall i, x_i \in K)$$

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- Since the construction was centrally symmetric, all we have to do in order for the total variation distance between two induced measures to be small is to make sure that the distribution of |x₁| is approximately the same in the two families.
- ► To do this, we use the liberty of choosing the shape of the cut, and the intensity of the poisson processes.

Possible further research

In fact, our proof gives a slightly stronger result:

Theorem

There exists $\varepsilon>0$ and a number $n_0\in\mathbb{N}$ such that for all $n>n_0$, there does not exist an algorithm which takes e^{n^ε} points generated randomly according to the uniform measure in a convex body $K\subset\mathbb{R}^n$, which determines Vol(K) up to e^{n^ε} with probability more than e^{-n^ε} to be correct.

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▶ The volume radius of a convex body $K \subset \mathbb{R}^n$ is defined as $Volrad(K) = Vol(K)^{\frac{1}{n}}$. Clearly, it is much easier to estimate the volume radius, but is it possible (say, up to 1.001)?

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- The above question is equivalent to the question whether or not the isotropic constant of a body can be estimated using a polynomial number of points.

What about a general convex body?

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- ► This result gives us two families of convex bodies that cannot be distinguished if we only have a polynomial number of random points. What can be said about the distribution of n¹⁰⁰ random points in a general convex body?
- If we first apply a random rotation to the body, then the marginal distribution of a single point is spherically symmetric and therefore all of the information is contained in the Gramm matrix.
- ► The recent thin shell results imply that the main diagonal entries are all concentrated around n.
- What about the off diagonal entries?
 - Is their distribution gaussian?
 - ► For a small enough number of points, the off diagonal entries may be distributed in the same way for all convex bodies.

Thank you!