

# Perfect Simulation and Non-monotone Markovian Systems

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# Discrete Event System

System description:  $(\mathcal{X}, \pi^0, \mathcal{E}, p, \phi)$

- ▶ Finite state space  $\mathcal{X}$ .

Without loss of generality,  $\mathcal{X} = \{1, \dots, N\}$ .

- ▶ Probability measure  $\pi^0$  on  $\mathcal{X}$ :

$\pi_x^0 \geq 0$ ,  $x \in \mathcal{X}$  is the probability that the system is in state  $x$  at time 0.

- ▶ Finite set of events  $\mathcal{E}$ .

- ▶ Probability measure  $p$  on  $\mathcal{E}$ :

$p_e > 0$ ,  $e \in \mathcal{E}$  is the probability of event  $e$ .

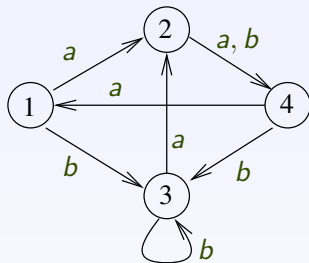
- ▶ Transition function  $\phi : \mathcal{X} \times \mathcal{E} \rightarrow \mathcal{X}$ .

# Discrete Event System (II)

Evolution of the system (over  $n$  steps):

1. Choose initial state  $X_0$  with probability measure  $\pi^0$ .
2. For  $i = 1$  to  $n$  do:
  - ▶ Choose an event  $e_i \in \mathcal{E}$  with probability measure  $p$
  - ▶  $X_i := \phi(X_{i-1}, e_i)$

## Example



Let  $p_a = 1/3$ ,  $p_b = 2/3$ , and  $\pi^0 = (1/4, 1/4, 1/4, 1/4)$ .

A possible **trajectory** of the system is

$1 - 3 - 3 - 2 - 4 - 1 - 3 - 3 - \dots$  starting from state 1 and for sequence of events  $bbababb \dots$

## Remarks

Random sequence  $\{X_n\}_{n \in \mathbb{N}}$  is a discrete time Markov chain (DTMC) with transition probability matrix:

$$P_{i,j} \stackrel{\text{def}}{=} \mathbb{P}(X_n = j | X_{n-1} = i) = \sum_{e \in \mathcal{E}} p_e \mathbf{1}_{\phi(i,e)=j}.$$

Furthermore, every DTMC can be represented in a form  $(\mathcal{X}, \pi^0, \mathcal{E}, p, \phi)$ . For a chain with  $N$  states, we can construct an event representation with at most  $N^2$ , with complexity  $O(N^2)$ .

# Sampling the Steady-state

Assumption:  $\{X_n\}_{n \in \mathbb{N}}$  is ergodic.

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Complexity of computing  $\pi$ :  $O(N^3)$  (where  $N = |\mathcal{X}|$ ).

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## Question

*How to avoid computing  $\pi$ ?*



# Monte-Carlo Simulation

Algorithm:

- ▶ Sample  $X_0$  from  $\pi^0$ .
- ▶ For  $i = 1$  to  $n$ :
  - ▶ Sample  $e_i$  from  $p$ .
  - ▶  $X_i = \phi(X_{i-1}, e_i)$ .

Output: a sample from the probability measure  $\pi^0 P^n$ .

Complexity:  $O(\mathcal{C}(\phi)n)$ .

(Remark: sampling from discrete probability measure can be done in  $O(1)$  using alias method [Walker, 74].)

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Inconvenient: approximation.

Error estimation is difficult: depends on the second eigenvalue of  $P$  which is hard to compute [Brémaud, Glynn, Whitt, Hordijk].

# Perfect Simulation

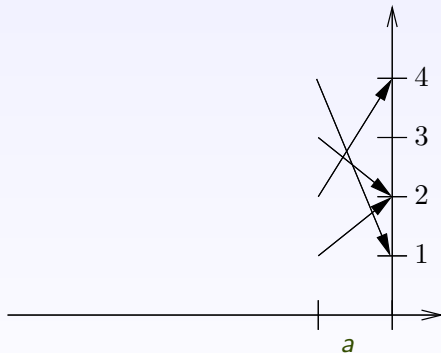
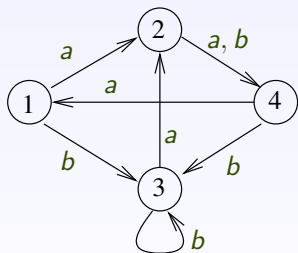
Goal:

- ▶ unbiased samples of  $\pi$  without computing it (nor  $P$ ).
- ▶ finite stopping time.

