



## Markov chains, mixing time and connections with reconfiguration

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Combinatorial Reconfiguration workshop, Banff

### **Reconfiguration:**

(my current understanding, based on Nishimura's 2018 survey)

We have

- a (finite?) set  $\Omega$  of solutions to a problem instance
- a notion of **adjacency** of solutions
- the **reconfiguration graph**  $\mathcal{G}$  with vertex set  $\Omega$  and edges given by **adjacency**

Usually (I imagine)  $\Omega$  is **exponentially large** as a function of the "size"  $n$  of the problem instance.

**Reconfiguration:** many structural questions

- **reachability:**  
Given  $X, Y \in \Omega$ , is there a path from  $X$  to  $Y$  in  $\mathcal{G}$ ?
- **connectivity:** Is  $\mathcal{G}$  connected?
- **shortest paths:**  
Given  $X, Y \in \Omega$ , what is length of  
shortest path from  $X$  to  $Y$  in  $\mathcal{G}$ ?
- **diameter:** What is the diameter of  $\mathcal{G}$ ?

**Reconfiguration:** many algorithmic questions, e.g.

- algorithms to decide **reachability**, **connectivity**,
- algorithms to find **shortest paths** or calculate **diameter**
- find “**best**” solution reachable from a given starting point

Related questions of **computational complexity**, **parameterised complexity** etc. Even **reachability** might be intractable!



**Markov chains:** (discrete time, finite state space)

We have

- a finite set  $\Omega$ , called a state space
- some allowed transitions, with the next transition chosen randomly according to some rule
- a (directed) graph  $\mathcal{G}$  stores the set of all possible transitions

(We'll be more precise soon.)

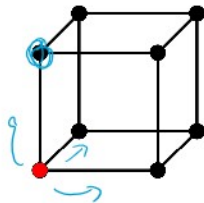
Usually  $\Omega$  is exponentially large as a function of some parameter  $n$ .

A Markov chain  $\mathcal{M}$  on state space  $\Omega$  is a stochastic process  $X_0, X_1, X_2, \dots$  which is memoryless:

$$\Pr(X_{t+1} = y \mid X_0 = x_0, \dots, X_t = x_t) = \Pr(X_{t+1} = y \mid X_t = x_t)$$

for all  $t \in \mathbb{N}$  and  $x_0, \dots, x_t, y \in \Omega$ .

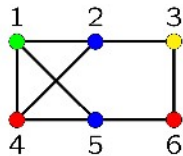
Here each  $X_t = X_t(n)$  is a random element of a set  $\Omega = \Omega_n$  which is usually finite but exponentially large with respect to some parameter  $n$ .



Example: A Markov chain for graph colourings.

Let  $k \geq 3$  be a fixed integer and let  $G = (V, E)$  be a graph. Write  $[k] = \{1, 2, \dots, k\}$ .

A  $k$ -colouring of  $G$  is a function  $\sigma : V \rightarrow [k]$  such that if  $\{x, y\} \in E$  then  $\sigma(x) \neq \sigma(y)$ .

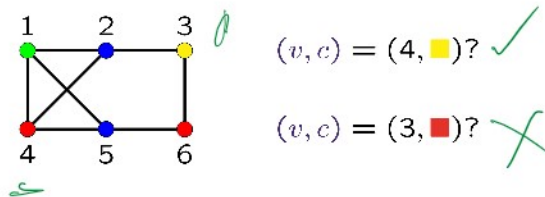


Let  $\Omega_k(G)$  be the set of all  $k$ -colourings of  $G$ .

A simple Markov chain on  $\Omega_k(G)$  has the following transition procedure: from the current state  $X \in \Omega_k(G)$  do

- Choose  $v \in V$  uniformly at random,
- Choose  $c \in [k]$  uniformly at random,
- Recolour  $v$  with  $c$  to give a new state  $X'$ , if possible. Otherwise, let  $X' = X$ .

This is the Glauber dynamics or single-site update chain.



We say that the Markov chain  $\mathcal{M}$  is **time-homogeneous** if the transition probabilities from a given state  $x$  are **independent of  $t$** .

For a **time-homogeneous** chain, the transition probabilities can be stored in the transition matrix  $P$  of  $\mathcal{M}$ , with **rows and columns indexed by  $\Omega$**  and entries

$$P(x, y) = \Pr(X_{t+1} = y \mid X_t = x).$$

Matrix  $P$  is **stochastic**: all rows sum to 1.

