# Distribution Testing and Probabilistic Programming: A Match made in Heaven 

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## Probabilistic Programs and Distributions

```
program P
    X[1] = Bernoulli( (p1);
    X[2] = Bernoulli (p2);
    X[n] = Bernoulli( }\mp@subsup{p}{n}{})\mathrm{ ;
    observe(X[2] + X[3] + X[7] > 2);
    return Y
```

- Semantics of discrete probabilistic programs: Distributions over $\{0,1\}^{n}$
- Distance Measures
- Closeness
- $d_{\infty}(P, Q)=\max _{x \in\{0,1\}^{n}}|P(x)-Q(x)|$
- Farness
- $d_{T V}(P, Q)=\frac{1}{2} \sum_{x \in\{0,1\}^{n}}|P(x)-Q(x)|$
- Hardness of probabilistic program distance computation
- Grey-box Testing
- The Boon of Testers


## Hardness of probabilistic program distance computation

## Product Distributions

- Represented by list of probabilities: $\left\{p_{1}, p_{2}, \ldots p_{n}\right\}$
- $P(x)=\prod_{x_{i}=1} p_{i} \prod_{x_{i}=0}\left(1-p_{i}\right)$


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Question How hard it is to compute $d_{T V}(P, Q)$ when $P$ and $Q$ are product distributions

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Question How hard it is to compute $d_{T V}(P, Q)$ when $P$ and $Q$ are product distributions

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program P
    X[1] = Bernoulli ( (p1);
    X[2] = Bernoulli (p2);
    X[n] = Bernoulli ( }\mp@subsup{p}{n}{})
    return X;
```

