



# Chasing Errors by Diagnostics

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$$P_n(t) = \text{Prob}(\mathcal{X}_A(t) = n)$$

$$\frac{dP_n(t)}{dt} = \sum_{m=0}^{\infty} \left( W(n | m, t) P_m(t) - W(m | n, t) P_n(t) \right); \quad n, m \in \mathbb{N}$$

$$\frac{dP_n(t)}{dt} = w_{n-1}^+ P_{n-1}(t) + w_{n+1}^- P_{n+1}(t) - (w_n^+ + w_n^-) P_n(t)$$

$$w_n^+ \equiv \lambda_n \quad \text{for } n \rightarrow n + 1 \quad \text{and}$$

$$w_n^- \equiv \mu_n \quad \text{for } n \rightarrow n - 1$$

Master equation in chemical reaction kinetics

$$P(\tau, \boldsymbol{\mu}) = P_1(\tau) \cdot P_2(\boldsymbol{\mu}|\tau)$$

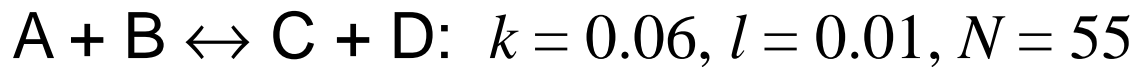
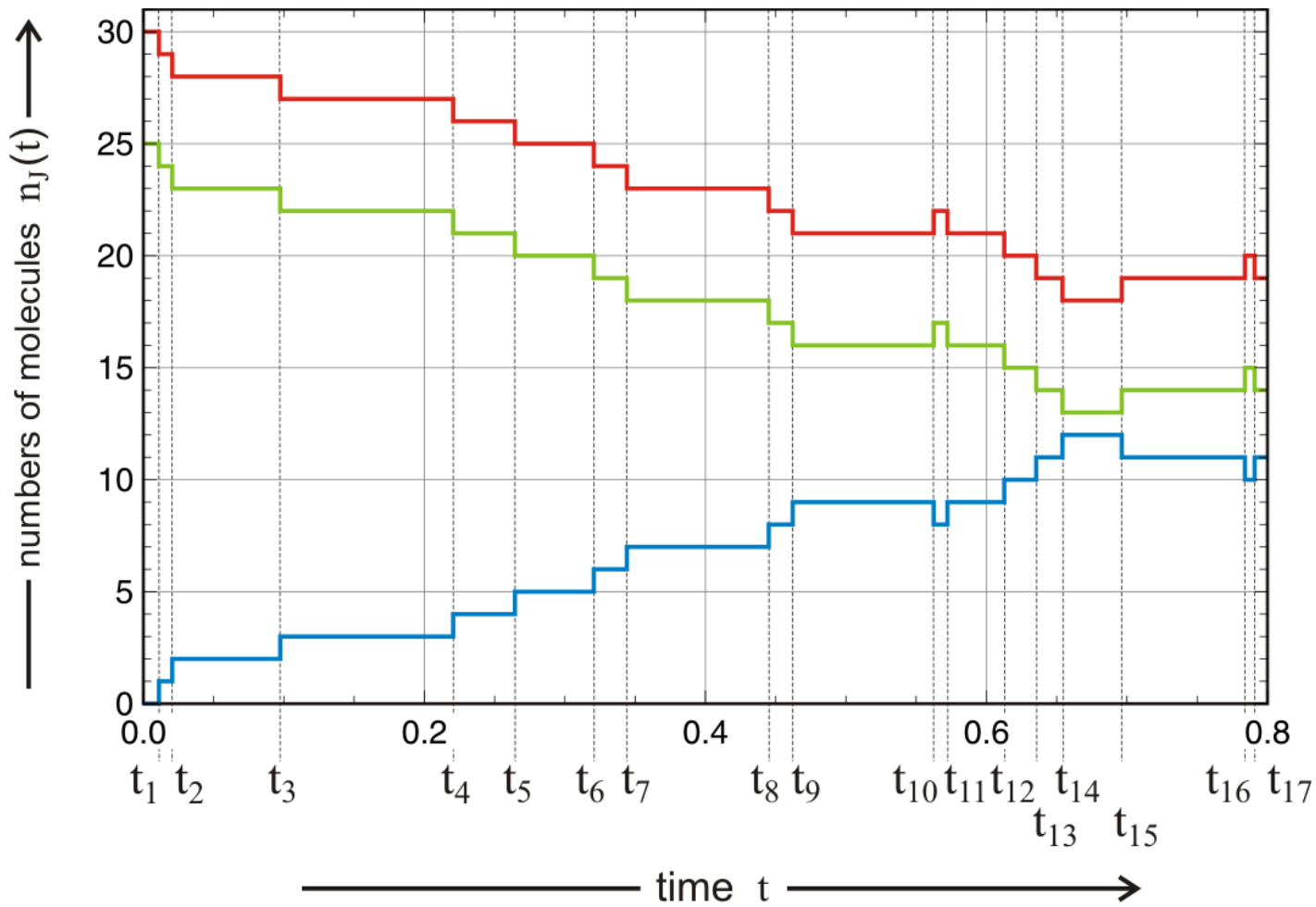
$$P_1(\tau) = \sum_{\boldsymbol{\mu}=1}^K P(\tau, \boldsymbol{\mu}) \quad P_2(\boldsymbol{\mu}|\tau) = P(\tau, \boldsymbol{\mu}) / \sum_{\boldsymbol{v}}^K P(\tau, \boldsymbol{v})$$

$$P_1(\tau) = \alpha \exp(-\alpha \tau), \quad 0 \leq \tau < \infty \quad \text{and}$$

$$P_2(\boldsymbol{\mu}|\tau) = P_2(\boldsymbol{\mu}) = \alpha_{\boldsymbol{\mu}} / \alpha, \quad \boldsymbol{\mu} = 1, \dots, K$$

$$\alpha_{\boldsymbol{\mu}} \equiv \gamma_{\boldsymbol{\mu}} h_{\boldsymbol{\mu}}(\mathbf{n}) \quad \text{and} \quad \alpha = \sum_{\boldsymbol{\mu}=1}^K \alpha_{\boldsymbol{\mu}} \equiv \sum_{\boldsymbol{\mu}=1}^K \gamma_{\boldsymbol{\mu}} h_{\boldsymbol{\mu}}(\mathbf{n})$$

