

# On sampling and approximate counting

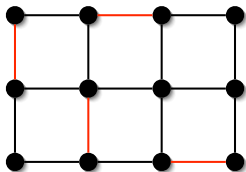
Mark Jerrum

Queen Mary, University of London

Collège de France, 9th January 2018

# Example 1: Matchings (monomer-dimer)

Instance: a graph  $G = (V, E)$ .



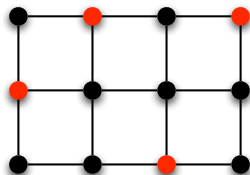
A *matching* is a collection  $M \subseteq E$  of vertex-disjoint edges.

$$\pi(M) = \lambda^{|M|} / Z_{\text{match}}, \quad \text{where } Z_{\text{match}} = Z_{\text{match}}(G, \lambda) = \sum_M \lambda^{|M|}.$$

Task: Sample from  $\pi$ , efficiently.

## Example 2: Independent sets (hard-core gas)

Instance: a graph  $G = (V, E)$ .



An *independent set* is a subset  $S \subseteq V$  of non-adjacent vertices.

$$\pi(S) = \lambda^{|S|} / Z_{IS}, \quad \text{where } Z_{IS} = Z_{IS}(G, \lambda) = \sum_S \lambda^{|S|}.$$

Task: As before.

# Estimating the partition function

A related computational task is estimating the *partition functions*  $Z_{\text{match}}(G, \lambda)$  and  $Z_{\text{IS}}(G, \lambda)$  for a specified graph  $G$  and *activity*  $\lambda$ , within specified relative error.

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## Metatheorem

Sampling combinatorial structures and estimating the associated partition function are of equivalent computational difficulty.

# Computational complexity

Viewpoint: A computational problem is *tractable* if it can be solved in a number of steps that scales as a polynomial in the size of the instance (e.g., number of vertices).

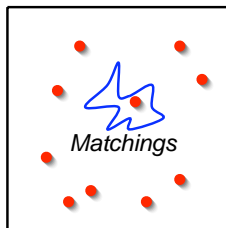
The study of counting problems was initiated by Leslie Valiant.



Leslie Valiant

- Exact evaluation of the partition function is intractable ( $\#P$ -complete) in both examples.
- Sampling configurations is tractable in one example and intractable in the other.
- Estimation of the partition function is tractable in one example and intractable in the other.

# Approach 1: Rejection sampling (Dart throwing)

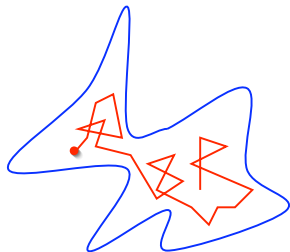


*All subsets of E*

- Until success:
  - ▶ Choose  $M \subseteq E$  uniformly at random.
  - ▶ If  $M$  is a matching, output  $M$ .

Correct distribution (for  $\lambda = 1$ ), but exponential running time.

## Approach 2: Markov chain Monte Carlo (MCMC)



- $M := \emptyset$ .
- Repeat for sufficiently many steps:
  - ▶ Choose  $e \in E$  uniformly at random, and let  $M' := M \oplus \{e\}$ .
  - ▶ If  $M'$  is a matching then  $M := M'$ ; otherwise, do nothing.
- Return  $M$ .



# Mixing time

The trial just described defines the transition probabilities  $P$  of a *Markov chain* on state space

$$\Omega = \{\text{All matchings in } G\}.$$

This Markov chain converges to a stationary distribution  $\pi$  that is uniform on  $\Omega$ .

We are interested in the *mixing time*  $\tau$  of the Markov chain, i.e., the time to convergence to near stationarity.



Andrei A. Markov

# Canonical paths/Multi-commodity flow

For every pair of states  $x, y \in \Omega$ , define a *canonical path*  $\gamma_{xy}$  from  $x$  to  $y$  using valid transitions of the MC.