```
program Q
```

program Q
Y[1] = Bernoulli(q}\mp@subsup{q}{1}{})
Y[1] = Bernoulli(q}\mp@subsup{q}{1}{})
Y[2] = Bernoulli(q2);
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Y[n] = Bernoulli(q})\mathrm{ ;
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Given two product distributions $P$ and $Q$, computation of $d_{T V}(P, Q)$ is \#P-hard.

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- We reduce from \#Knapsack, i.e., we construct $P$ and $Q$ such that $\#\{x: P(x)=Q(x)\}$ will solve \#Knapsack
- Now given $P$ and $Q$, we construct $P^{\prime}$ and $Q^{\prime}$ where $p_{i}^{\prime}=p_{i}$ and $q_{i}^{\prime}=q_{i}$ for $i \leq n$ and $p_{n+1}^{\prime}=\frac{1}{2}+\gamma$ and $q_{n+1}^{\prime}=\frac{1}{2}-\gamma$ such that whenever $P(x)>Q(x)$ we have $P(x)\left(\frac{1}{2}-\gamma\right)>Q(x)\left(\frac{1}{2}+\gamma\right)$

$$
\begin{aligned}
d_{T V}\left(P^{\prime}, Q^{\prime}\right)=\sum_{x \in\{0,1\}^{n}} & \left(\left|P(x)\left(\frac{1}{2}+\gamma\right)-Q(x)\left(\frac{1}{2}-\gamma\right)\right|\right. \\
& \left.+\left|P(x)\left(\frac{1}{2}-\gamma\right)-Q(x)\left(\frac{1}{2}+\gamma\right)\right|\right) \\
& =d_{T V}(P, Q)+2 \gamma \sum_{x: P(x)=Q(x)} P(x)
\end{aligned}
$$

## Grey-box Testing

## Basic Access

- We can run the program: sample!


## What does Complexity Theory Tell Us



Figure: $\mathcal{U}$ : Reference Distribution


Figure: $\mathcal{A}$ : $1 / 2$-far from uniform

## What does Complexity Theory Tell Us



Figure: $\mathcal{U}$ : Reference Distribution


Figure: $\mathcal{A}$ : $1 / 2$-far from uniform

- If $<\sqrt{S} / 100$ samples are drawn then with high probability you see only distinct samples from either distribution.


## Theorem (Batu-Fortnow-Rubinfeld-Smith-White (JACM 2013))

Testing whether a distribution is $\epsilon$-close to uniform has query complexity $\Theta\left(\sqrt{|S|} / \epsilon^{2}\right)$. [Paninski (Trans. Inf. Theory 2008)]

- If the output of a program is represented by 3 doubles, then $S>2^{100}$


## Beyond Basic Access

## Definition (Conditional Sampling)

Given a distribution $\mathcal{A}$ on $S$ one can

- Specify a set $T \subseteq S$,
- Draw samples according to the distribution $\left.\mathcal{A}\right|_{T}$, that is, $\mathcal{A}$ under the condition that the samples belong to $T$.


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Conditional sampling is at least as powerful as drawing normal samples. But how more powerful is it?

## Testing Uniformity Using Conditional Sampling




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An algorithm for testing uniformity using conditional sampling:
(1) Draw two elements $x$ and $y$ uniformly at random from the domain. Let $T=\{x, y\}$.
(2) In the case of the "far" distribution, with probability $\frac{1}{2}$, one of the two elements will have probability 0 , and the other probability non-zero.
(3) Note $\sqrt{|T|}=\sqrt{2}$ is a constant.
(9) Now a constant number of conditional samples drawn from $\left.\mathcal{A}\right|_{T}$ is enough to identify that it is not uniform.

What about other distributions?



## What about other distributions?



Previous algorithm fails in this case:
(1) Draw two elements $\sigma_{1}$ and $\sigma_{2}$ uniformly at random from the domain. Let $T=\left\{\sigma_{1}, \sigma_{2}\right\}$.
(2) In the case of the "far" distribution, with probability almost 1 , both the two elements will have probability same, namely $\epsilon$.
(3) Probability that we will be able to distinguish the far distribution from the uniform distribution is very low.

Testing Uniformity Using Conditional Sampling




## Testing Uniformity Using Conditional Sampling




(1) Draw $\sigma_{1}$ uniformly at random from the domain and draw $\sigma_{2}$ according to the distribution $\mathcal{A}$. Let $T=\left\{\sigma_{1}, \sigma_{2}\right\}$.
(2) In the case of the "far" distribution, with constant probability, $\sigma_{1}$ will have "low" probability and $\sigma_{2}$ will have "high" probibility.
(3) We will be able to distinguish the far distribution from the uniform distribution using constant number of samples from $\left.\mathcal{A}\right|_{T}$.
(9) The constant depend on the farness parameter.

## Barbarik

Input: A Distribution under test $\mathcal{A}$, a reference uniform distribution $\mathcal{U}$, a tolerance parameter $\varepsilon>0$, an intolerance parmaeter $\eta>\varepsilon$, a guarantee parameter $\delta$
Output: ACCEPT or REJECT with the following guarantees:

- if the generator $\mathcal{A}$ is $\varepsilon$-close (in $d_{\infty}$ ), i.e., $d_{\infty}(\mathcal{A}, \mathcal{U}) \leq \varepsilon$ then Barbarik ACCEPTS with probability at least $(1-\delta)$.
- If the generator $\mathcal{A}$ is $\eta$-far in $\left(d_{T V}\right)$, i.e., $d_{T V}(\mathcal{A}, \mathcal{U})>\eta$, then Barbarik REJECTS with probability at least $1-\delta$.