The matrix  $P$  is **too large** to work with directly. (e.g. can't find **eigenvalues in polynomial time**).

**AIM: sampling  $\Omega$**

We can also define the **directed graph  $\mathcal{G}$**  underlying the Markov chain, with vertex set  $\Omega$  and  $(x, y)$  a directed edge if and only if  $P(x, y) > 0$ , for all  $x, y \in \Omega$ .

A **self-loop** is a directed edge  $(x, x)$  in  $\mathcal{G}$ .



[Similar to "adjacency", "reconfiguration graph".]

If the state space  $\Omega$  is **connected under moves of  $\mathcal{M}$**  then we say that  $\mathcal{M}$  is **irreducible**.

[This is (strong) "connectivity of the reconfiguration graph".]

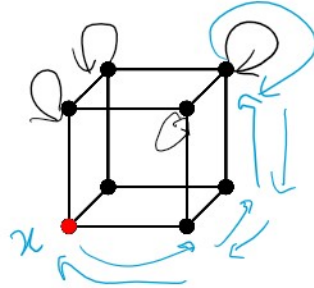
We don't usually worry about **shortest paths, diameter**.

Write  $P_x^t(\cdot)$  for the distribution of state  $X_t$  of the Markov chain  $\mathcal{M}$ , after  $t$  steps from initial state  $X_0 = x$ .

Say  $\mathcal{M}$  is **aperiodic** if for all  $x \in \Omega$ ,

$$\gcd\{t \in \mathbb{Z}^+ \mid P_x^t(x) > 0\} = 1.$$

One self-loop is sufficient to guarantee aperiodicity, if  $\mathcal{M}$  is **irreducible**.



### Classical Markov chain theory

A row-vector  $\pi$  with nonnegative entries which add up to 1 is a **stationary distribution** of Markov chain  $\mathcal{M}$  if

$$\pi P = \pi.$$

We say that Markov chain  $\mathcal{M}$  is **ergodic** if it is **irreducible** and **aperiodic**.

If  $\mathcal{M}$  is ergodic then  $\mathcal{M}$  has a **unique stationary distribution**  $\pi$  such that  $\pi(x) > 0$  for all  $x \in \Omega$ , and

$$\lim_{t \rightarrow \infty} P_x^t(y) = \pi(y)$$

for all  $x, y \in \Omega$ .

See for example: book by Levin, Peres & Wilmer (2009).



How to find  $\pi$ ?

We say that Markov chain  $\mathcal{M}$  satisfies the **detailed balance equations** with respect to row vector  $\psi$  if

$$\psi(x) P(x, y) = \psi(y) P(y, x) \quad \text{for all } x, y \in \Omega.$$

We also say that  $\mathcal{M}$  is **reversible**, or **time-reversible**, with respect to  $\psi$ .

If  $\mathcal{M}$  is **ergodic** and satisfies **detailed balanced** with respect to some vector  $\psi \neq \mathbf{0}$ , then the **unique stationary distribution**  $\pi$  is given by **normalising**  $\psi$ .

In this case, the underlying graph  $\mathcal{G}$  is really **undirected**.

eg **reconfiguration graph**.

Example: A Markov chain for **graph colourings**.

Fact: The **Glauber dynamics**  $\mathcal{M}_k(G)$  is **irreducible** on  $\Omega_k(G)$  when  $k \geq \Delta(G) + 2$ , where  $\Delta(G)$  is the **maximum degree** of  $G$ .

Also  $\mathcal{M}_k(G)$  is **aperiodic** as  $P(x, x) \geq 1/k$  for any  $x \in \Omega_k(G)$ . So  $\mathcal{M}_k(G)$  is **ergodic**.

The transition probabilities satisfy

$$P(x, y) = \frac{1}{kn} = P(y, x)$$

whenever  $x, y \in \Omega_k(G)$  differ on a **single vertex**, and  $P(x, y) = 0$  for all other  $x \neq y$ .

The chain satisfies **detailed balance** with respect to  $(1, 1, \dots, 1)$ , so its **stationary distribution** is **uniform**.

We use the **detailed balanced** equations to **design** our Markov chains, so they have the desired stationary distribution.

In particular, if  $\pi$  is uniform then  $P$  must be **symmetric**.

We also need to prove **irreducibility**; that is, the underlying graph  $\mathcal{G}$  is **connected**.