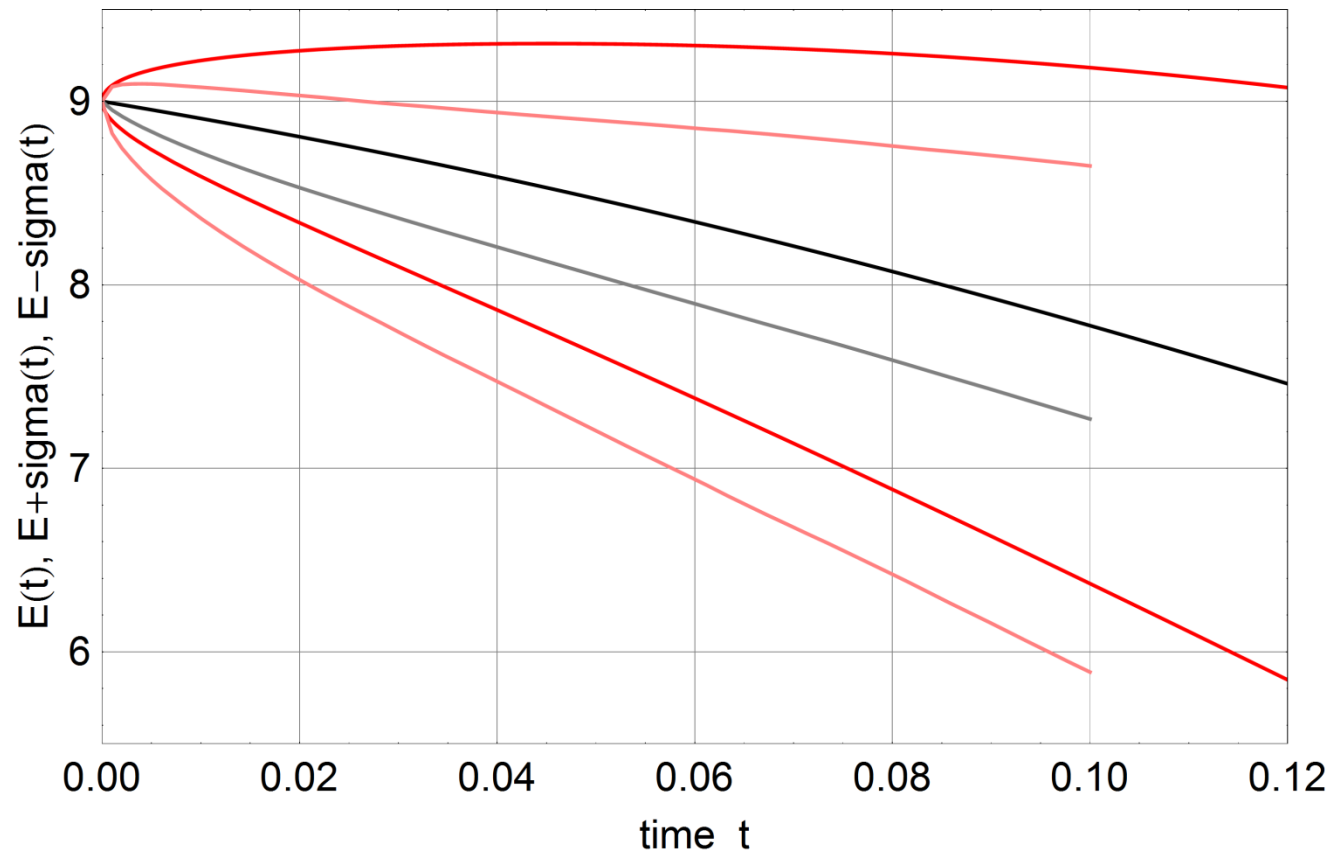
The stochastic simulation algorithm (SSA, Gillespie et al.)



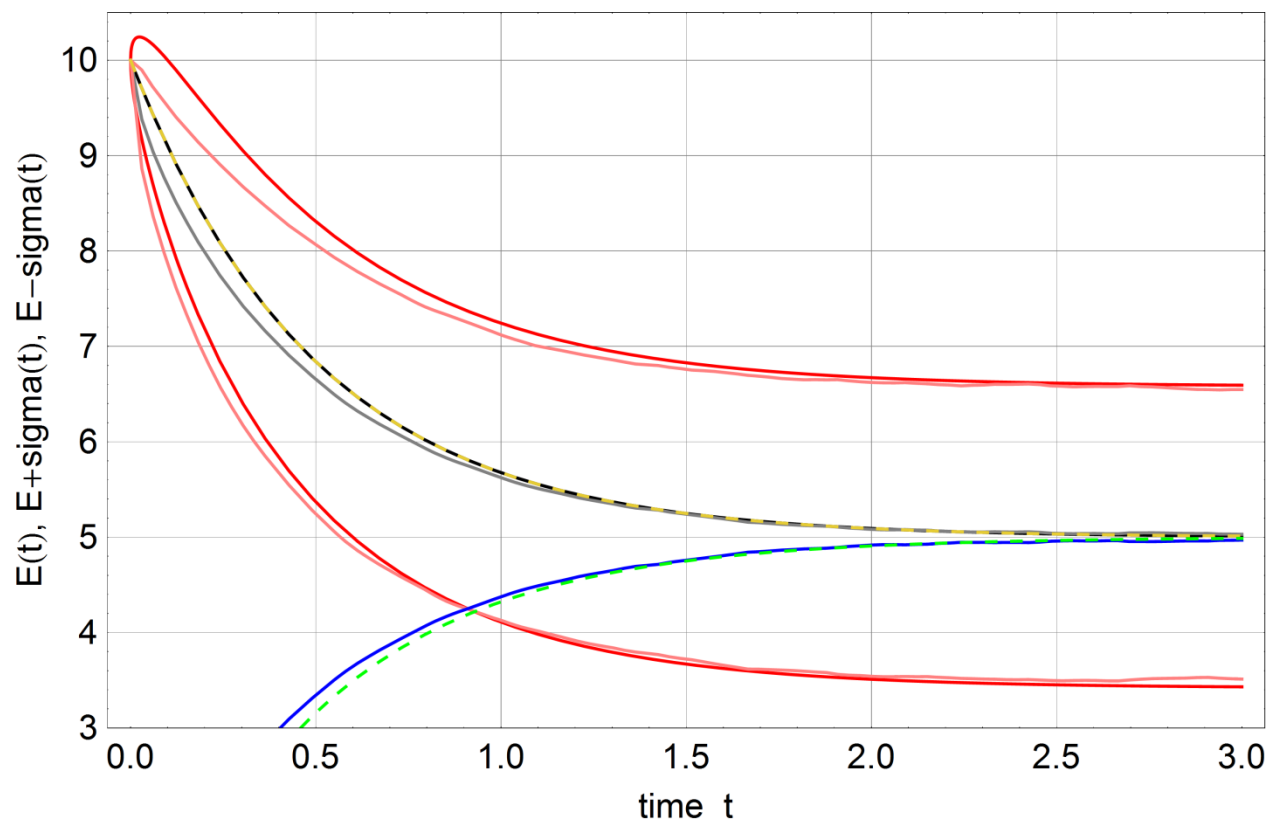
## Simulation tool: xSSA-lite

xSSA is a module of the xCellerator simulation package for systems biology of whole cells.

Shapiro, B.E., Levchenko, A., World, E.M.M.B.J., Mjolsness, E.D.: Cellerator: Extending a computer algebra system to include biochemical arrows for signal transduction simulations. *Bioinformatics* **19**, 677–678 (2003)







$A \leftrightarrow B: k = 5, l = 5, N = 10\ 000$

Table 1: **Simulation errors of xSSA-lite in the xCellerator package for the reactions  $A \rightleftharpoons B$** . Shown are the equilibrium expectation values of the random variable  $\bar{E}(\mathcal{X}_A(t))$  and the width of the one- $\sigma$  confidence interval  $2\sigma(\mathcal{X}_A(t))$  obtained by simulation and calculation. Sample size:  $N = 1000$  trajectories.

