Analysis of Markovian Population Models Dissertation Defense

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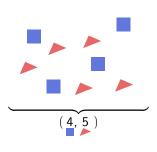
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Motivation

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

Semantics



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



- continuous time
- exponential jump times / CTMC dynamics

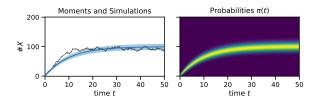
Stationary Distribution - Foster-Lyapunov Functions

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

Example $x \xrightarrow{0.1 \cdot \# X} \varnothing, \quad \alpha_1(x) = 0.1x \\ \varnothing \xrightarrow{10} X, \quad \alpha_2(x) = 10$

- ▶ 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.

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

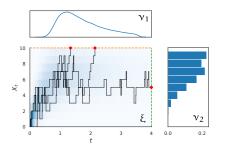
Martingale Process

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

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

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

Linear Moment Constraints



- $\tau = \inf\{X_t \geqslant H \mid t \geqslant 0\} \wedge T$
- \triangleright exp. occupation measure ξ
- ightharpoonup exit location measures v_1 , v_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 mus 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.

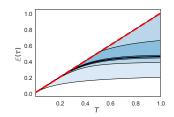
Semi-Definite Program (SDP)

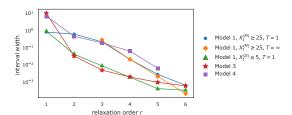
An optimization problem with

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

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

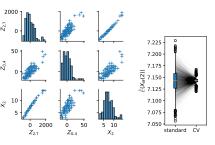
Results





- ▶ fast convergence of bounds with increasing order
- numerically challenging (inherent stiffness)

Using Correlated RVs with Known Expected Value



- ▶ improve MC estimates using Z_T
- lacktriangle 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

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

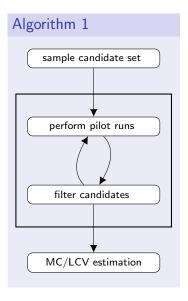
highly correlated variates

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

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

[BBW19; BBW22]

Selection by Filtering



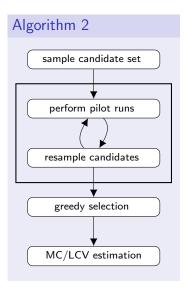
Filter criteria:

1. low target correlation

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

2. redundancy criterion

Selection by Resampling



Filter criteria:

1. low target correlation

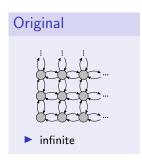
$$\rho_{i\nu} < \text{max}\left(0.1, \frac{\text{max}_j \; \rho_{j\nu}}{k_{\text{min}}}\right)$$

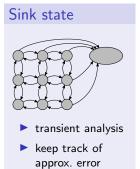
2. redundancy criterion

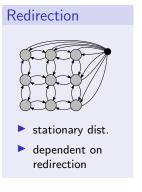
Results

best example?

Finite-Space Projection

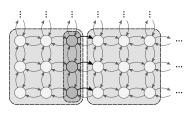




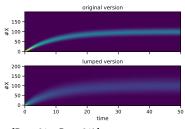


State-Space Aggregation

Treating Hyper-Cubes of States as One



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



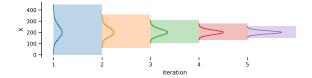
- resulting distribution more "flat"
- locate main probability mass

Stationary Distribution

Iterative Refinement Algorithm

A simple refinement based on approximate solutions:

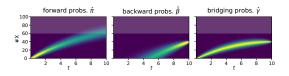
- 1. start with macro-states of size 2k
- 2. compute approximate distribution
- 3. remove states with low probability
- 4. split the remaining states
- 5. go to step 2



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

Bridging Problem

Dynamical Analysis Under Initial and Terminal Constraints



Forward Probabilities

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

Backward Probabilities

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

Bridging Probabilities

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

[Bac+21a]

Bridging Problem

Refinement

► Show that bridge=forward*backwards

Importance Sampling

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+21b]
 - bridging distribution [Bac+21a]
 - 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. "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.
- [Bac+21b] 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.