**Aperiodicity** is easy: just ensure  $P(x, x) > 0$  for all  $x \in \Omega$ .

Now we have a **very well behaved** Markov chain which **converges** to its stationary distribution.... eventually.

**Q: How quickly?**

For probability distributions  $\sigma, \mu$  on  $\Omega$ , the total variation distance between  $\sigma$  and  $\mu$  is

$$d_{TV}(\sigma, \mu) = \frac{1}{2} \sum_{x \in \Omega} |\sigma(x) - \mu(x)|. = \max_{A \subseteq \Omega} \sigma(A) - \mu(A).$$

Define the **mixing time**  $\tau(\varepsilon)$  of  $\mathcal{M}$  by

$$\tau(\varepsilon) = \max_{x \in \Omega} \min\{t \mid d_{TV}(P_x^t, \pi) < \varepsilon\}.$$

Convergence is monotonic.

Here  $\varepsilon > 0$  is a small **user-defined tolerance**.

We say that the Markov chain  $\mathcal{M}$  is **rapidly mixing** if  $\tau(\varepsilon)$  is bounded above by some **polynomial in  $\log |\Omega|$  and  $\log(\varepsilon^{-1})$** .



This is a strong condition: we want to get exponentially close to the stationary distribution, over an exponentially large state space, in polynomial time.

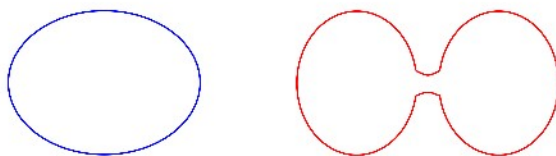
Linear algebra  $\Rightarrow$  eigenvalues of  $P$  control convergence of  $\mathcal{M}$ . But  $\Omega$  is too big to allow direct computation of the eigenvalues in polynomial time.

Other methods:

- coupling, path coupling [Probabilistic]
- geometric arguments [\*]
- functional inequalities,  
e.g. Poincaré inequality, log-Sobolev inequality

### Geometry of the state space

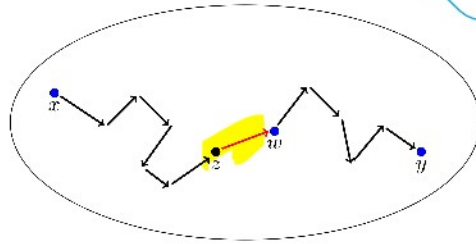
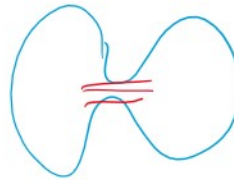
Which state space do you think encourages rapid mixing?



Constrictions in the state space make it difficult for the chain to escape: exponential mixing time!

Lack of constrictions allows chain to mix freely. Results by Jerrum & Sinclair (1987) make this precise: conductance.

Method ~~1~~: Canonical paths



- For all pairs  $(x, y) \in \Omega^2$ , define a path  $\gamma_{xy}$  from  $x$  to  $y$ , where each step is a transition of the Markov chain.
- Analyse the congestion of the set of all paths: are any transitions heavily loaded? Then apply Sinclair (1992).

NB: Canonical paths might not be shortest paths.

Instead, we want to avoid having too many paths going through the same edge of  $\mathcal{G}$ .

(These conditions almost seem orthogonal!?, somehow??)

Let  $\mathcal{M}$  be time-homogeneous, ergodic, reversible Markov chain with  $N = |\Omega|$ . The eigenvalues of the transition matrix  $P$  are real and satisfy

$$1 = \lambda_0 > \lambda_1 \geq \dots \geq \lambda_{N-1} > -1$$

and the mixing time of  $\mathcal{M}$  is controlled by

$$\lambda_{\max} = \max\{\lambda_1, |\lambda_{N-1}|\}.$$

If  $\pi^* = \min\{\pi(x) \mid x \in \Omega\}$  then

$$\tau(\varepsilon) \leq (1 - \lambda_{\max})^{-1} (\log(1/\pi^*) + \log(\varepsilon^{-1})).$$

See Sinclair (1992).

The quantity  $(1 - \lambda_{\max})^{-1}$  is the relaxation time of  $\mathcal{M}$ .

$$\frac{1}{1 - \lambda_1} \quad \text{or} \quad \frac{1}{1 - |\lambda_{N-1}|} = \frac{1}{1 + \lambda_{N-1}}$$

Typically  $\lambda_{\max} = \lambda_1$ . This can be guaranteed by making the chain lazy, that is, replacing  $P$  with  $(I + P)/2$ .