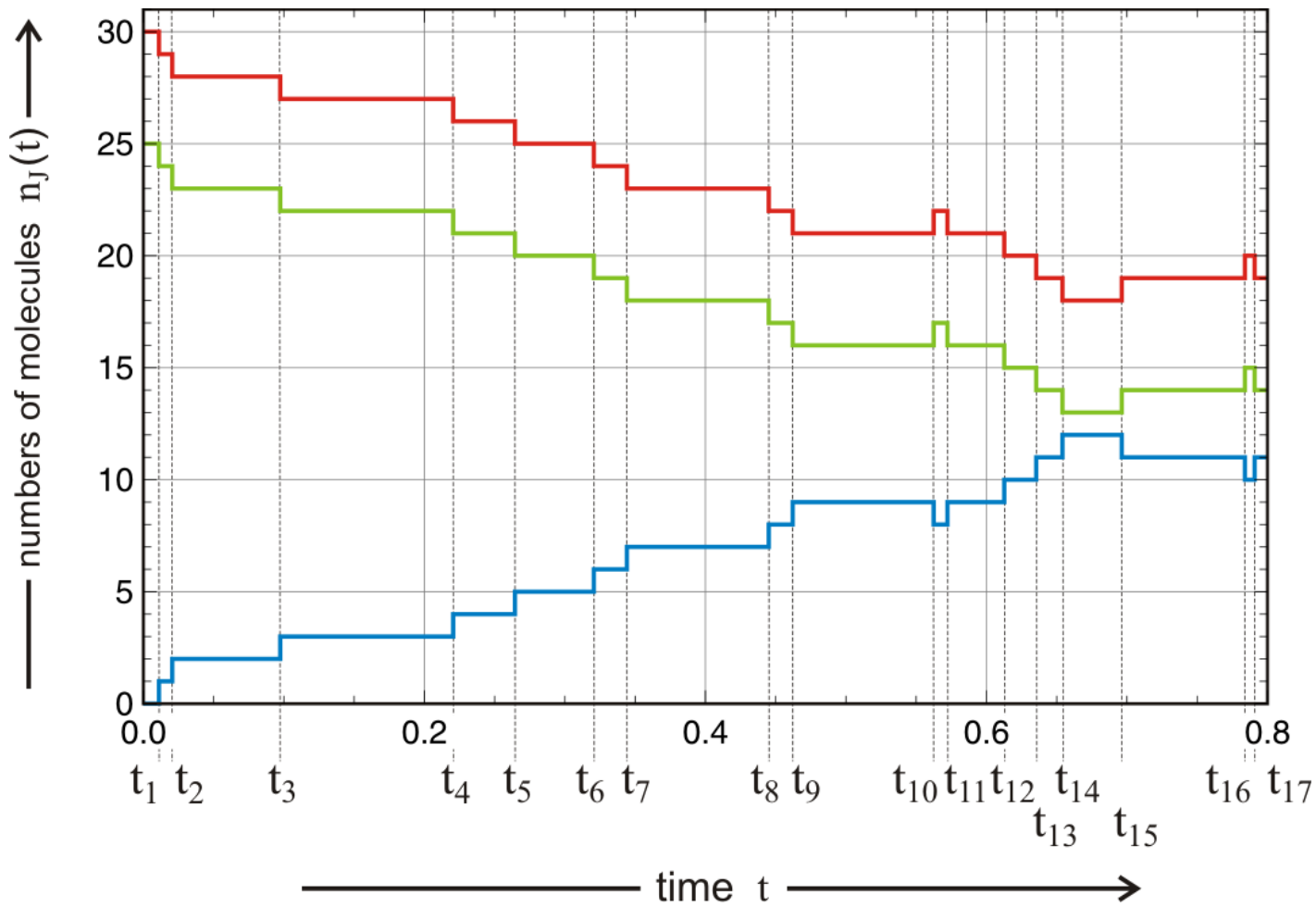
parameters			simulation			calculation		
$n_0$	$k$	$l$	$\bar{E}(\mathcal{X}_A)$	$\bar{E}(\mathcal{X}_B)$	$2\bar{\sigma}(\mathcal{X})$	$\bar{E}(\mathcal{X}_A)$	$\bar{E}(\mathcal{X}_B)$	$2\bar{\sigma}(\mathcal{X})$
10	5	5	5.005	4.995	3.068	5	5	3.162
10	4	6	5.995	4.005	3.011	6	4	3.098
10	3	7	6.977	3.023	2.778	7	3	2.898
10	2	8	7.949	2.051	2.382	8	2	2.529
10	1	9	8.883	1.117	1.676	9	1	1.897

Binomial distribution:  $\bar{E}(\mathcal{X}_A) = n_0 \frac{l}{l+k}$  and  $\overline{\text{var}}(\mathcal{X}_A) = n_0 \frac{l \cdot k}{(l+k)^2}$

Expectation values are wrong for **asymmetric reaction rates**  
equilibrium constants  $K \neq 1$ .

Variances or standard deviations are calculated **too small**.