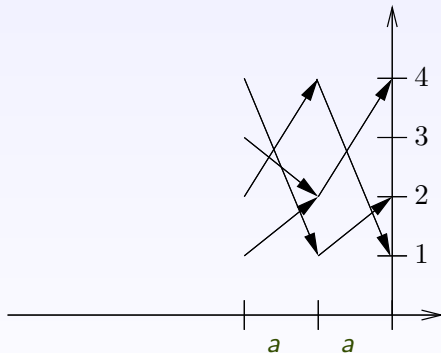
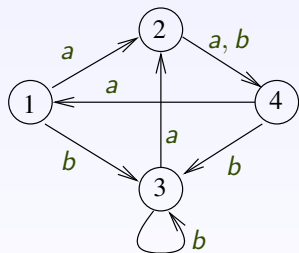
First results (theoretical and existential) [Borovkov 75, Glynn 96]

Propp and Wilson (1996) proposed backward coupling algorithm.

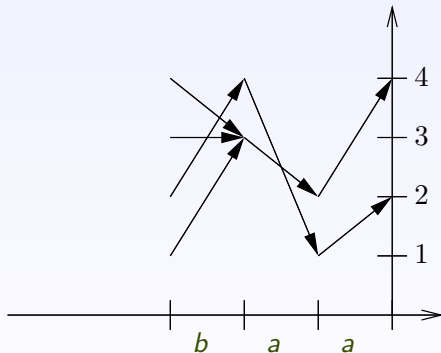
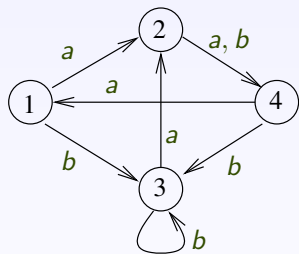
# Backward coupling



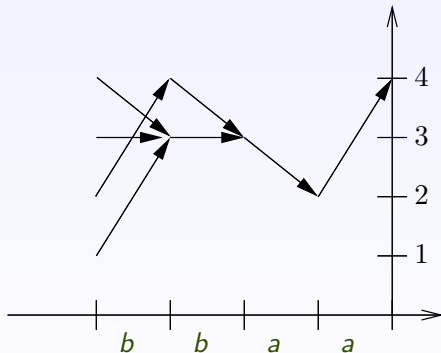
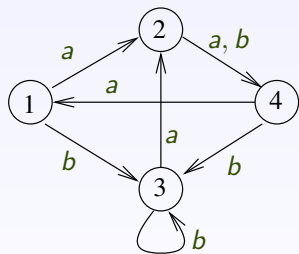
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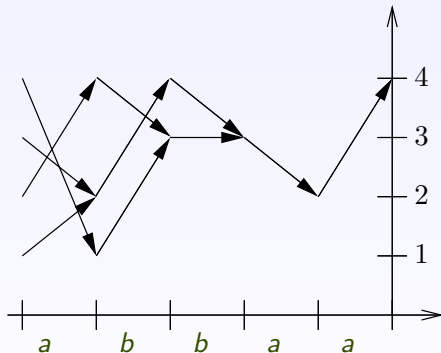
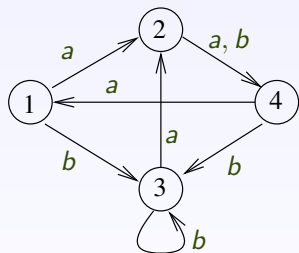
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## Backward coupling (II)

$$\Phi^n(x, e_{1 \rightarrow n}) \stackrel{\text{def}}{=} \Phi(\dots \Phi(\Phi(x, e_1), e_2), \dots, e_n).$$

$$\text{For } A \subset \mathcal{X}, \Phi^n(A, e_{1 \rightarrow n}) \stackrel{\text{def}}{=} \{\Phi^n(x, e_{1 \rightarrow n}), x \in A\}.$$

Theorem ([Propp and Wilson (1996)])

*There exists  $\ell \in \mathbb{N}$  such that*

$$\lim_{n \rightarrow \infty} |\Phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})| = \ell \text{ almost surely.}$$

*The system couples if  $\ell = 1$ . In that case, the value of  $\Phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})$  is steady state distributed.*

*Coupling time:  $\tau^b \stackrel{\text{def}}{=} \min\{n \in \mathbb{N} : |\Phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})| = 1\}$ .*

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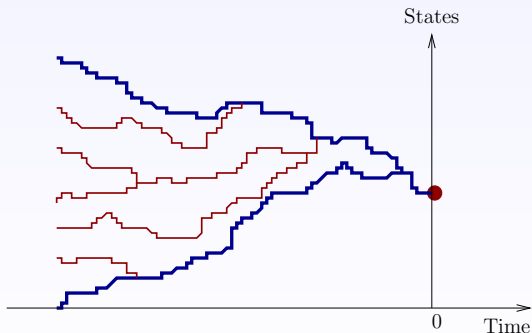
Coupling time:  $\tau^b \stackrel{\text{def}}{=} \min\{n \in \mathbb{N} : |\Phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})| = 1\}$ .