“Congestion constant”  $\rho$ :

$$\sum_{\gamma_{xy} \ni (z, z')} \pi(x)\pi(y) |\gamma_{xy}| \leq \rho \pi(z)P(z, z'), \quad \forall z, z'.$$

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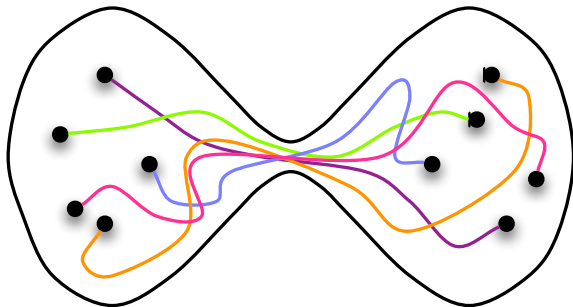
Theorem (Diaconis, Stroock;  
Sinclair)

$$\tau = O(\rho \log \pi_{\min}^{-1}).$$



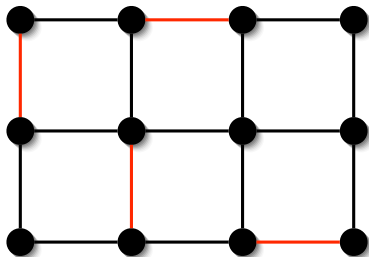
Alistair Sinclair

# Low congestion implies no “bottleneck”



## Richer set of transitions

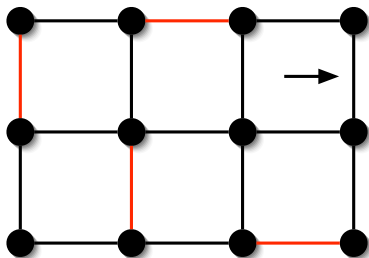
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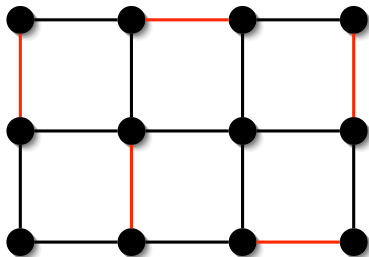


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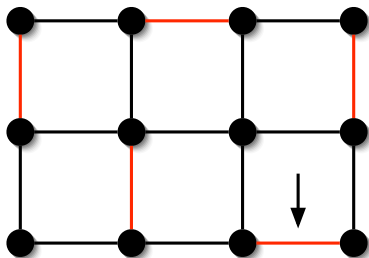
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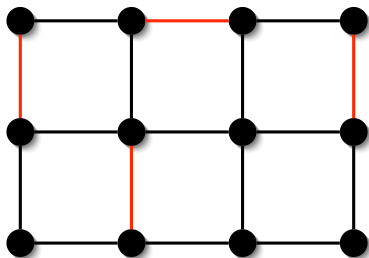
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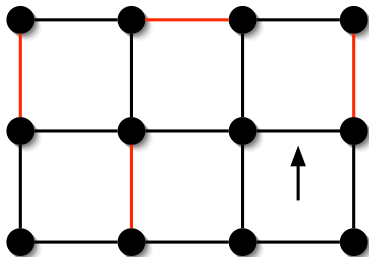
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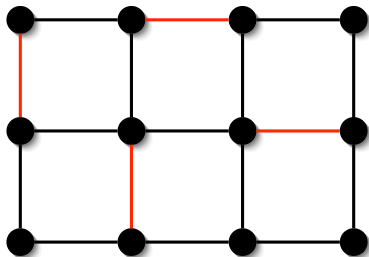
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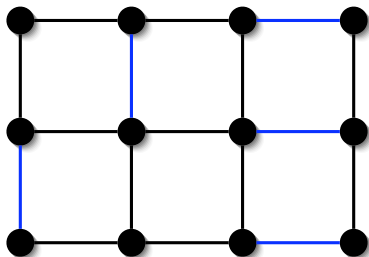
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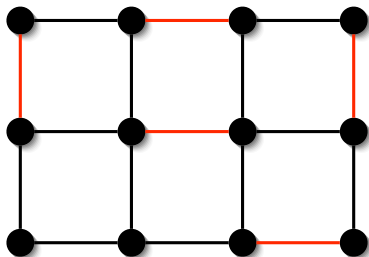
# Canonical paths for matchings

To get from the blue matching...