## Sample complexity

## Theorem

Given $\varepsilon, \eta$ and $\delta$, Barbarik need at most $K=\widetilde{O}\left(\frac{1}{(\eta-\varepsilon)^{4}}\right)$ samples for any input formula $\varphi$, where the tilde hides a poly logarithmic factor of $1 / \delta$ and $1 /(\eta-\varepsilon)$.

- $\varepsilon=0.6, \eta=0.9, \delta=0.1$
- Maximum number of required samples $K=1.72 \times 10^{6}$
- Independent of the number of variables
- To Accept, we need $K$ samples but rejection can be achieved with lesser number of samples.


## Extensions to general case

Setting Given $P$ and $Q$, test whether $P$ and $Q$ are $\varepsilon$-close (in $d_{\infty}$ ) or $\eta$-far (in $d_{T V}$ )

- For all $\sigma_{1}, \sigma_{2} \in 0,1^{n}, \frac{P\left(\sigma_{1}\right)}{P\left(\sigma_{2}\right)}$ is known: A prior distribution conditioned by constraints: dependence on $\max _{\sigma_{1}, \sigma_{2} \in 0,1^{n}} \frac{P\left(\sigma_{1}\right)}{P\left(\sigma_{2}\right)}$ (MPC20; PM21)
- Arbitrary $P$ and $Q$
(DCM22 - under submission)


## The Boon of Testers

## Where are the Samplers?

- Implicit representation of a set $S$ : Set of all solutions of $\varphi$.
- Given a CNF formula $\varphi$, a Sampler $\mathcal{A}$, outputs a random solution of $\varphi$.


## Definition

A CNF-Sampler, $\mathcal{A}$, is a randomized algorithm that, given a $\varphi$, outputs a random element of the set $S$, such that, for any $\sigma \in S$

$$
\operatorname{Pr}[\mathcal{A}(\varphi)=\sigma]=\frac{1}{|S|},
$$

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- Uniform sampling has wide range of applications in automated bug discovery, pattern mining, and so on.
- Several samplers available off the shelf: tradeoff between guarantees and runtime


## Constrained Sampling

- Input formula: $F$ over variables $X$
- Challenge: Conditional Sampling over $T=\left\{\sigma_{1}, \sigma_{2}\right\}$.
- Construct $G=F \wedge\left(X=\sigma_{1} \vee X=\sigma_{2}\right)$


## Constrained Sampling

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- Challenge: Conditional Sampling over $T=\left\{\sigma_{1}, \sigma_{2}\right\}$.
- Construct $G=F \wedge\left(X=\sigma_{1} \vee X=\sigma_{2}\right)$
- Most of the samplers enumerate all the points when the number of points in the Domain are small
- Need way to construct formulas whose solution space is large but every solution can be mapped to either $\sigma_{1}$ or $\sigma_{2}$.


## Kernel

Input: A Boolean formula $\varphi$, two assignments $\sigma_{1}$ and $\sigma_{2}$, and desired number of solutions $\tau$
Output: Formula $\hat{\varphi}$
(1) $\tau=|\operatorname{Sol}(\hat{\varphi})|$
(2) $\operatorname{Supp}(\varphi) \subseteq \operatorname{Supp}(\hat{\varphi})$
(3) $z \in \operatorname{Sol}(\hat{\varphi}) \Longrightarrow z_{\downarrow} S \in\left\{\sigma_{1}, \sigma_{2}\right\}$
(1) $\left|\left\{z \in \operatorname{Sol}(\hat{\varphi}) \mid z_{\downarrow \operatorname{Supp}(\varphi)}=\sigma_{1}\right\}\right|=\left|\left\{z \in \operatorname{Sol}(\hat{\varphi}) \mid z_{\downarrow \operatorname{Supp}(\varphi)} \cap \sigma_{2}\right\}\right|$
(5) $\varphi$ and $\hat{\varphi}$ has similar structure

## Non-adversarial Sampler

Let $(\hat{\varphi})$ obtained from $\operatorname{kernel}\left(\varphi, \sigma_{1}, \sigma_{2}, N\right)$ such that there are only two set of assignments to variables in $\varphi$ that can be extended to a satisfying assignment for $\hat{\varphi}$

## Definition

The non-adversarial sampler assumption states that the distribution of the projection of samples obtained from $\mathcal{A}(\hat{\varphi})$ to variables of $\varphi$ is same as the conditional distribution of $\mathcal{A}(\varphi)$ restricted to either $\sigma_{1}$ or $\sigma_{2}$

- If $\mathcal{A}$ is a uniform sampler for all the input formulas, it satisfies non-adversarial sampler assumption
- If $\mathcal{A}$ is not a uniform sampler for all the input formulas, it may not necessarily satisfy non-adversarial sampler assumption


## Barbarik

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Output: ACCEPT or REJECT with the following guarantees:

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- If the generator $\mathcal{A}$ is $\eta$-far in $\left(d_{T V}\right)$, i.e., $d_{T V}(\mathcal{A}, \mathcal{U})>\eta$ and if non-adversarial sampler assumption holds then Barbarik REJECTS with probability at least $1-\delta$.


## Experimental Setup

- Three state of the art (almost-)uniform samplers
- UniGen2: Theoretical Guarantees of almost-uniformity
- SearchTreeSampler: Very weak guarantees
- QuickSampler: No Guarantees
- The study (in 2018) that proposed Quicksampler perform unsound statistical tests and claimed that all the three samplers are indistinguishable