Inconvenient: Complexity  $O(\tau^b \mathcal{C}(\phi) N)$ .

# Monotone systems

Assumption: state space is partially ordered ( $\prec$ ) and transition function is monotone:

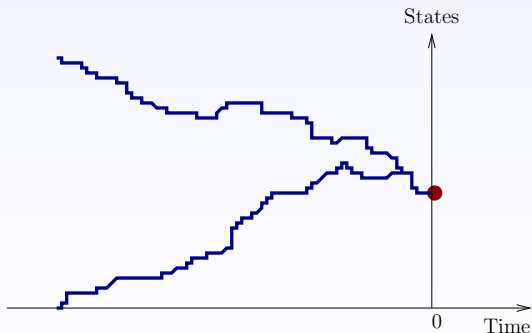
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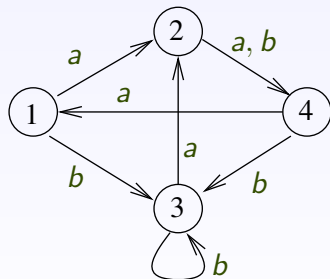
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# Non-monotone case

## Question

*What to do with non-monotone events?*



## Non-monotone case (II)

Assumption:  $(\mathcal{X}, \prec)$  is a complete lattice.

Let  $T \stackrel{\text{def}}{=} \sup \mathcal{X}$  and  $B \stackrel{\text{def}}{=} \inf \mathcal{X}$ .

New transition function  $\Gamma : \mathcal{X} \times \mathcal{X} \times \mathcal{E} \rightarrow \mathcal{X} \times \mathcal{X}$

$$\Gamma_1(m, M, e) \stackrel{\text{def}}{=} \inf_{m \prec x \prec M} \phi(x, e)$$

$$\Gamma_2(m, M, e) \stackrel{\text{def}}{=} \sup_{m \prec x \prec M} \phi(x, e).$$

### Theorem

If  $\Gamma^n(B, T, e_{-n+1 \rightarrow 0})$  hits the diagonal  $\mathcal{D}$  (i.e. states of the form  $(x, x)$ ) in finite time:  $\tau^e \stackrel{\text{def}}{=} \min \left\{ n : \Gamma^n(B, T, e_{-n+1 \rightarrow 0}) \in \mathcal{D} \right\}$ ,  
then  $\Gamma^{\tau^e}(B, T, e_{-\tau^e+1 \rightarrow 0})$  has the steady state distribution  $\pi$ .

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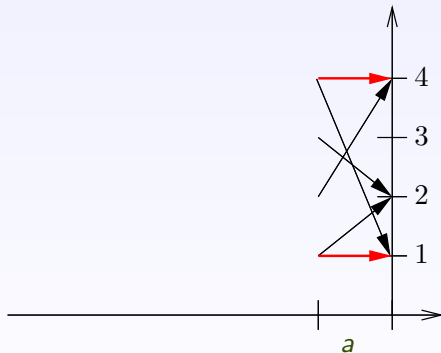
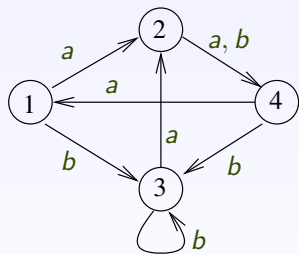
$$\Gamma_2(m, M, e) \stackrel{\text{def}}{=} \sup_{m \prec x \prec M} \phi(x, e).$$

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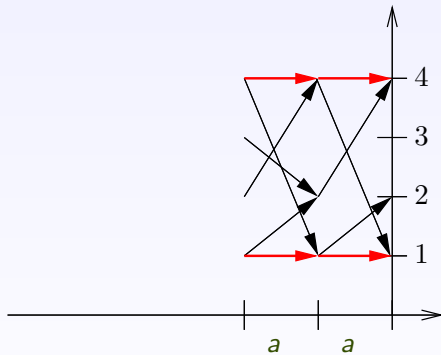
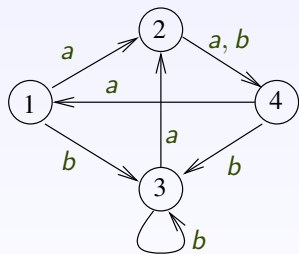
Proof: If  $(m_0, M_0) \stackrel{\text{def}}{=} \Gamma^n(B, T, e_{-n+1 \rightarrow 0})$ , then the set  $\phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})$  is included in  $\{x : m_0 \prec x \prec M_0\}$ . If the latter is reduced to one point, so is the set  $\phi^n(\mathcal{X}, e_{-n+1 \rightarrow 0})$ .