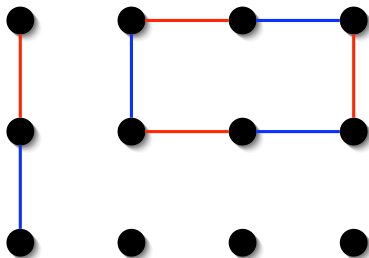
# Canonical paths for matchings

... to the red matching...



# Canonical paths for matchings

... first superimpose red and blue (symmetric difference)...

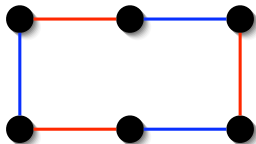


and then “unwind” each component (path or cycle).



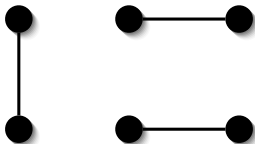
# “Unwinding” a cycle

The cycle:



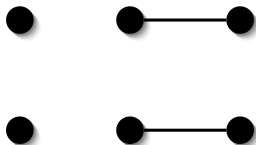
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Initial matching:



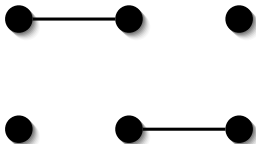
# “Unwinding” a cycle

After 1 step:



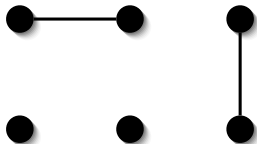
# “Unwinding” a cycle

After 2 steps:



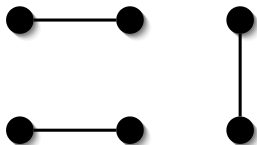
# “Unwinding” a cycle

After 3 steps:



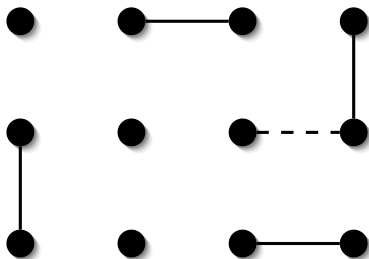
# “Unwinding” a cycle

After 4 steps (final matching):



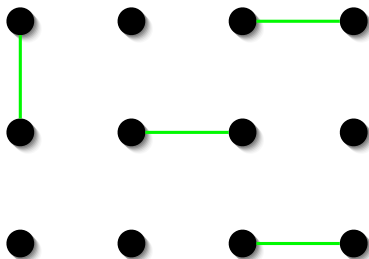
# Encoding a canonical path through a transition

A transition:



# Encoding a canonical path through a transition

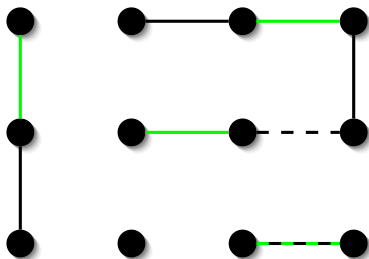
An encoding (matching):





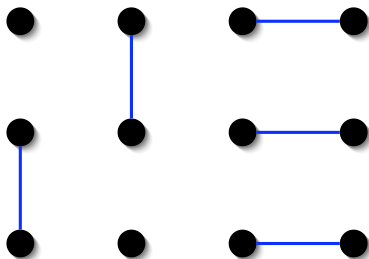
# Encoding a canonical path through a transition

Superposition reveals the initial and final matching:



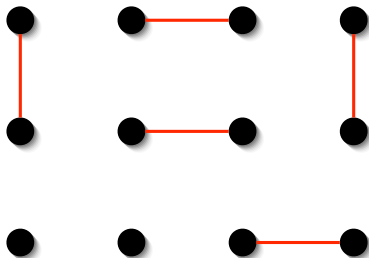
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Superposition reveals the **initial** and final matching:



# Encoding a canonical path through a transition

Superposition reveals the initial and **final** matching:



# Calculating the congestion

The encoding argument shows that the number of canonical paths passing through a given transition is roughly equal to the size of the state space.

