# Analysis of Markovian Population Models Dissertation Defense

Michael Backenköhler

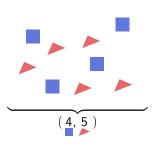
Saarland Informatics Campus

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### Motivation

- stochastic concentration changes
- systems biology: switches,
- ▶ applications: queueing, metabolic networks, bio-switches, traffic etc.
- ▶ goal: do *rigorous* analysis on such models

#### **Semantics**



- ► state space ~ population sizes
- often huge to infinite



- continuous time
- exponential jump times / CTMC dynamics

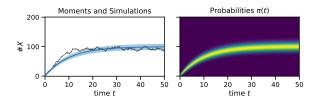
Stationary Distribution - Foster-Lyapunov Functions

lacktriangle ergodic chains converge to unique distribution  $(t o \infty)$ 

#### 

- how does this distribution look like for infinite state-spaces?
- use Foster-Lyapunov function to bound sets
- ▶ locally augment functions for tighter sets / bounds

#### Moment Dynamics



#### Moment formula

Moments such as mean E  $(X_t)$  and variance E  $(X_t^2) - E(X_t)^2$  are described by (often linear) ODEs.

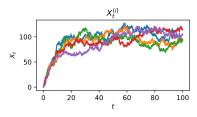
$$\frac{d}{dt} E\left(f(\vec{X}_t)\right) = \sum_{j=1}^{n_R} E\left(\left(f(\vec{X}_t + \vec{v}_j) - f(\vec{X}_t)\right) \alpha_j(\vec{X}_t)\right)$$

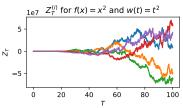
#### Martingale Process

- ▶ multiply time-weighting:  $w(t) = t^k$ ,  $k \in \mathbb{N}$  or  $w(t) = \exp(\lambda t)$
- ▶ analytic integration and resulting martingale process

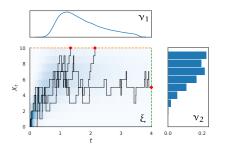
$$Z_T := w(T)f(\vec{X}_T) - w(0)f(\vec{X}_0) - \int_0^T \frac{dw(t)}{dt}f(\vec{X}_t) dt$$
$$-\sum_{j=1}^{n_R} \int_0^T w(t)(f(\vec{X}_t + \vec{v}_j) - f(\vec{X}_t))\alpha_j(\vec{X}_t) dt$$

▶ known expectation:  $E(Z_T) = 0$ ,  $\forall T \ge 0$ 





Linear Moment Constraints



- $\tau = \inf\{X_t \geqslant H \mid t \geqslant 0\} \land T$
- ► exp. occupation measure ξ
- $\blacktriangleright$  exit location measures  $\nu_1$ ,  $\nu_2$

$$\begin{split} 0 = E\left(Z_{T}\right) = T^{k} \overbrace{E\left(X_{\tau}^{m}; \tau = T\right)}^{\nu_{1}} + H^{m} \overbrace{E\left(\tau^{k}; \tau < T, X_{\tau} = H\right)}^{\nu_{2}} \\ - 0^{k} x_{0}^{m} + \sum_{i} c_{i} \underbrace{E\left(\int_{0}^{\tau} t^{k_{i}} X_{t}^{m_{i}} \ dt\right)}_{E} \end{split}$$

Moment Matrices

#### Moment Matrices

The moment matrix must be positive semi-definite.

$$\mathsf{E} \begin{pmatrix} X^0 & X^1 & X^2 & \dots & X^n \\ X^1 & X^2 & \ddots & \ddots & \vdots \\ \vdots & \ddots & \ddots & \ddots & \vdots \\ X^n & X^{n+1} & \dots & \dots & X^{2n} \end{pmatrix} \succeq \mathsf{0}\,,$$

where  $M \succeq 0$  iff.  $\forall v \in \mathbb{R}^n . v^T M v \geqslant 0$ .

### Example

Let 
$$M = \begin{pmatrix} 1 & E(X) \\ E(X) & E(X^2) \end{pmatrix}$$
. Then  $\det M = E(X^2) - E(X)^2 = \sigma^2 \geqslant 0$ .

Semi-Definite Program

- measure support can be restricted using semi-definite constraints
- resulting SDPs can be solved using off-the-shelf software.