# Example

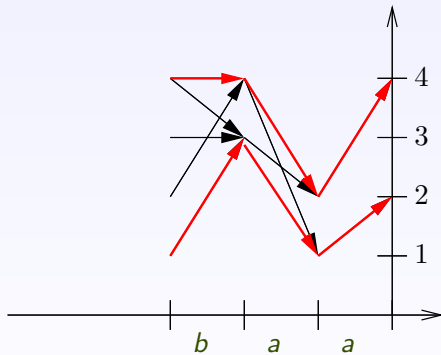
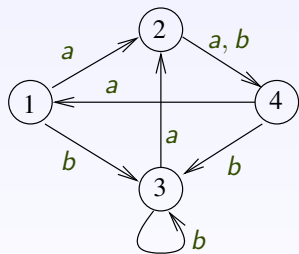




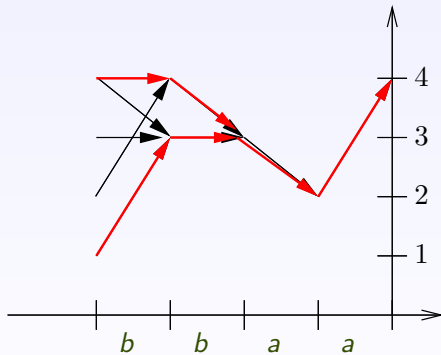
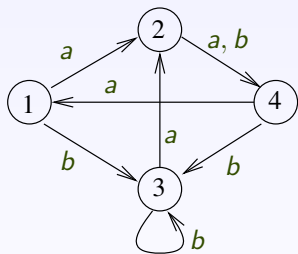
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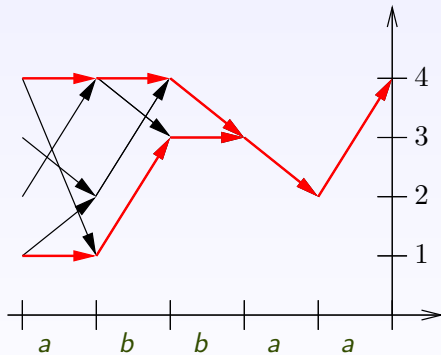
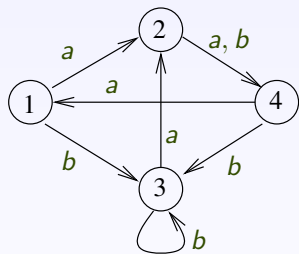
# Example



# Example



# Example



# Envelope perfect simulation

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**Data:** -  $\Phi$ ,  $\{e_{-n}\}_{n \in \mathbb{N}}$   
-  $\Gamma$  the pre-computed envelope function

**Result:** A state  $x^* \in \mathcal{X}$  generated according to the stationary distribution of the system

**begin**

```
 $n = 1; M := T; m := B;$   
repeat  
  for  $i = n - 1$  downto  $0$  do  
     $(m, M) := \Gamma(m, M, e_{-i});$   
   $n := 2n;$   
until  $M = m;$   
 $x^* := M;$   
return  $x^*;$ 
```

**end**

---

Complexity:  $O(\mathcal{C}(\Gamma)\tau^e)$  (to compare with  $O(\mathcal{C}(\phi)N\tau^b)$ ).

# Comments

1. Everything works the same if  $\Gamma_1$  (resp.  $\Gamma_2$ ) is replaced by a lower (resp. upper) bound on the infimum (res. supremum).
2. The definition of the envelopes is based on the constructive definition  $\Phi$  of the Markov chain. For a new event representation  $\Phi'$  of the Markov chain envelopes are modified accordingly.
3. If the function  $\Phi(., e)$  is non-decreasing for all event  $e$ , then for any  $m \leq M$ ,  $\Gamma_1(m, M, e) = \Phi(m, e)$  and  $\Gamma_2(m, M, e) = \Phi(M, e)$ , so that Algorithm EPSA coincides with the classical monotone perfect simulation algorithm for monotone Markov chains.