Pursuing the calculation in more detail yields:

## Theorem (J. & Sinclair)

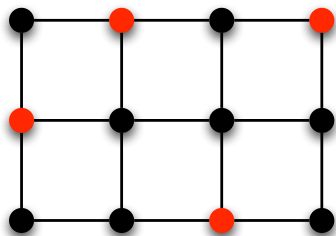
$\rho = O(nm\bar{\lambda}^2)$ , where  $n = |V|$ ,  $m = |E|$  and  $\bar{\lambda} = \max\{\lambda, 1\}$ .

## Corollary

$\tau = O(nm^2\bar{\lambda}^2)$ .

# Return to independent sets

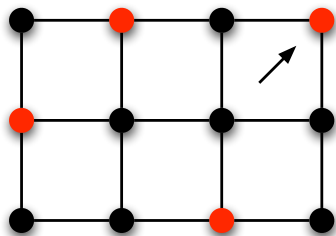
Consider an analogous set of transitions.



E.g., [Luby and Vigoda, Bubley and Dyer].

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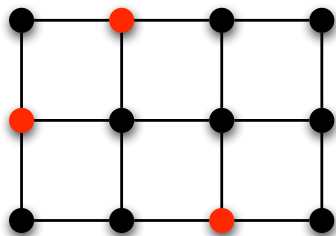
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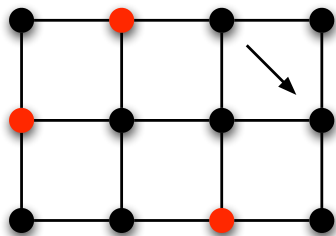
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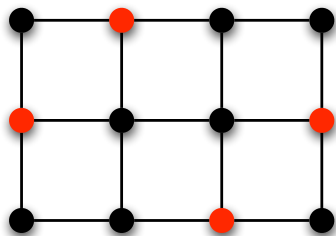


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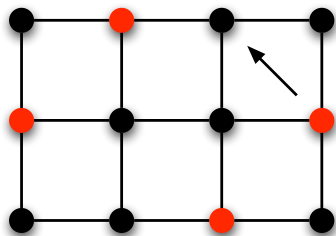
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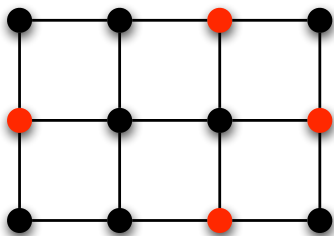
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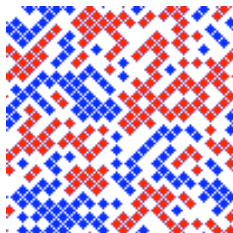
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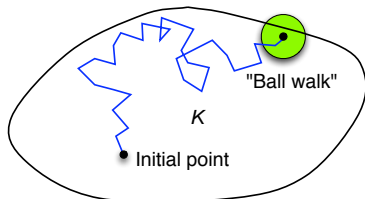
# Clustering at higher activities



- The diamonds are the vertices in the independent set; the colour red/blue indicates parity.
- At high enough  $\lambda$  there is a bias towards red or blue. (E.g., red “islands” in a blue “sea”.) This leads to a constriction in the state space.

## Example 3: sampling from a convex body

[Dyer, Frieze & Kannan, 1991], [Lovász & Simonovits, 1997].



Exploit the geometry directly (c.f. Cheeger inequality).

Mixing is rapid, provided  $K \subseteq \mathbb{R}^n$  is not “long and thin”, specifically,  
 $\tau = \text{poly}(n, \text{diam } K)$ .

Technically, surprisingly hard!

## Some other successes

- Proper colourings of a bounded degree graph, a.k.a. antiferromagnetic Potts model [Various].
- Partition function of the Ferromagnetic Ising model [J. and Sinclair].
- Bases of a “balanced” matroid [Feder and Mihail].
- Linear extensions of a partial order. [Khachiyan and Karzanov], [Bubley and Dyer].
- Feasible solutions to an instance of the knapsack problem [Morris and Sinclair].
- Perfect matchings in a bipartite graph [J., Sinclair and Vigoda].

# A renaissance

MCMC was to a close approximation the only game in town. That situation has changed in the last ten years with a rapid expansion of techniques. Can be explored in the context of the independent sets model.