However, a method of Diaconis & Saloff-Coste (1993) can be applied directly to bound  $(1 + \lambda_{N-1})^{-1}$ , without resorting to laziness.

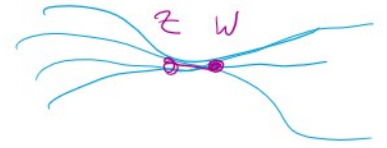
As a special case, if every state has a self-loop then

$$(1 + \lambda_{N-1})^{-1} \leq \frac{1}{2} \max_{x \in \Omega} P(x, x)^{-1}.$$

The point is, we can focus on  $\lambda_1$ .

The quantity  $1 - \lambda_1$  is called the spectral gap.

Sinclair (1992): Let  $\mathcal{M}$  be time-homogenous, ergodic and reversible with stationary distribution  $\pi$ .



Let  $\Gamma = \{\gamma_{xy} \mid x, y \in \Omega\}$  be a set of canonical paths for  $\mathcal{M}$ . Define the congestion

$$\bar{\rho} = \bar{\rho}(\Gamma) = \max_{z, w \in E(G)} \frac{1}{\pi(z)P(z, w)} \sum_{\gamma_{xy} \ni z, w} \pi(x)\pi(y) |\gamma_{xy}|$$

where  $|\gamma_{xy}|$  is the length of the path  $\gamma_{xy}$ .

Then

$$(1 - \lambda_1)^{-1} \leq \bar{\rho}.$$

$$\begin{aligned} \pi(z)P(z, w) &= \pi(w)P(w, z) \\ &= Q(z, w) \end{aligned}$$

So we want an upper bound on  $\bar{\rho}$  which is polynomial in  $\log |\Omega|$ .

**Theorem (Sinclair, 1992)**

Suppose that  $\mathcal{M}$  is time-homogenous, ergodic and reversible, and let  $\Gamma$  be a set of canonical paths for  $\mathcal{M}$ . Then

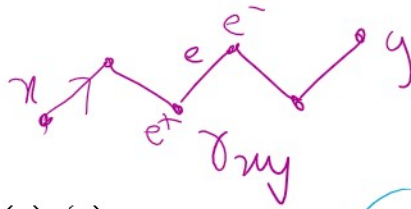
$$(1 - \lambda_1)^{-1} \leq \bar{\rho}.$$

**Proof.** Let  $L = I - P$ , so that the eigenvalues of  $L$  are  $\mu_i = 1 - \lambda_i$ . The variational characterisation of  $\mu_1$  is

$$\mu_1 = \inf_{\psi} \frac{\sum_{x, y \in \Omega} (\psi(x) - \psi(y))^2 \pi(x)P(x, y)}{\sum_{x, y \in \Omega} (\psi(x) - \psi(y))^2 \pi(x)\pi(y)}$$

smallest positive eigenvalue of  $L$

with the infimum taken over all non-constant functions  $\psi : \Omega \rightarrow \mathbb{R}$ .



Now

Denom of expression

$$\begin{aligned} & \sum_{x,y} (\psi(x) - \psi(y))^2 \pi(x)\pi(y) \\ &= \sum_{x,y} \pi(x)\pi(y) \left( \sum_{e \in \gamma_{xy}} |\psi(e^+) - \psi(e^-)| \right)^2 \\ &\leq \sum_{x,y} \pi(x)\pi(y) |\gamma_{xy}| \sum_{e \in \gamma_{xy}} (\psi(e^+) - \psi(e^-))^2, \end{aligned}$$

telescoping sum along canonical path

writing  $e = e^-e^+$  for each  $e \in \gamma_{xy}$ . The final line uses the Cauchy-Schwarz inequality.