# Problems

- ▶ The envelopes may not couple even if the trajectories do.  
Example: a single queue with batch arrivals of size 3 and batch services of size 2. (Notation:  $(+3, -2)$  queue.)  
If the whole batch cannot be accepted, the batch is rejected (blocking).

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If the whole batch cannot be accepted, the batch is rejected (blocking).
- ▶ When the envelopes couple, the coupling time of envelopes can be much longer.  
Example: as above, with individual and batch arrivals.
- ▶ The complexity of envelope computation might be too high.  
Complexity of EPISA:  $O(\mathcal{C}(\Gamma) \cdot \tau^e)$ .  
 $\mathcal{C}(\Gamma)$  should not depend on  $N!$

# Queuing networks

Most of the events are piece-wise space homogeneous (i.e.  $\phi(x, e) = x + v_R$  for  $x$  in region  $R$ ) and we often have:  $\mathcal{C}(\Gamma) \sim \mathcal{C}(\phi)$ .

Difference between PSA and EPSA in  $N_{\mathcal{T}^b}$  and  $\mathcal{T}^e$ .

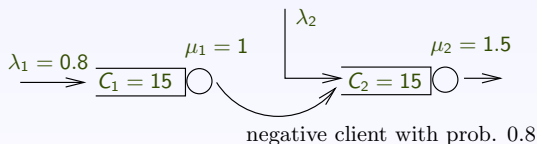


Figure: A network with negative customers.

## Queuing networks (II)

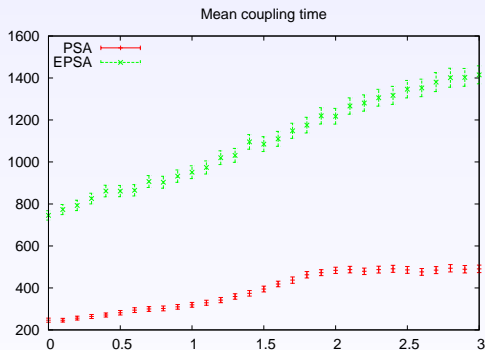
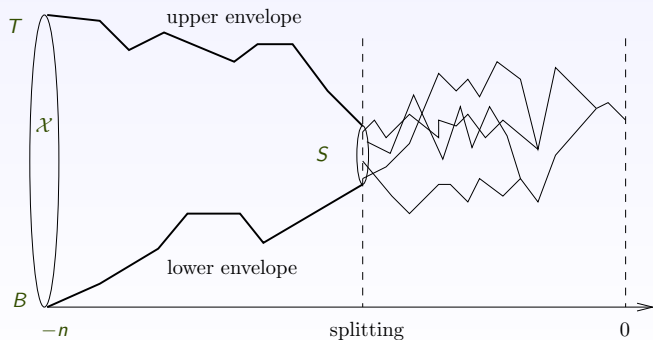


Figure: Mean coupling times of PSA and EPSA algorithms for the network in Figure 1 as a function of  $\lambda_2$ .

# Beyond envelopes

When the coupling time for envelopes is too long (or if they do not couple):

- ▶ bounds
- ▶ splitting



# Example

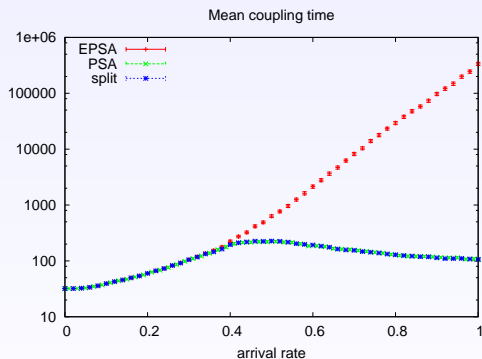
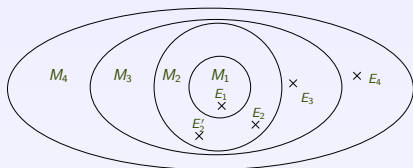


Figure: Mean coupling times for PSA, EPSA and EPSA with splitting for a  $(+2, +3, -1)$  queue.

# Classes



## Classes:

- ▶  $M_1$  - monotone MC
- ▶  $M_2$  - non-monotone MC, where envelope perfect simulation can be used efficiently
- ▶  $M_3$  - envelopes do couple but take a much larger time
- ▶  $M_4$  - envelopes do not couple (bounds, splitting)

## Examples:

- ▶  $E_1$  - a network of finite queues with monotone routing.
- ▶  $E_2$  - a network as  $E_1$  with negative customers
- ▶  $E_2'$  - a network as  $E_1$  with fork and join nodes
- ▶  $E_3$  - a network with individual customers and batches
- ▶  $E_4$  - a network of queues with only batches larger than two.