

Importance sampling simulation of the fork-join queue

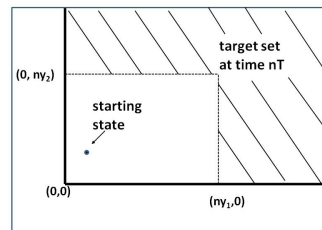
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The rare event

$\{S(k) = (S_1(k), S_2(k)) : k = 0, 1, \dots\}$
 is the discrete-time Markov chain
 analogon of the fork-join queue by
 embedding at jump times;
 $S(k)$ represents the backlogs at the
 queues.



Problem

Estimate by simulation:

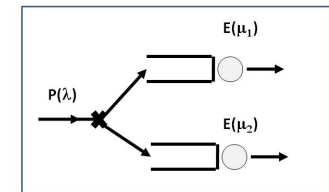
$$\gamma_n(x, y, T) = \mathbb{P}(S_1(nT) \geq ny_1 \text{ or } S_2(nT) \geq ny_2 | S(0) = nx),$$

for fixed scaled initial state $x = (x_1, x_2) \in \mathbb{R}_+^2$, fixed scaled threshold $y = (y_1, y_2) \in \mathbb{R}_+^2$, fixed scaled horizon $T > 0$, and parameter $n \rightarrow \infty$.

The fork-join queue

Model

- Poisson (λ) arrivals;
- an arriving job splits in two subjobs;
- two independent single server queues;
- exponential service times with rate μ_1 and μ_2 , resp;
- for stability $\lambda < \min(\mu_1, \mu_2)$.



Folklore application: two bathrooms, one for men and one for women, and arrivals of couples.

Original motivation: machine with parallel processors (Hahn&Flatto 1984).

More general: allow individual arrivals (Wright 1992; Schwartz-Weiss 1992).

The rarity set

The scaled set of interest:

$$D = \{\eta \in \mathbb{R}_+^2 : \eta_1 \geq y_1 \text{ or } \eta_2 \geq y_2\},$$

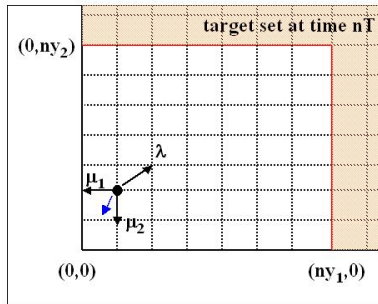
i.e.

$$\gamma_n(x, y, T) = \mathbb{P}(S(nT)/n \in D | S(0) = nx).$$

The difficulty for importance sampling is twofold:

- the rarity set is not convex (Dupuis&Wang 2007);
- the rarity set cannot be decomposed into two disjoint sets such that the separate probabilities are estimated by efficient importance sampling estimators (Glassermann&Wang 1997).

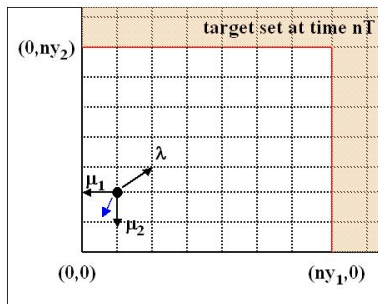
A sample path of the fork-join queue



The blue arrow indicates the 'natural' drift. We show the transition rates; for the discrete-time Markov chain these are normalized to probabilities.



A sample path after change of measure

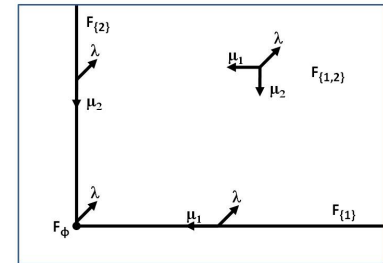


The idea of the importance sampling scheme: until a certain time nT it follows the original transition probabilities.



Face-homogeneous random walk

The fork-join queue is a face-homogeneous random walk on \mathbb{Z}_+^2 with four **faces**.



The transition probabilities $p_{s,s+d}$ are constant for s in the same face F_Λ . We might associate a random walk $(S_\Lambda(k))_{k=0}^\infty$ with jump variable X_Λ with probabilities $p_\Lambda(j) \doteq p_{s,s+j}$.