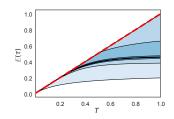
### Semi-Definite Program (SDP)

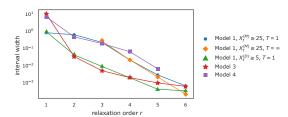
An optimization problem with

- 1. linear constraints on moments and
- 2. positive semi-definite constraints on certain matrices.
- alternative: linear Hausdorff constraints instead of semi-definite constraints

$$\int_{[0,1]^{\vec{\pi}}} \vec{x}^{\vec{\ell}} (1 - \vec{x})^{\vec{k}} d\mu(\vec{x}) \geqslant 0$$

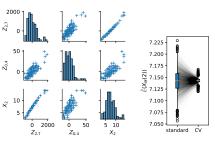
#### Results





- fast convergence of bounds with increasing order
- ► SDPs are usually solved within seconds
- numerically challenging (inherent stiffness)
- scaling state-space / model size is difficult

Using Correlated RVs with Known Expected Value



- ▶ improve MC estimates using Z<sub>T</sub>
- ightharpoonup use correlations between  $Z_T$  and  $X_T$
- $\begin{tabular}{l} E\left(X_T+bZ_T\right) \text{ instead of } E\left(X_T\right) \text{ (recall } E\left(Z_T\right)=0\text{)} \end{tabular}$

#### **Linear Control Variates**

Given a control variate vector  $\vec{Z}$ , the estimator

$$\hat{V} - (\hat{\Sigma}_{\vec{Z}}^{-1} \hat{\Sigma}_{\vec{Z}V})^{\mathsf{T}} \hat{\vec{Z}}$$

has lower or equal variance as  $\hat{V}$ .

[BBW19]

Efficiency Trade-off

### cost: slowdown

$$c_{\text{old}}/c_{\text{new}}$$

- ightharpoonup computing  $\int_0^T w(t) X_t^m dt$
- computing the estimate

#### benefit: variance reduction

$$\sigma_{\text{new}}^2 \big/ \sigma_{\text{old}}^2$$

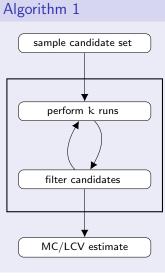
highly correlated variates

Approach: Assess correlations between k candidates and the target RV V

$$\begin{bmatrix} 1 & \dots & \rho_{1k} & \rho_{1\nu} \\ \vdots & \ddots & \vdots & \vdots \\ \rho_{k1} & \dots & 1 & \rho_{k\nu} \\ \rho_{\nu1} & \dots & \rho_{\nu k} & 1 \end{bmatrix}.$$

[BBW19; BBW22]

Selection by Filtering

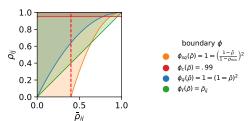


#### Filter criteria:

1. low target correlation

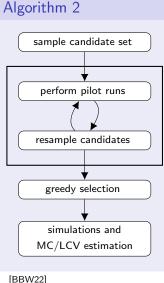
$$\rho_{i\nu} < \text{max}\left(0.1, \frac{\text{max}_{j} \; \rho_{j\nu}}{k_{\text{min}}}\right)$$

2. various redundancy heuristics: criteria based on  $\rho_{ij}$  and  $\rho_{i\nu}$ 



[BBW19]

Selection by Resampling



Filter criteria:

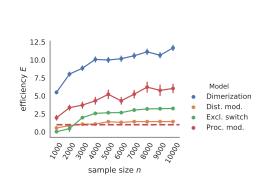
1. resampling proportional to improvement

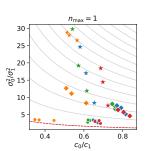
$$\gamma_{k\nu} = (1-\rho_{k\nu}^2)^{-1}$$

2. selection by best improvement of one CV

$$\underset{1\leqslant i\leqslant |P_{\mathsf{all}}|}{\text{arg max}}\,\hat{\gamma}_{i\nu}\prod_{\substack{1\leqslant j\leqslant |P_{\mathsf{all}}|\\ (m_j,\lambda_j)\in P^*}}\hat{\gamma}_{ij}^{-1}$$