Started with Weitz [2006] who exploited decay of correlations to produce a deterministic algorithm. Uses  $O(\log n)$  depth explorations from a vertex  $v$  to estimate the probability that  $v$  is in a random independent set.

## Theorem (Weitz, 2006)

*If  $\lambda < \lambda_c(d)$  then there is an FPRAS for  $Z_{IS}(\lambda)$  on graphs of maximum degree  $d$ .*

## A renaissance (continued)

Here  $\lambda_c(d)$  is the location of the phase transition (unique versus multiple Gibbs measures) in an infinite regular tree of degree  $d$ . This result prompts the daring conjecture that the phase transition in the usual **physical** sense coincides with a phase transition for **computational** tractability. Indeed:

Theorem (Sly, 2010; Galanis, Ge, Štefankovič, Vigoda and Yang, 2012; Sly and Sun, 2012)

*If  $\lambda > \lambda_c(d)$  then  $Z_{IS}(\lambda)$  is NP-hard to approximate, even for graphs of maximum degree  $d$ .*

So, remarkably, for graphs of bounded degree  $d$ , the critical value  $\lambda_c(d)$  is also the boundary between computational tractability and intractability.



## A renaissance (continued)

Another significant line of attack was introduced by Barvinok, and taken forward by Patel and Regts.

The idea is to consider a Taylor expansion of  $\log Z$  about some point where the partition function is easy to evaluate. Then by enumerating small substructures, one can estimate the first few coefficients in the expansion. This in turn gives a good estimate for  $Z$  at a hard-to-evaluate point.

For this to work we need there to be no zeros in the neighbourhood in which we are performing the Taylor expansion.

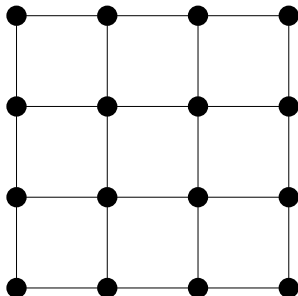
Zeros of the partition function  $Z$  are associated with phase transitions: another connection to the physics of spin systems!

# Independent sets by Partial Rejection Sampling

A further recent approach to efficient sampling is *Partial rejection sampling* [J. & Guo, 2017]:

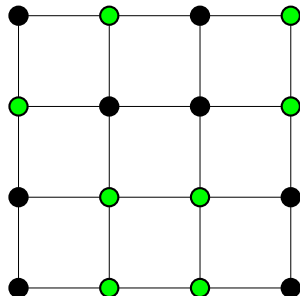
# Independent sets by Partial Rejection Sampling

1. Randomize each vertex (in/out). Consider the connected components induced by the in-vertices.
2. Let **Bad** be the set of vertices in connected components of size at least 2.
3. **Resample** = **Bad**  $\cup$   $\partial$ **Bad**.
4. Resample variables in set **Resample**. Check independence.