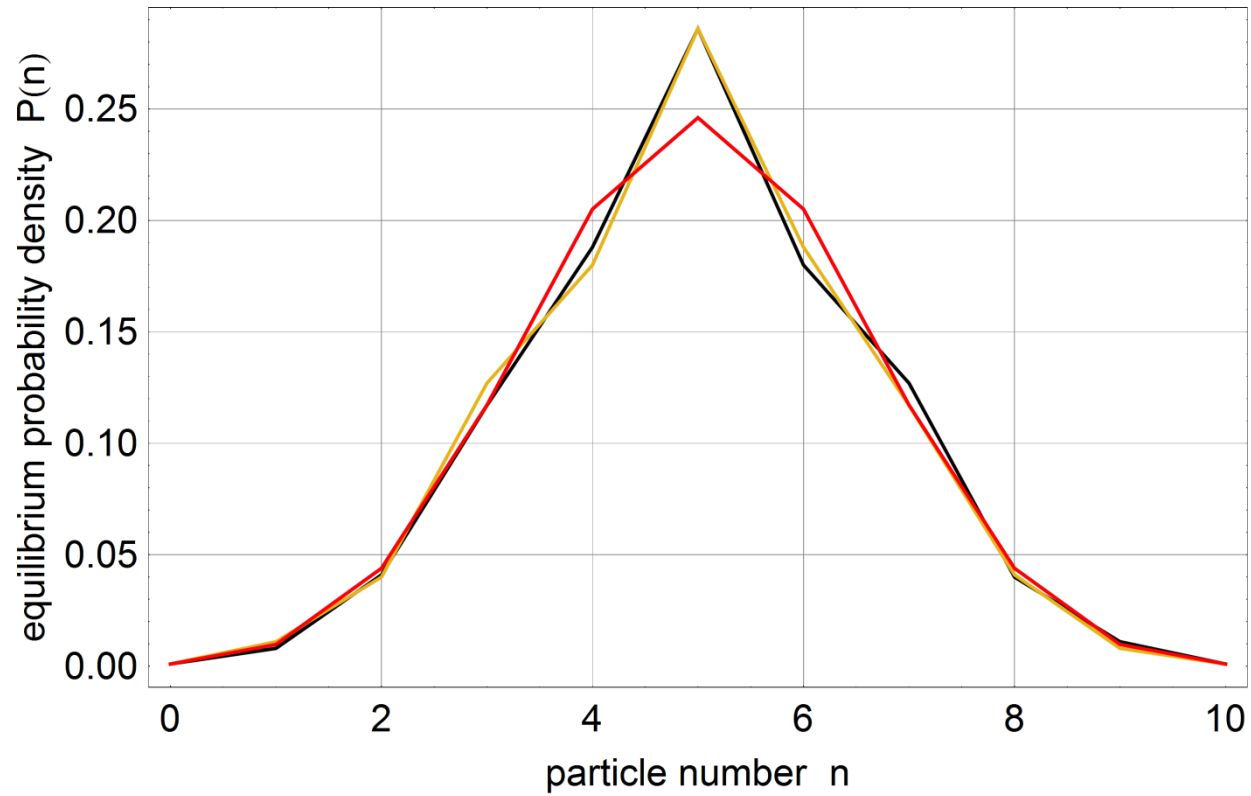
Diagnostics of simulation errors



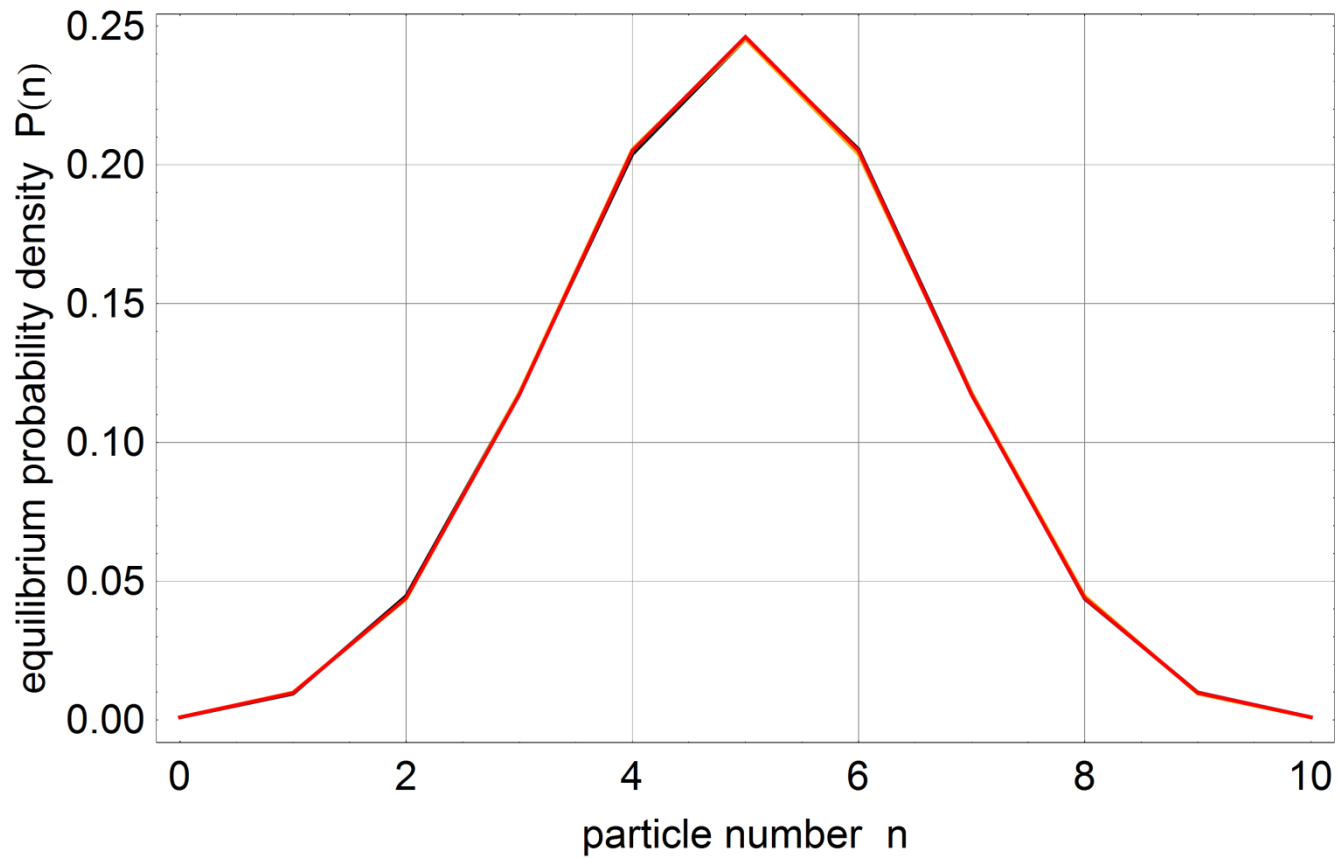
Endpoint statistics

Table 2: **Endpoint statistics of xSSA-lite in the xCellerator package for the reactions  $A \rightleftharpoons B$** . Shown are the equilibrium expectation values of the random variable  $\bar{E}(\mathcal{X}_A(t))$  and the width of the one- $\sigma$  confidence interval  $2\bar{\sigma}(\mathcal{X}_A(t))$  obtained by simulation and calculation. Sample size:  $N = 1000$  trajectories.

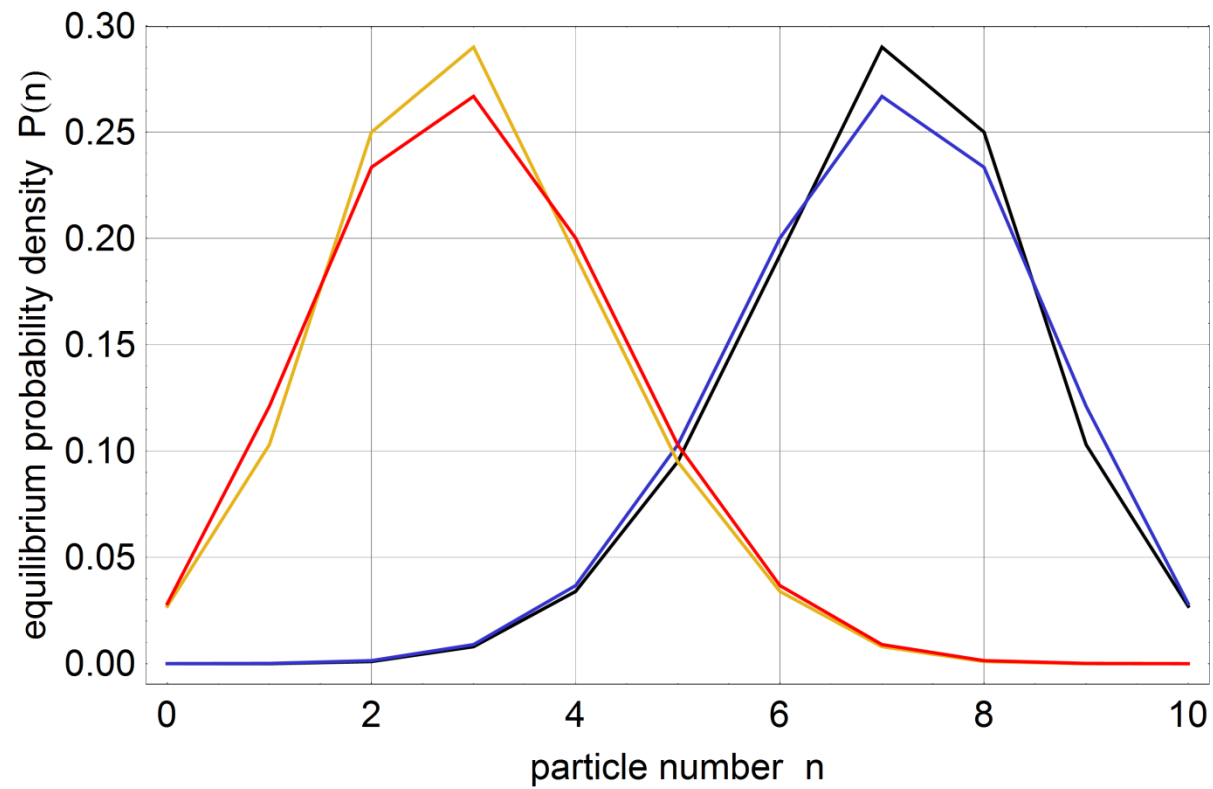
parameters			simulation			calculation		
$n_0$	$k$	$l$	$\bar{E}(\mathcal{X}_A)$	$\bar{E}(\mathcal{X}_B)$	$2\bar{\sigma}(\mathcal{X})$	$\bar{E}(\mathcal{X}_A)$	$\bar{E}(\mathcal{X}_B)$	$2\bar{\sigma}(\mathcal{X})$
10	5	5	5.001	4.999	3.166	5	5	3.162
10	3	7	6.998	3.002	2.899	7	3	2.898
10	1	9	8.996	1.004	1.903	9	1	1.897