#### Results





#### boundary $\phi$

- $\phi_{sq}(\bar{\rho}) = 1 \left(\frac{1-\bar{\rho}}{1-\rho_{min}}\right)^2$
- $\phi_c(\bar{\rho}) = .99$
- $\phi_q(\bar{\rho}) = 1 (1 \bar{\rho})^2$
- $\phi_{\ell}(\bar{\rho}) = \bar{\rho}_{ij}$

#### Model

- Dimerization
- Exclusive Switch
- Dist. Modification

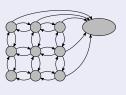
# Finite-Space Projection

### Original



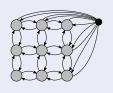
- very large/infinite
- impossible to analyze

### Sink state



- transient analysis
- keep track of approx. error

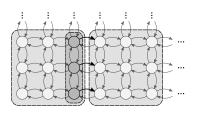
#### Redirection



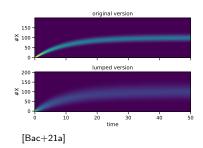
- stationary dist.
- dependent on redirection

# State-Space Aggregation

Treating Hyper-Cubes of States as One



- hyper-cube macro-states
- assumption: uniform dist. within
- closed-form transition rates

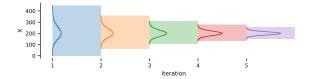


- resulting distribution more "flat"
- locate main probability mass

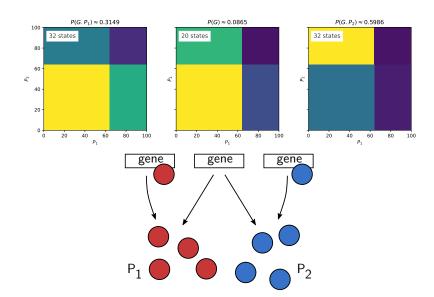
#### Iterative Refinement Algorithm

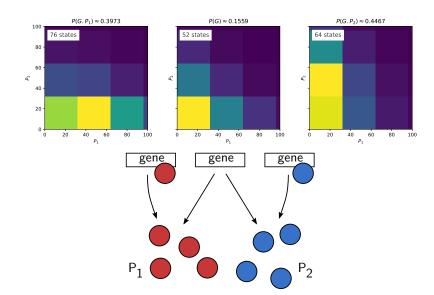
A simple refinement based on approximate solutions:

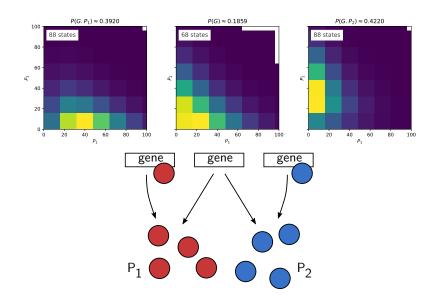
- 1. start with macro-states of size 2k
- 2. compute approximate distribution
- 3. remove states with low probability
- 4. split the remaining states
- 5. go to step 2

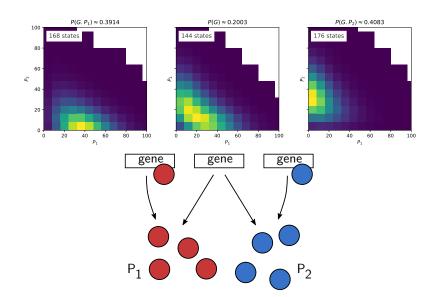


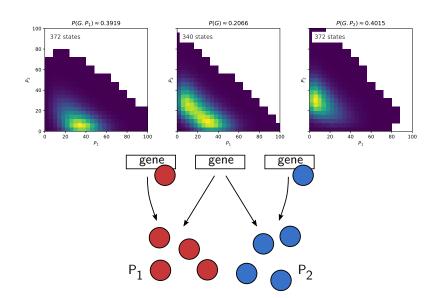
 $[\mathsf{Bac}{+}21\mathsf{b};\,\mathsf{Bac}{+}21\mathsf{a}]$ 