Importance sampling scheme

Our importance sampling scheme will be a mixture of two sets of **exponentially shifted** jump probabilities of the jump variables X_Λ .

For any $\theta \in \mathbb{R}^2$, the θ -shifted jump X_Λ^θ has jump probabilities

$$p_\Lambda^\theta(j) = e^{(\theta,j) - \psi_\Lambda(\theta)} p_\Lambda(j),$$

where $\psi_\Lambda(\cdot)$ is the log moment generating function of jump variable X_Λ .

This gives us a set of 4 jump (or transition) probability densities.

We have two of such sets, and before we simulate a sample path, we choose randomly a set.

Sample path large deviations

Define continuous processes $(S^{[n]}(t))_{0 \leq t \leq T}$, $n = 1, 2, \dots$, by **scaling**:

$$S^{[n]}(t) = S(nt)/n \text{ for } t = 0, 1/n, 2/n \dots, T,$$

and linear interpolation in the other points.

Consider absolute continuous functions $\phi : [0, T] \rightarrow \mathbb{R}_+^2$. Then (Ignatiouk 2005)

$$-\lim_{\epsilon \downarrow 0} \lim_{n \rightarrow \infty} \frac{1}{n} \log \mathbb{P} \left(\sup_{0 \leq t \leq T} |S^{[n]}(t) - \phi(t)| < \epsilon \right) = \int_0^T \ell_{\Lambda(\phi(t))}(\phi'(t)) dt \doteq I(\phi),$$

where $\ell_{\Lambda} : \mathbb{R}^2 \rightarrow [0, \infty]$ are so-called **locate rate functions** (see forthcoming slides).

The local rate functions

The local rate function $\ell_{\Lambda}(v)$ is the convex conjugate of a certain convex function $\psi : \mathbb{R}^2 \rightarrow \mathbb{R}$:

$$\ell_{\Lambda}(v) = \sup_{\theta \in \mathbb{R}^2} (\langle \theta, v \rangle - \psi(\theta)).$$

It can be determined numerically via the method in (Ignatiouk 2001).

The optimizer θ_v is called the optimal **shift factor** associated with speed v .

When θ_v is used to exponentially shift the jump probabilities of the **internal** jump variable $X_{\{1,2\}}$, the speed vector v corresponds with the drift of the shifted jump variable $X_{\{1,2\}}^{\theta_v}$, thus restricted for being a convex combination of the jumps $(-1, 0)$, $(0, -1)$, $(1, 1)$.

Clearly, the speed vectors in the **boundary** faces $F_{\{1\}}$ and $F_{\{2\}}$ are restricted to be 0 in the perpendicular direction and between -1 and 1 in the parallel direction. (They do not correspond with drifts!)

Denote by V_{Λ} the set of feasible speed vectors v in face F_{Λ} .

Sample path large deviations (cont'd)

Notice that for the fork-join queue problem

$$\gamma_n(x, y, T) = \mathbb{P}(S(nT)/n \in D | S(0)/n = x) = \mathbb{P}(S^{[n]} \in E),$$

where E is an appropriate set of absolute continuous paths $\phi : [0, T] \rightarrow \mathbb{R}_+^2$ with specifically $\phi(0) = x$ and $\phi(T) \in D$ (the rarity set).

Let $\tilde{E} \subset E$ be the subset of **piecewise linear paths** of the following form.

$\phi = \phi_{\tau, v}$: it follows the **natural drift** until time τ and then it goes **straight** at constant **speed** $v = \phi'(t)$ to a point in the rarity set D .

One can show that

$$\lim_{n \rightarrow \infty} \frac{1}{n} \log \mathbb{P}(S^{[n]} \in E) = \lim_{n \rightarrow \infty} \frac{1}{n} \log \mathbb{P}(S^{[n]} \in \tilde{E}) = -I(\tilde{E}),$$

where $I(\tilde{E}) = \inf_{\tau, v} I(\phi_{\tau, v})$, and $I(\phi_{\tau, v}) = (T - \tau)\ell_{\Lambda}(v)$ assuming that the second part of the path runs entirely in face F_{Λ} .

The paths with constant speed

We restrict to starting state $x = (0, 0)$.

Consider paths that stay in 0 during τ time units.