$A \leftrightarrow B: k = 5, l = 5, N = 1000$

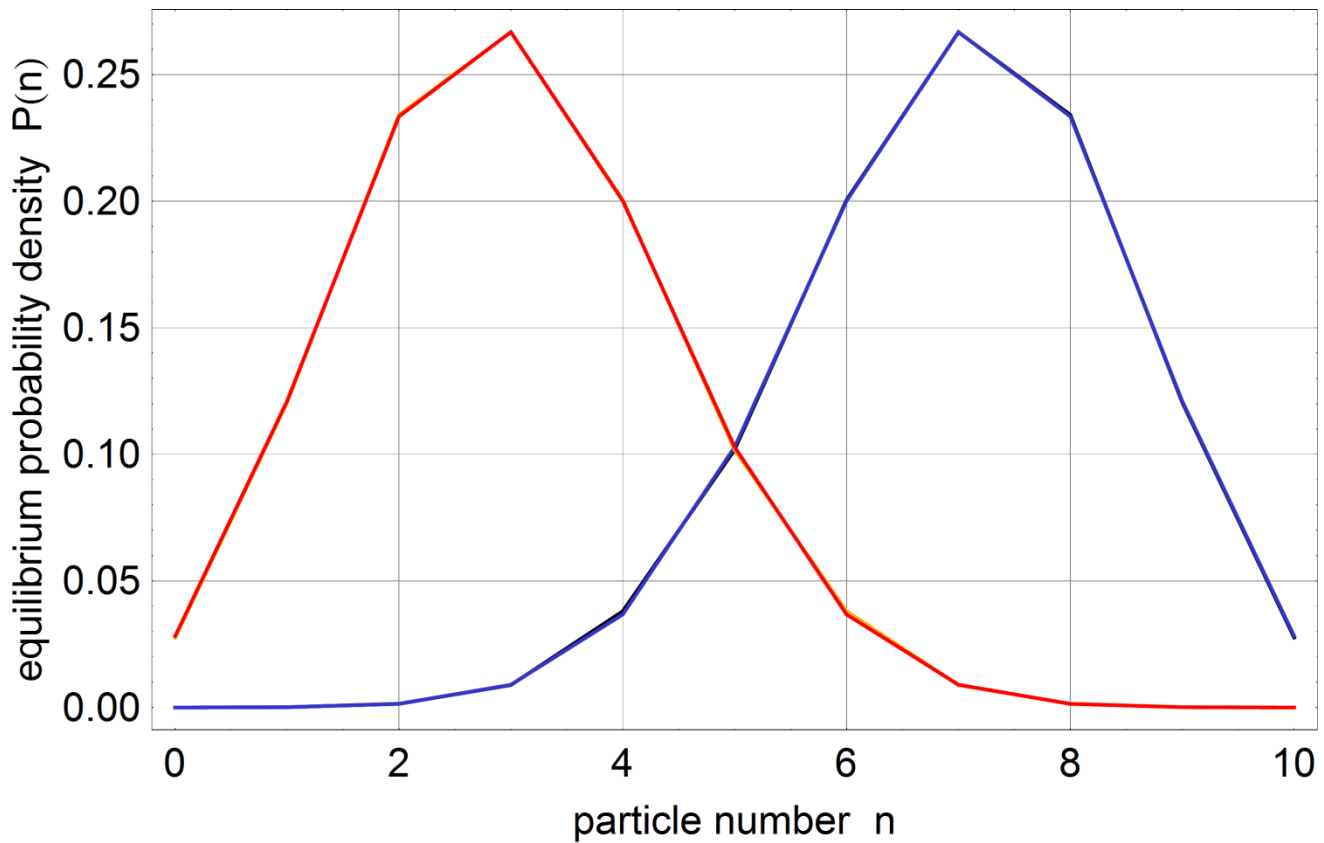


$A \leftrightarrow B: k = 5, l = 5, N = 100\,000$

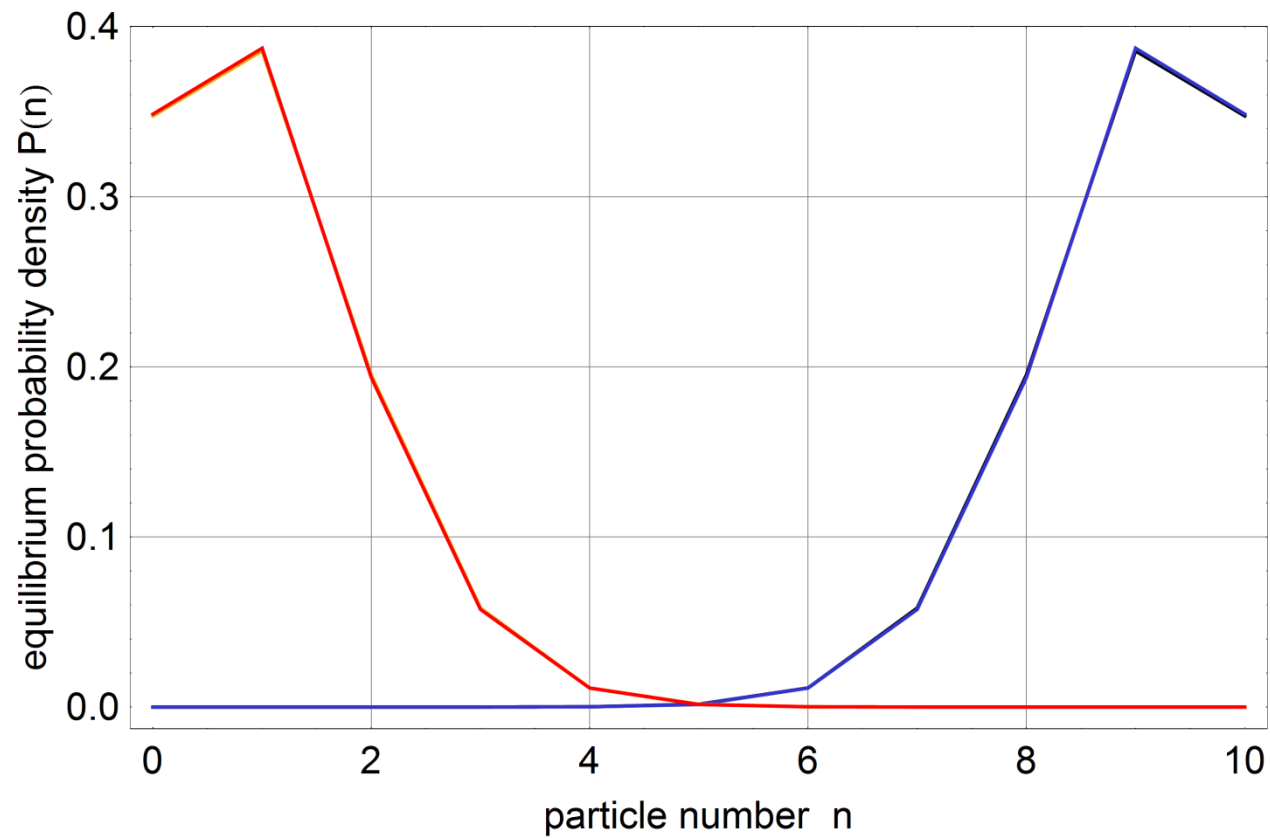


$A \leftrightarrow B: k = 3, l = 7, N = 1000$

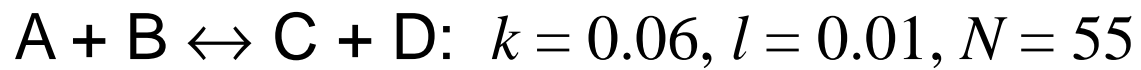
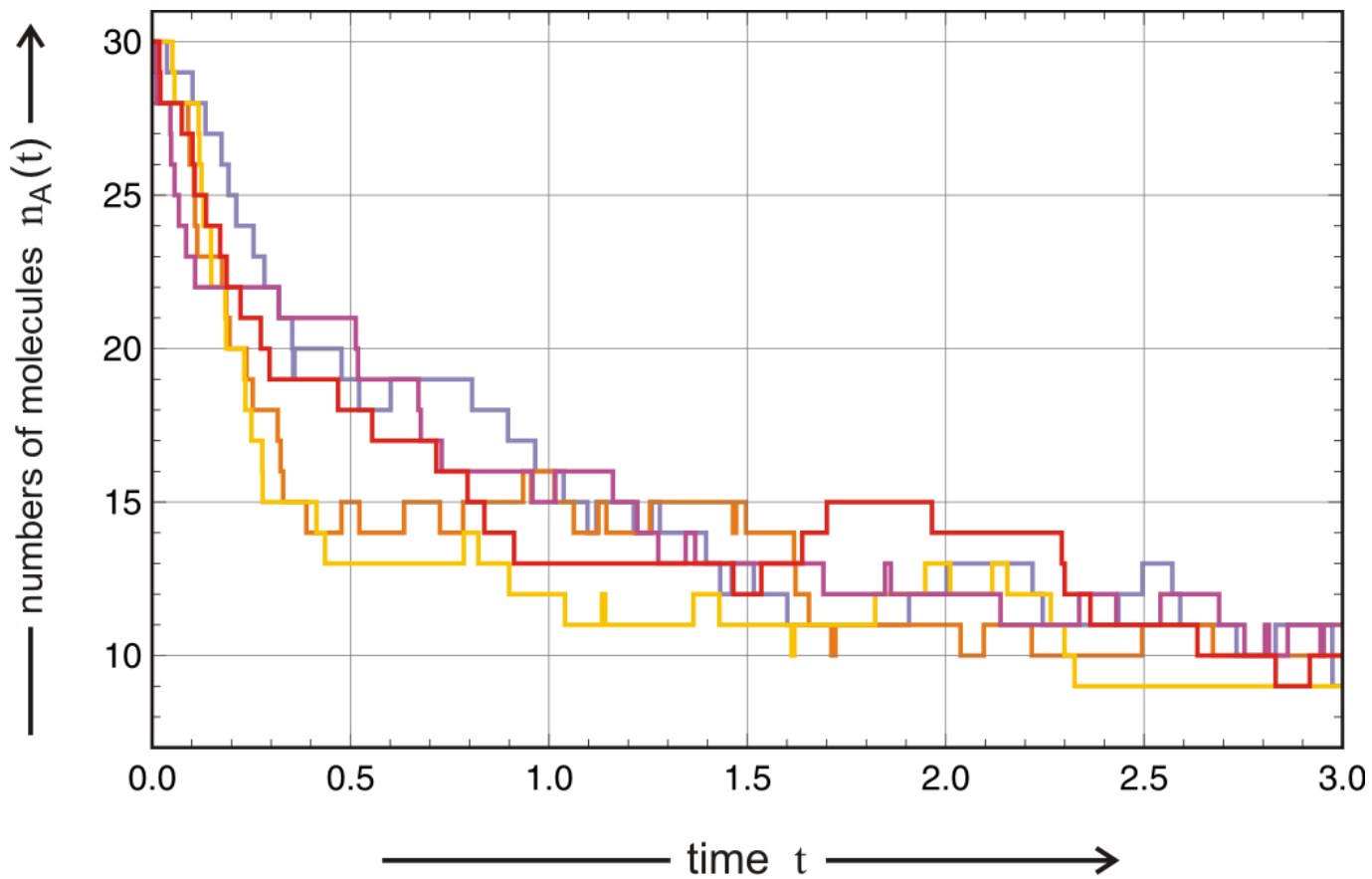


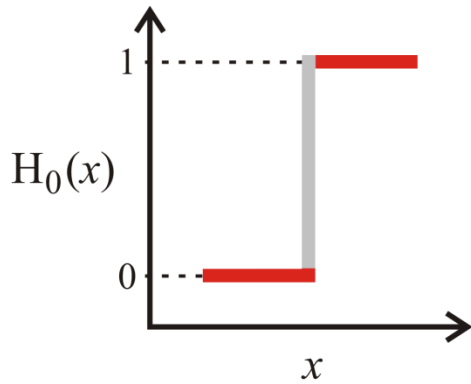