Exchanging the order of summation gives

$$\begin{aligned} & \sum_{x,y} (\psi(x) - \psi(y))^2 \pi(x)\psi(y) \\ &\leq \sum_e (\psi(e^+) - \psi(e^-))^2 \sum_{\gamma_{xy} \ni e} \pi(x)\pi(y) |\gamma_{xy}| \\ &\leq \sum_e (\psi(e^+) - \psi(e^-))^2 \pi(e^-) P(e^-, e^+) \bar{\rho}(\bar{\Gamma}) \\ &= \bar{\rho}(\bar{\Gamma}) \sum_{x,y} (\psi(x) - \psi(y))^2 \pi(x) P(x, y), \end{aligned}$$

which implies that

$$1 \leq \bar{\rho}(1 - \lambda_1),$$

completing the proof.  $\square$

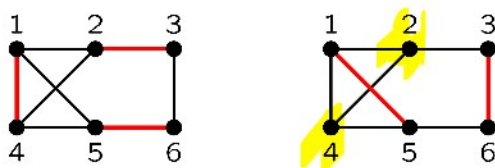
numerator

Example:

A Markov chain for **perfect and near-perfect matchings**.

Let  $G = (V, E)$  be a graph. A **matching** in  $G$  is a set of edges  $M \subseteq E$  such that no vertex is incident with **more than one edge of  $M$** .

If  $|V| = 2n$  and  $|M| = n$  then  $M$  is a **perfect matching**. Matchings with  $n - 1$  edges are called **near-perfect**.



Let  $\mathcal{P}$  be the set of all **perfect matchings** of  $G$  and let  $\mathcal{N}$  be the set of all **near-perfect matchings** of  $G$ . **Broder (1986)** introduced a Markov chain with state space  $\mathcal{P} \cup \mathcal{N}$ .

From current state  $M \in \mathcal{P} \cup \mathcal{N}$ ,

- with probability  $\frac{1}{2}$  let  $M' = M$ ; otherwise: *lazy*
  - Choose  $e = uv \in E(G)$  **uniformly at random**.
  - If  $M \in \mathcal{P}$  and  $e \in M$  then let  $M' = M \setminus e$ .



- If  $M \in \mathcal{N}$  and both  $u, v$  are **unmatched** in  $M$  then let  $M' = M + e$ .



- If  $M \in \mathcal{N}$  and **exactly one** of  $u, v$  are **matched** in  $M$ , with matching edge  $e'$ , then let  $M' = (M \setminus e') + e$ .



The **Broder chain** is **irreducible** (honest!), and **aperiodic** (in fact it is **lazy**).

If  $P(M, M')$  is nonzero for distinct  $M, M'$  then

$$P(M, M') = \frac{1}{2m} = P(M', M),$$

where  $m$  is the number of edges of  $G$ .

So the stationary distribution is **uniform**.

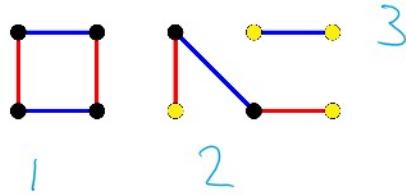
**Broder (1986)** applied **coupling** to try to analyse this chain, but his proof was **incorrect**. **Jerrum & Sinclair (1989)** used **canonical paths**.



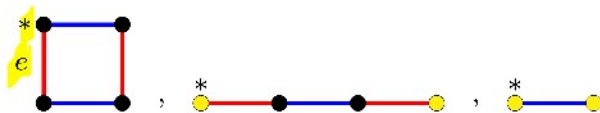
Let  $M, M'$  be two elements of  $\mathcal{P} \cup \mathcal{N}$ .



The symmetric difference  $M \oplus M'$  is the disjoint union of cycles and at most two paths.



We process the components of  $M \oplus M'$  in some canonical order, and let the start vertex of each cycle (respectively, path) be the least labelled vertex (respectively, endvertex) in the cycle/path.