Let $V_{\Lambda}(\tau)$ be the set of feasible speed vectors v in face F_{Λ} such that $(T - \tau)v \in D$, i.e.,

$$V_{\Lambda}(\tau) = \{v \in V_{\Lambda} : \phi_{\tau, v} \in \tilde{E}\}.$$

Hence, there is a one-to-one correspondence

$$\begin{aligned} \tilde{E} &\leftrightarrow \bigcup_{\tau \geq 0} V_{\{1,2\}}(\tau) \cup \bigcup_{\tau \geq 0} V_{\{1\}}(\tau) \cup \bigcup_{\tau \geq 0} V_{\{2\}}(\tau) \\ &= V_{\{1,2\}}(0) \cup V_{\{1\}}(0) \cup V_{\{2\}}(0) \doteq V(0). \end{aligned}$$

Efficient importance sampling

The idea is to choose (better: find) a set of speeds $v^{(1)}, \dots, v^{(m)}$ and associated optimal shift factors $\theta^{(1)}, \dots, \theta^{(m)}$, such that

$$V(0) \subset \bigcup_{i=1}^m \mathcal{H}(v^{(i)}),$$

where

$$\mathcal{H}(v) = \{w \in V(0) : \langle \theta_v, w \rangle \geq \langle \theta_v, v \rangle\}.$$

Then, any mixture importance sampling scheme with exponentially shifted probability densities using shift factors $\theta^{(1)}, \dots, \theta^{(m)}$ is asymptotically optimal (Bucklew 1990, 2004).

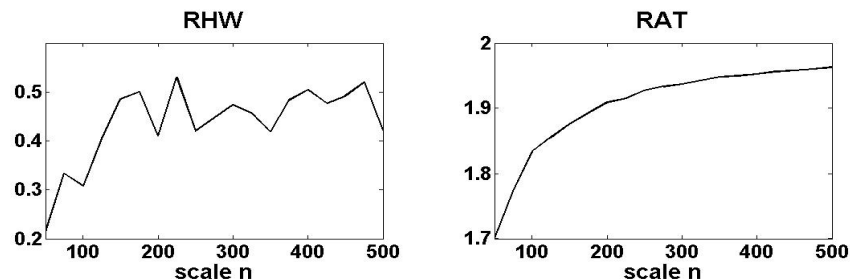
Example (cont'd)

Results for scalings $n = 25\text{--}500$ with sample size $k = 50000$ for plotting the relative half width of the 95% confidence interval for estimator $\hat{\gamma}_n$,

$$\text{RHW} = 1.96 \sqrt{\text{Var}[\hat{\gamma}_n]} / \mathbb{E}[\hat{\gamma}_n],$$

$$\text{and ratio RAT} = \log \mathbb{E}[(\hat{\gamma}_n)^2] / \log \mathbb{E}[\hat{\gamma}_n].$$

(efficient estimators have RAT that converge to 2).



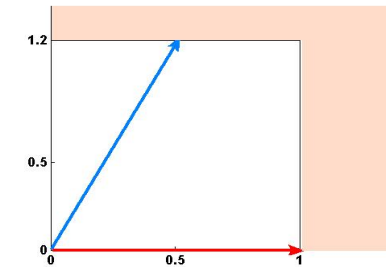
Example

$$\lambda = 1, \mu_1 = 1.5, \mu_2 = 2, x = (0, 0), y = (1, 1.2), T = 10.$$

We were able to find a solution of two speed vectors such that

$$V(2) \doteq \bigcup_{\tau \geq 2} V_{\{1,2\}}(\tau) \cup \bigcup_{\tau \geq 2} V_{\{1\}}(\tau) \cup \bigcup_{\tau \geq 2} V_{\{2\}}(\tau) \subset \mathcal{H}(v^{(1)}) \cup \mathcal{H}(v^{(2)}).$$

We mix them 0.8 (red path) and 0.2 (blue path).



Conclusion and further research

- We have developed an importance sampling scheme which is a mixture of two time-dependent (state-independent) exponentially shifted densities.
- Excellent simulation results.
- Need to prove that using $V(2)$ in stead of $V(0)$ still gives asymptotical optimality. (The large deviations asymptotic still holds.)
- Further investigations include other starting points, and other algorithms, for instance with mixing transition probabilities that depend on state and time.

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