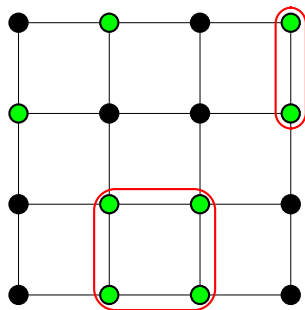
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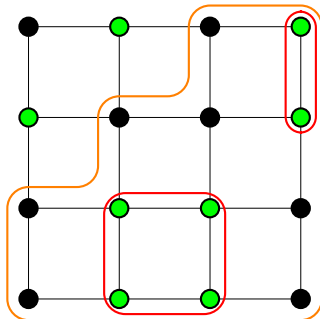
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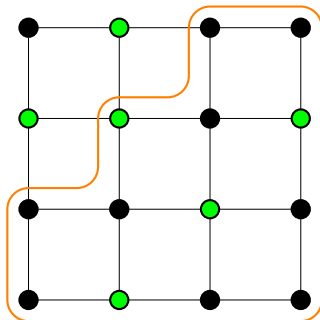
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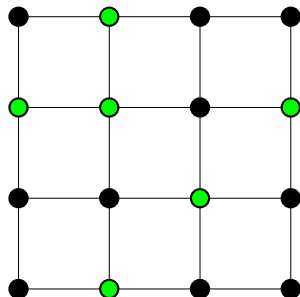
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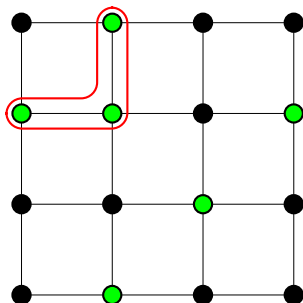
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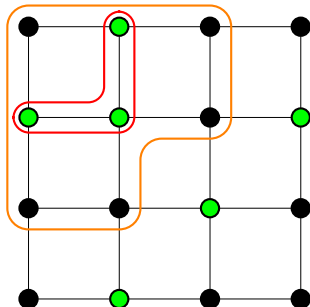
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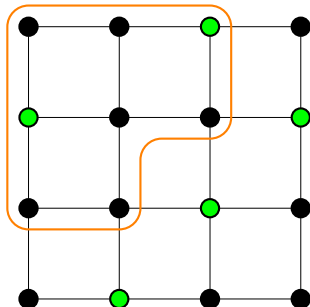
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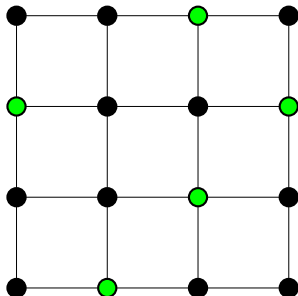
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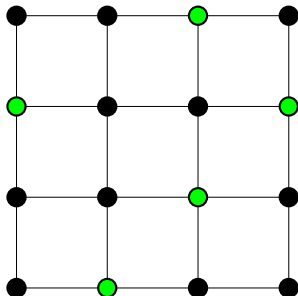
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3. **Resample** = **Bad**  $\cup$   $\partial$ **Bad**.
- 4. Resample variables in set **Resample**. Check independence.



# Independent sets by Partial Rejection Sampling

1. Randomize each vertex (in/out). Consider the connected components induced by the in-vertices.
2. Let **Bad** be the set of vertices in connected components of size at least 2.
3. **Resample** = **Bad**  $\cup$   $\partial$ **Bad**.
4. Resample variables in set **Resample**. Check independence.



When the algorithm stops, it yields a uniform independent set.

# Unblocking sets

The key property of the set **Resample** is that it is **unblocking** under the current assignment  $\sigma$  to the variables.

## Definition

A set  $\mathcal{U}$  of variables is **unblocking** under  $\sigma$  if  $\sigma[\mathcal{U}]$  determines the truth value of all clauses that *share* variables with  $\mathcal{U}$  (and not just the clauses containing *only* variables from  $\mathcal{U}$ )

The resampling set from the independent set example was unblocking.

In applications we also require that  $\mathcal{U}$  is “adapted” to  $\sigma$ .

# Partial rejection sampling

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## Algorithm 1 Partial Rejection Sampling

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PRS( $V, \Phi$ ) //  $\Phi$  is a formula on variable set  $V$

Sample, from the product distribution, an assignment  $\sigma$  to the variables in  $V$

**while**  $\text{Bad}(\sigma) \neq \emptyset$  **do**

$S \leftarrow \bigcup \{\text{var}(C) : C \in \text{Bad}(\sigma)\}$

    Resample all variables in  $U = \text{Resample}(S; \sigma)$

**end while**

---

Note 1. The procedure **Resample** must not probe variables outside of  $U$  while computing  $U$ . This is the condition of being adapted.

Note 2. A previous approach to rejection sampling that tries to preserve randomness is the “Randomness Recycler” of Fill and Huber.

# A selection of open problems

- Is there a polynomial-time algorithm for sampling perfect matchings in a *general* graph?
- Is there an algorithm for sampling perfect matchings in a bipartite graph that is efficient in practice?
- Is there a polynomial-time algorithm for sampling contingency tables?
- Can one sample proper colourings efficiently when  $q \geq (1 + \varepsilon)d$ ? ( $q$  is the number of colours and  $d$  the maximum degree.)
- Is the bases-exchange walk rapidly mixing for all matroids?