## Results-I

| Instances | \#Solutions | UniGen2 |  | SearchTreeSampler |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Output | \#Samples | Output | \#Samples |
| 71 | $1.14 \times 2^{59}$ | A | 1729750 | R | 250 |
| blasted_case49 | $1.00 \times 2^{61}$ | A | 1729750 | R | 250 |
| blasted_case50 | $1.00 \times 2^{62}$ | A | 1729750 | R | 250 |
| scenarios_aig_insertion1 | $1.06 \times 2^{65}$ | A | 1729750 | R | 250 |
| scenarios_aig_insertion2 | $1.06 \times 2^{65}$ | A | 1729750 | R | 250 |
| 36 | $1.00 \times 2^{72}$ | A | 1729750 | R | 250 |
| 30 | $1.73 \times 2^{72}$ | A | 1729750 | R | 250 |
| 110 | $1.09 \times 2^{76}$ | A | 1729750 | R | 250 |
| scenarios_tree_insert_insert | $1.32 \times 2^{76}$ | A | 1729750 | R | 250 |
| 107 | $1.52 \times 2^{76}$ | A | 1729750 | R | 250 |
| blasted_case211 | $1.00 \times 2^{80}$ | A | 1729750 | R | 250 |
| blasted_case210 | $1.00 \times 2^{80}$ | A | 1729750 | R | 250 |
| blasted_case212 | $1.00 \times 2^{88}$ | A | 1729750 | R | 250 |
| blasted_case209 | $1.00 \times 2^{88}$ | A | 1729750 | R | 250 |
| 54 | $1.15 \times 2^{90}$ | A | 1729750 | R | 250 |

## Results-II

| Instances | \#Solutions | UniGen2 |  | QuickSampler |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
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## Beyond Simply Testing

How can we use the availability of Barbarik to design a good sampler?

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Wishlist

- Sampler should pass the Barbarik test.
- Sampler should be at least as fast as STS and QuickSampler.
- Sampler should perform good on real world applications.


## CMSGen

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- Processing techniques: bounded variable elimination, local search, or symmetry breaking, and many more.
- Can change solution space of instances.
- Restart at static intervals.
- Helps to generate samples which are very hard to find.
- Test-Driven Development of CMSGen.
- Parameters of CMSGen are decided with the help of Barbarik
- Iterative process.
- Based on feedback from Barbarik, change the parameters.
- Uniform-like-sampler.
- Lack of theoretical analysis.


## Testing of Samplers

- Samplers without guarantees (Uniform-like Samplers):
- STS (Ermon, Gomes, Sabharwal, Selman, 2012)
- QuickSampler (Dutra, Laeufer, Bachrach, Sen, 2018)
- Sampler with guarantees:
- UniGen3 (Chakraborty, Meel, and Vardi 2013, 2014,2015)

|  | QuickSampler | STS | UniGen3 |
| :--- | :---: | :---: | :---: |
| ACCEPTs | 0 | 14 | 50 |
| REJECTs | 50 | 36 | 0 |

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## QuickSampler STS UniGen3 CMSGen

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- Sampler should pass the Barbarik test. $\checkmark$
- Sampler should be at least as fast as STS and QuickSampler.
- Sampler should perform good on real world applications.


## CMSGen vs. Other State-of-the-Art Samplers

- 70 Benchmarks arising from:
- probabilistic reasoning, (Chakraborty, Fremont, Meel et al.,2015)
- bounded model checking. (Clarke, Biere, Raimi, Zhu, 2001)
- bug synthesis. (Roy, Pandey, Dolan-Gavitt,Hu, 2018)
- Runtime evaluation to generate 1000 samples.
- Timeout: 7200 seconds.


## CMSGen vs. Other State-of-the-Art Samplers (II)



## QuickSampler STS CMSGen <br> 33 <br> 37 <br> 52

- Sampler should pass the Barbarik test. $\checkmark$
- Sampler should be at least as fast as STS and QuickSampler. $\checkmark$
- Sampler should perform good on real world applications.


## Combinatorial Testing

- A powerful paradigm for testing configurable system.
- Challenge: To generate test suites that maximizes $t$-wise coverage.

$$
\mathrm{t} \text {-wise coverage: }=\frac{\# \text { of t-sized combinations in test suite }}{\text { all possible valid t-sized combinations }}
$$

- To generate the test suites use constraint samplers.
- Uniform sampling to have high $t$-wise coverage (Plazar, Acher, Perrouin et al., 2019).


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- To generate the test suites use constraint samplers.
- Uniform sampling to have high $t$-wise coverage (Plazar, Acher, Perrouin et al., 2019).
- Experimental Evaluations:
- Generate 1000 samples (test cases).
- 110 Benchmarks, Timeout: 3600 seconds
- 2-wise coverage $t=2$.


## Combinatorial Testing: The Power of CMSGen

Higher is better


## A Long and Winding Road

- Hardness of probabilistic program distance computation
- Distribution Testing
- The Boon of Testers

Open Source Tools
Barbarik https://github.com/meelgroup/barbarik
CMSGen https://github.com/meelgroup/cmsgen

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Where do we go from here?

- Exploring the tradeoffs: the cost of conditioning and query complexity
- Modular proofs
- Benchmark suite generation

These slides are available at https://tinyurl.com/meel-talk