$A \leftrightarrow B: k = 3, l = 7, N = 100\,000$

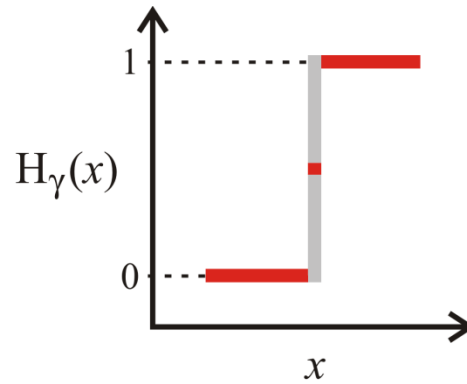


$A \leftrightarrow B: k = 1, l = 9, N = 100\,000$



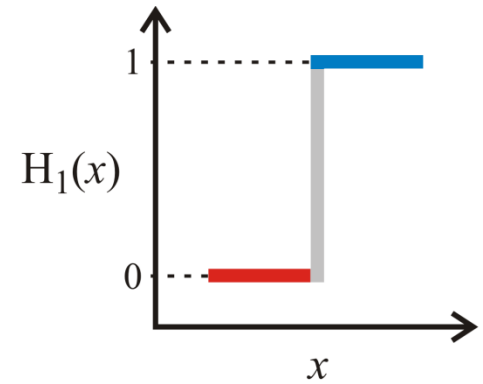


left-hand continuity



no continuity

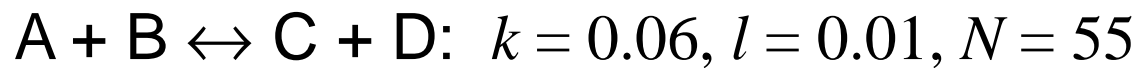
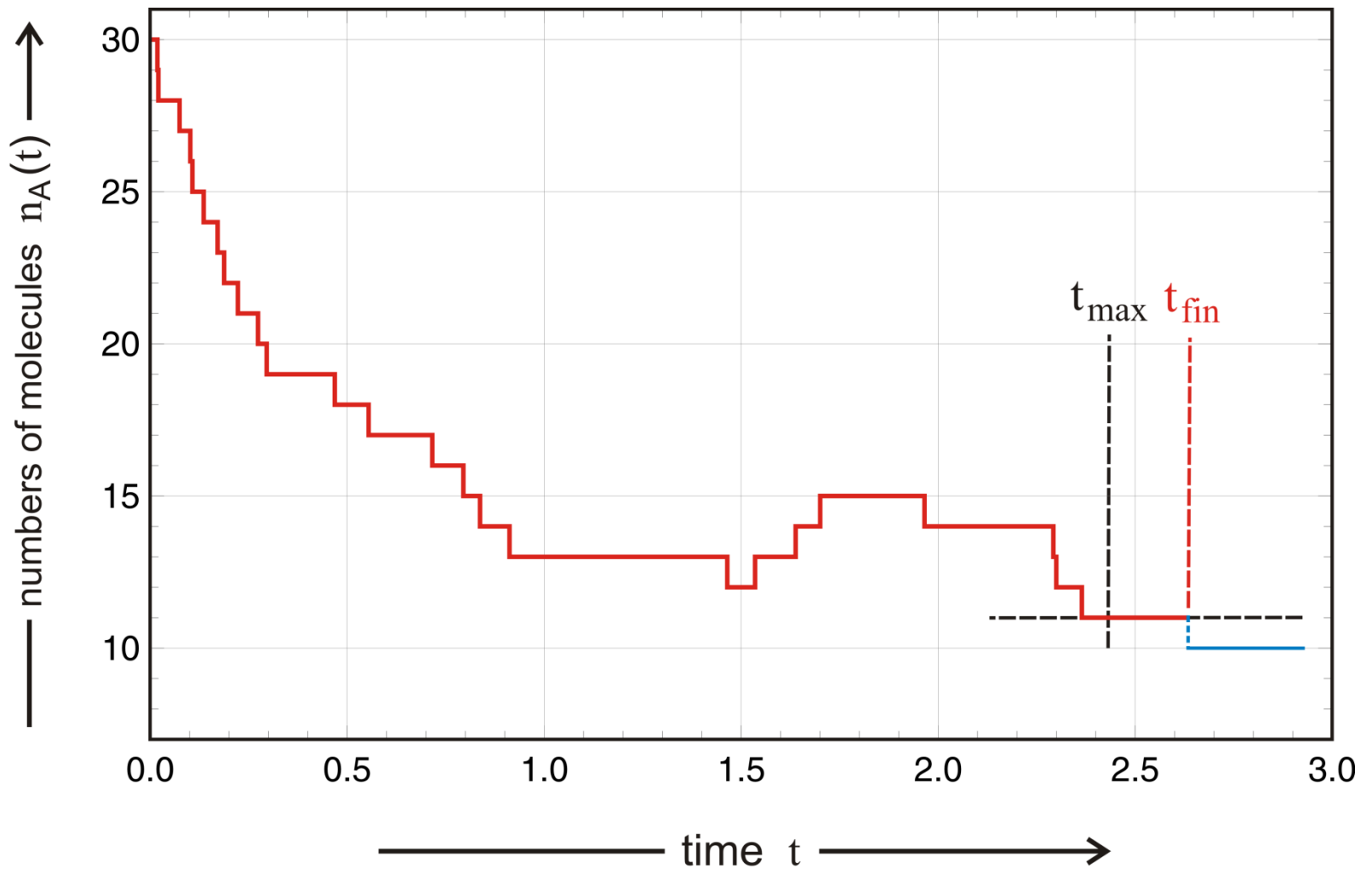
sign function