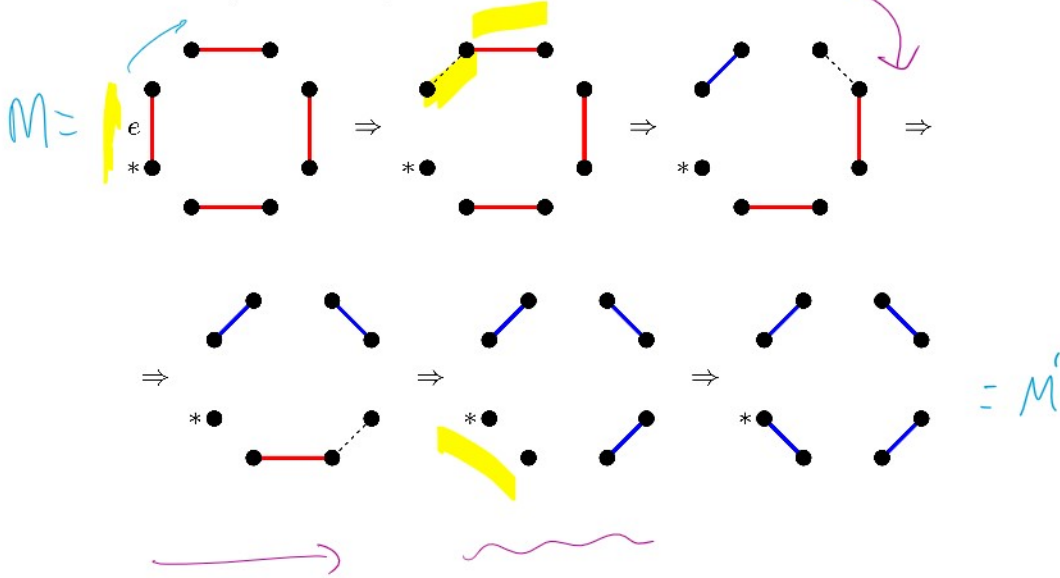
Denote by  $e = e(C)$  the edge of  $M$  incident with the start-vertex of each cycle  $C$ .

With each component we process, we add a few more steps to our canonical path from  $M$  to  $M'$ .



$M$  is a perfect matching

How to process a cycle:



Paths are processed similarly. This defines a **canonical path**  $\gamma_{XY}$  between each pair  $(X, Y) \in \mathcal{P} \cup \mathcal{N}$ .

Now we must **analyse** the set of canonical paths  $\Gamma = \{\gamma_{XY}\}$ . What is the **load** on each transition?

**Key tool:** define the **encoding**  $\eta_t(X, Y)$  of the transition  $t$  on the canonical path  $\gamma_{XY}$ , such that if you know  $t = (M, M')$  and  $\eta_t$  then you can **uniquely recover**  $(X, Y)$ .



Encodings used in **Jerrum & Sinclair (1989)**:

If  $t = (M, M')$  where  $M, M' \in \mathcal{N}$  and we are processing a cycle, then

$$\eta_t(X, Y) = (X \oplus Y \oplus (M \cup M')) \setminus \{e\},$$

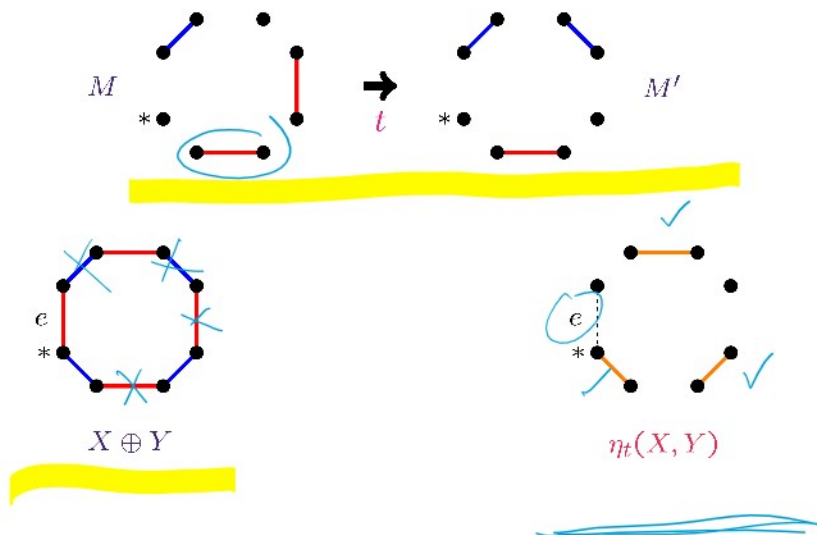
where  $e$  is the edge of  $X$  adjacent to the start-vertex of the cycle; and otherwise

$$\eta_t(X, Y) = X \oplus Y \oplus (M \cup M').$$

**Q:** Why remove  $e$ ?

**A:** This ensures that  $\eta_t(X, Y)$  belongs to  $\mathcal{P} \cup \mathcal{N}$ .

Example:  $\eta_t(X, Y) = (X \oplus Y \oplus (M \cup M')) \setminus \{e\}$



Facts:

- The encoding  $\eta_t(X, Y)$  is either a perfect or near-perfect matching of  $G$ .
- If you know  $t = (M, M')$  and  $\eta_t$  then you can uniquely recover  $(X, Y)$ .