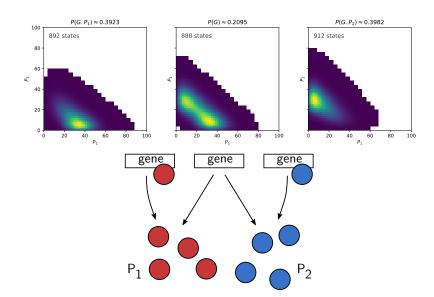


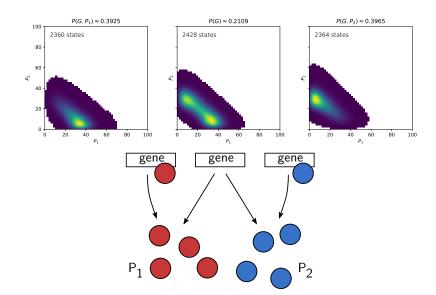






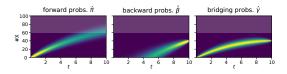






### **Bridging Problem**

Dynamical Analysis Under Initial and Terminal Constraints



#### Forward Probabilities $\pi$

How the process evolves with time:  $Pr(X_t = x \mid X_0 = 0)$ 

### Backward Probabilities β

Probability of ending up in a given state:  $Pr(X_T = 40 \mid X_t = x)$ 

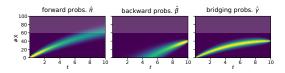
### Bridging Probabilities $\gamma$

In between:  $Pr(X_t = x | X_0 = 0, X_T = 40)$ 

[Bac+21b]

### **Bridging Problem**

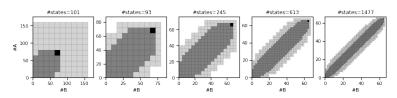
#### Refinement



bridging distribution:

$$\gamma(x_{\text{i}},t) = \pi(x_{\text{i}},t)\beta(x_{\text{i}},t)/\pi(x_{g},T)$$

- record intermediary times
- remove or split based on  $\hat{\gamma}(x_i, t)$



[Bac+21b]

### **Bridging Problem**

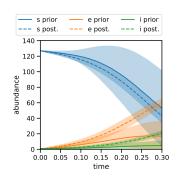
Bayesian Filtering in an SEIR model

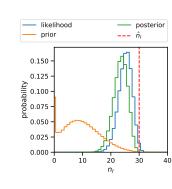
$$S + I \rightarrow E + I$$
  $E \rightarrow I$   $I \rightarrow R$ 

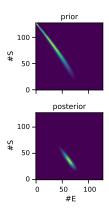
We know:

- initial state
- ightharpoonup uncertain measurement of I at T = 0.3

We are interested in the posterior at T.







[Bac+21b]

### Contributions

- ▶ local augmentation of Foster-Lyapunov functions
- bounding of mean first-passage times [BBW20]
- ▶ variance reduction for MC estimation [BBW19; BBW22]
- state-space aggregation scheme
  - stationary distribution [Bac+21a]
  - bridging distribution [Bac+21b]
  - importance sampling

### References I

[BBW20] Michael Backenköhler, Luca Bortolussi, and Verena Wolf. "Bounding Mean First Passage Times in Population Continuous-Time Markov Chains". In: 17th International Conference on Quantitative Evaluation of SysTems. Vol. 12289. Lecture Notes in Computer Science. Springer, 2020, pp. 155–174.

[BBW19] Michael Backenköhler, Luca Bortolussi, and Verena Wolf. "Control Variates for Stochastic Simulation of Chemical Reaction Networks". In: 17th International Conference on Computational Methods in Systems Biology. Vol. 11773. Lecture Notes in Computer Science. Springer, 2019, pp. 42–59.

[BBW22] Michael Backenköhler, Luca Bortolussi, and Verena Wolf. "Variance Reduction in Stochastic Reaction Networks using Control Variates". In: Principles of Systems Design – Essays Dedicated to Thomas A. Henzinger on the Occasion of His 60th Birthday. Vol. 13660. Lecture Notes in Computer Science. Springer, 2022.

### References II

- [Bac+21a] Michael Backenköhler et al. "Abstraction-Guided Truncations for Stationary Distributions of Markov Population Models". In: 18th International Conference on Quantitative Evaluation of SysTems. Vol. 12846. Lecture Notes in Computer Science. Springer, 2021, pp. 351–371.
- [Bac+21b] Michael Backenköhler et al. "Analysis of Markov Jump Processes under Terminal Constraints". In: 27th International Conference on Tools and Algorithms for the Construction and Analysis of Systems. Vol. 12651. Lecture Notes in Computer Science. Springer, 2021, pp. 210–229.