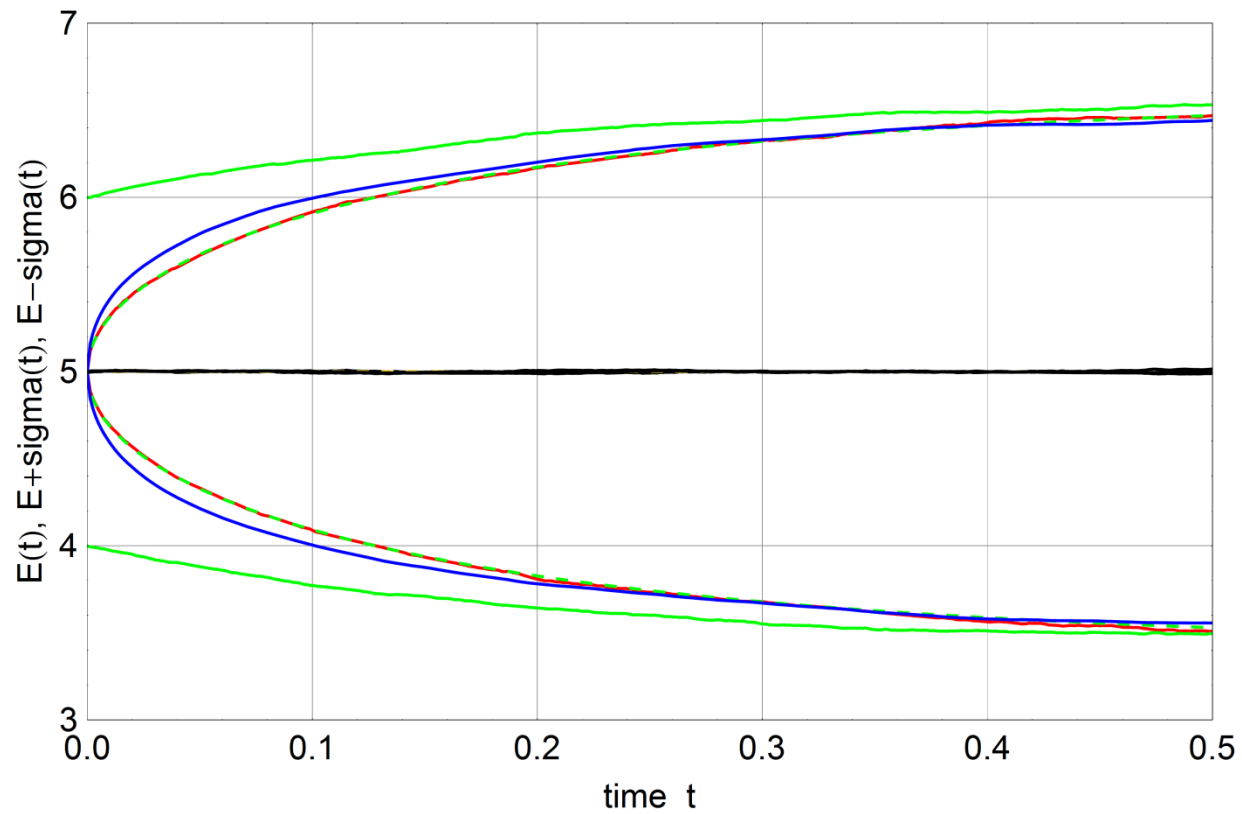


right-hand continuity

càdlàg convention

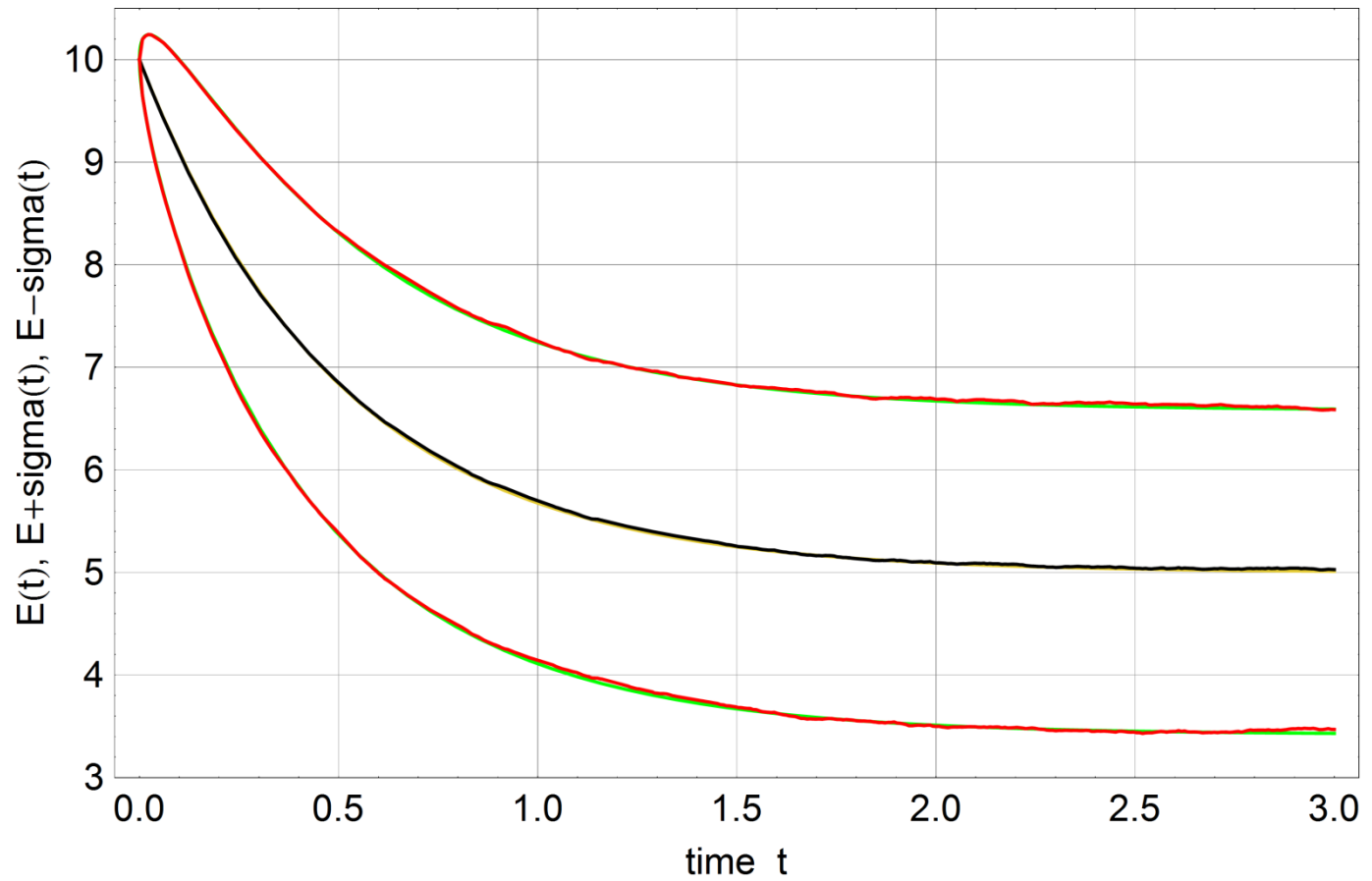
## Continuity in step functions



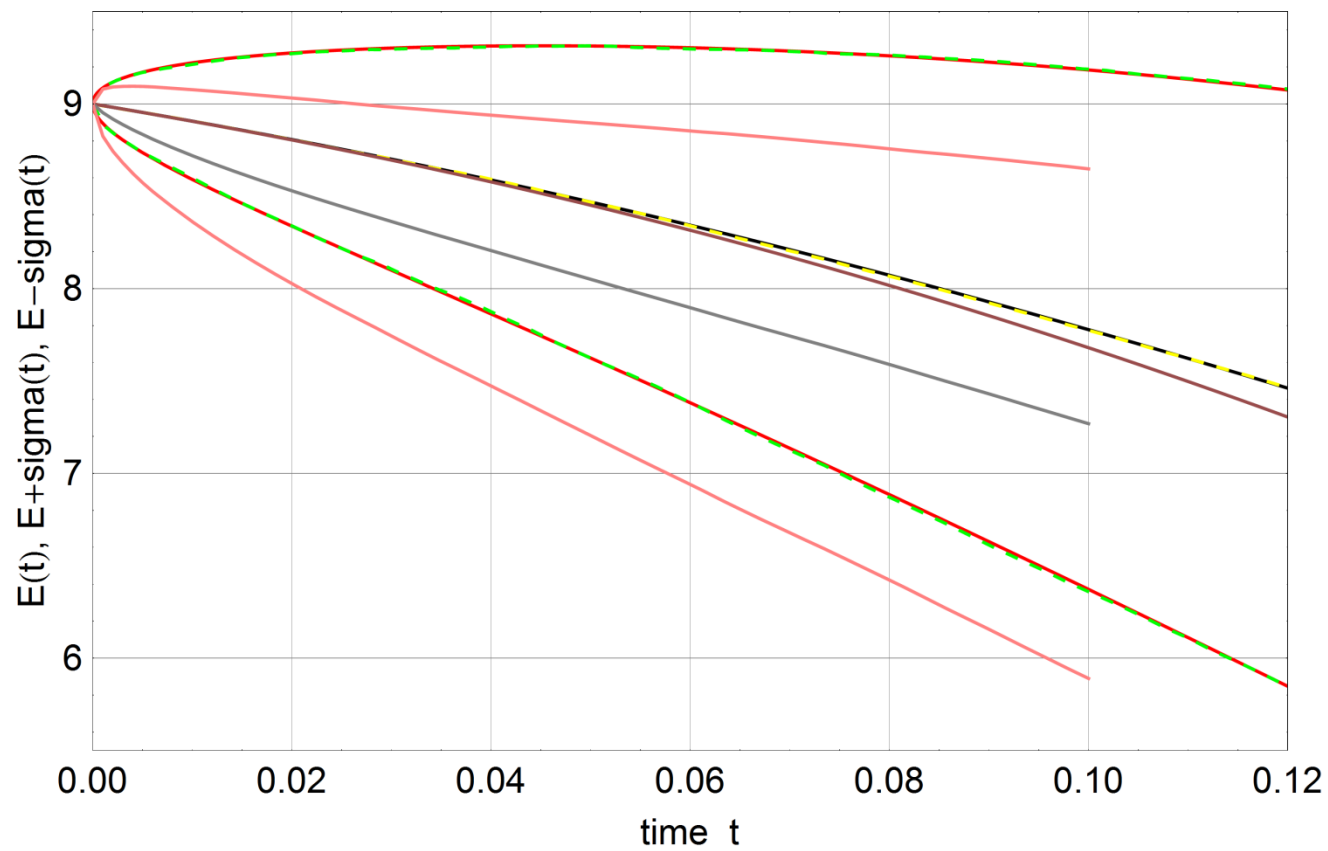


comparison of interpolation order zero and one