Hence no transition lies on more than  $|\mathcal{P} \cup \mathcal{N}|$  canonical paths.

Also, if  $G$  has  $2n$  vertices then  $|\gamma_{xy}| \leq 2n$  for all  $X, Y \in \mathcal{P} \cup \mathcal{N}$ .



Thus

$$\begin{aligned} \bar{\rho} &= \max_{zw \in E(G)} \frac{1}{\pi(z)P(z, w)} \sum_{\gamma_{xy} \ni zw} \pi(x)\pi(y) |\gamma_{xy}| \\ &\leq |\mathcal{P} \cup \mathcal{N}| 2m \cdot |\mathcal{P} \cup \mathcal{N}| \cdot \frac{2n}{|\mathcal{P} \cup \mathcal{N}|^2} \\ &= 4mn. \end{aligned}$$

Also

$$1/\pi^* = |\mathcal{P} \cup \mathcal{N}| \leq (2n)! \leq \exp(2n \log n).$$

Jerrum & Sinclair (1989): The Broder chain has mixing time

$$\tau(\varepsilon) \leq 4mn(2n \log n + \log(\varepsilon^{-1}))$$

where  $G$  has  $n$  vertices and  $m$  edges.

The canonical path method is a thing of beauty, when it works. But...

- \* Usually, it does NOT lead to tight bounds, and
- \* Finding a good set of canonical paths can be tricky.

Huang, Lu & Zhang (SODA 2016),  
 "Canonical paths for MCMC: from art to science".

They build on work of McQuillan (2013) to reduce the task of designing canonical paths to solving a system of linear equations.



Sinclair (1992) introduced the multicommodity flow method, where  $\pi(x)\pi(y)$  units of flow are split across a set of paths from  $x$  to  $y$  in  $\mathcal{G}$ .



This method has been used to analyse the switch chain for sampling graphs (or directed graphs or bipartite graphs) with given degree sequence, under various conditions.

Transitions:

From a given graph, choose a pair of non-incident edges randomly and replace them by a (randomly chosen) pair of edges without changing the degree sequence.



References:

- Several chapters by Mark Jerrum  
 "Counting, Sampling and Integrating: Algorithms and Complexity" (2003) *Monograph*
- Levin, Peres & Wilmer,  
 "Markov Chains and Mixing Times" (2009) *Book*

- Sinclair, Improved bounds for mixing rates of Markov chains and multicommodity flow," CPC **1** (1992), 351-370
- 

The switch chain is **ergodic** and its stationary distribution is **uniform**.

The switch chain has been shown to be **rapidly mixing** in various situations, by several authors, using **multicommodity flow**. (See for example my BCC 2021 talk.)

But the mixing time bounds are **just awful**, e.g. **Cooper, Dyer, Greenhill (2007)** proved that

$$\tau(\varepsilon) \leq d^{23} n^8 (dn \log(dn) + \log(\varepsilon^{-1}))$$

for  $d$ -regular graphs on  $n$  vertices.

**Q:** Is it really this bad?

**A:** Maybe not.

**Tikhomirov & Youssef, arXiv.2007.02729** proved a mixing time bound of

$$C dn (dn \log dn + \log(2\varepsilon^{-1}))$$

for the **switch chain** on  $d$ -regular **bipartite** graphs, where  $3 \leq d \leq n^c$ , for some constants  $c, C > 0$ .

This is a **huge** improvement on any previously-known bound.

Proof is **long & technical**, involves establishing a **Poincaré inequality** to bound the eigenvalues of the chain **directly**.

## Connections with approximate counting

If a problem is “self-reducible” then **approximate counting** can be reduced to **approximately uniform sampling**, e.g. using Markov chains.

Other approaches to **approximate counting** using **deterministic algorithms**:

- **correlation decay method** (Weitz, 2006)
- **polynomial interpolation method** (Barvinok, 2016)

Some related **computational complexity** questions, mostly related to **approximate counting**:

**Q:** Is the counting problem **#P-complete**?

**Dyer, Goldberg, Greenhill, Jerrum (2003)** defined **approximation-preserving reductions** (AP-reductions) and identified 3 classes of **approximate counting problems**:

- solvable in **randomized polynomial time** (RP),
- AP-interreducible with **#SAT**,
- AP-interreducible with **#BIS**

Here **#BIS** is the problem of counting **independent sets** in bipartite graphs.

\* **Thank you!** \*