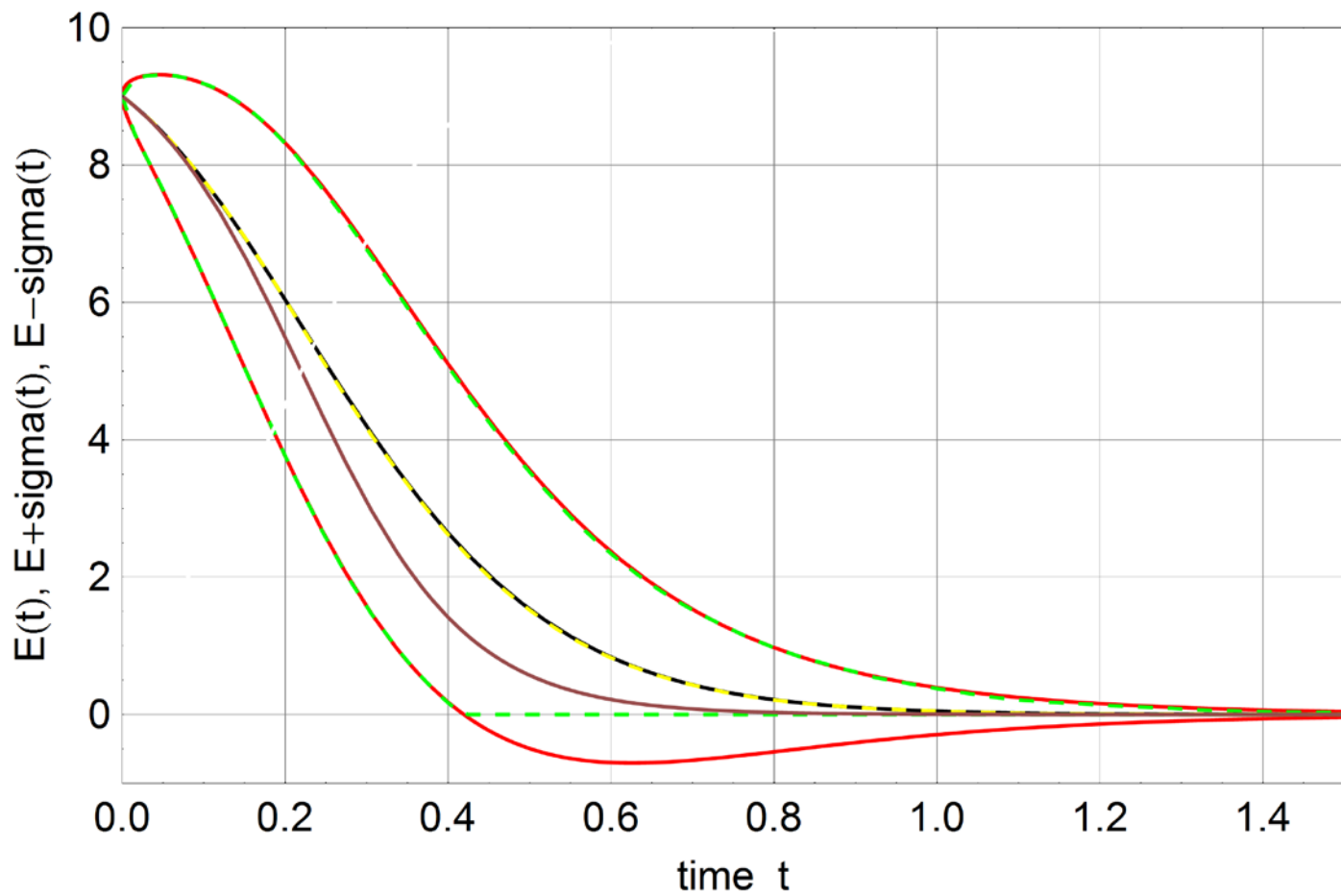
$A \leftrightarrow B$ :  $k = 5, l = 5, N = 10\,000$



$A \leftrightarrow B: k = 5, l = 5, N = 10\ 000$







Thank you for your attention!

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