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A Krylov-based FSP algorithm for solving the chemical master equation arising in the discrete modelling of biological systems

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Contents

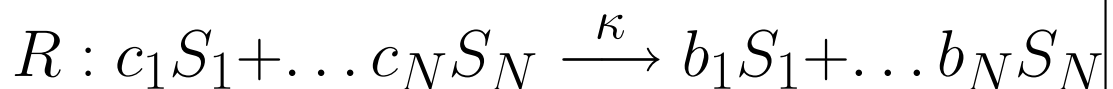
- Stochastic models for biochemical systems
- The Finite State Projection Algorithm (FSP)
- The Krylov FSP
- Numerical Results

Joint work with

- Kevin Burrage (UQ)
- Markus Hegland (ANU)
- Shev MacNamara (UQ)

Chemical reaction systems

- Well-stirred mixture.
- N molecular species: S_1, \dots, S_N .
- Constant temperature, fixed volume Ω
- M reaction channels: R_1, \dots, R_N



κ : reaction rate constant.

c_i, b_i : coefficients of reactants & products

- Dynamical state: $\mathbf{x}(t_0) = \mathbf{x}_0$, $\mathbf{x}(t) = (x_1(t), \dots, x_N(t))^T$, where $x_i(t)$ is the number of S_i molecules in the system.
- Propensity function: $\alpha_j(\mathbf{x})dt =$ the probability, given $\mathbf{x}(t) = \mathbf{x}$, that one reaction R_j will occur in Ω in $[t, t + dt)$.
- Stoichiometric vector: $\boldsymbol{\nu}_j$. When R_j occurs, $\boldsymbol{\nu}_j$ is used to update the state of the system by $\mathbf{x}(t + dt) = \mathbf{x}(t) + \boldsymbol{\nu}_j$.
 $\boldsymbol{\nu} = (b_1 - c_1, \dots, b_N - c_N)^T$.

Chemical Master Equation (CME)

Let $P(\mathbf{x}; t)$ be the probability of being in state \mathbf{x} at time t . Then it satisfies the discrete parabolic PDE

$$\begin{aligned} \frac{\partial P(\mathbf{x}; t)}{\partial t} = & \sum_{j=1}^M \alpha_j(\mathbf{x} - \boldsymbol{\nu}_j) P(\mathbf{x} - \boldsymbol{\nu}_j; t) \\ & - P(\mathbf{x}; t) \sum_{j=1}^M \alpha_j(\mathbf{x}) \end{aligned}$$

ODE formulation

Let $\mathbf{p}(t)$ be a vector of probabilities $P(\mathbf{x}; t)$, indexed by the states \mathbf{x} . Then

$$\dot{\mathbf{p}}(t) = \mathbf{A}\mathbf{p}(t)$$

where $\mathbf{A} = [a_{ij}]$ is the transition matrix populated by the propensities. The rows and columns of the matrix are indexed by the states, so the states have now been enumerated.

Solution techniques

- Stochastic Simulation Algorithm (SSA):
evolve $\mathbf{x}(t)$ by $\mathbf{x}(t + dt) = \mathbf{x}(t) + \boldsymbol{\nu}_j$.
Trajectories.
- CME Solver: $\mathbf{p}(t) = \exp(t\mathbf{A})\mathbf{p}_0$.

Issues

- Large state space
- Long integration domain

Finite State Projection Algorithm

FSP [Munsky/Khammash (J. Chem. Phys., 2006)]

$$\mathbf{A} = \left(\begin{array}{c|c} \mathbf{A}_k & * \\ \hline * & * \end{array} \right)$$
$$e^{t_f \mathbf{A}} \mathbf{v} \approx e^{t_f \mathbf{A}_k} \mathbf{v}_k$$

The states indexed by $\{1, \dots, k\}$ form the *finite state projection*, denoted \mathbf{X}_k .

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Theorem. Let

$$\Gamma_k = \mathbb{1}^T e^{t_f \mathbf{A}_k} \mathbf{p}_k(0), \quad \mathbb{1} = (1, \dots, 1)^T$$

Then

□ $\Gamma_k \leq \Gamma_{k+1} \leq \dots \leq 1.$

□ If $\Gamma_k \geq 1 - \epsilon$ then

$$\begin{pmatrix} e^{t_f \mathbf{A}_k} \mathbf{p}_k(0) \\ \mathbf{0} \end{pmatrix} \leq \mathbf{p}(t_f) \leq \begin{pmatrix} e^{t_f \mathbf{A}_k} \mathbf{p}_k(0) \\ \mathbf{0} \end{pmatrix} + \epsilon \mathbb{1}.$$

FSP Algorithm

```
ALGORITHM: FSP( $\mathbf{A}$ ,  $\mathbf{p}_0(0)$ ,  $t_f$ ,  $\epsilon$ ,  $\mathbf{X}_0$ )  
 $\mathbf{A}_0 := \text{submatrix}(\mathbf{X}_0)$ ;  
 $\Gamma_0 := \mathbb{1}^T \exp(t_f \mathbf{A}_0) \mathbf{p}_1(0)$ ;  
for  $k := 1, 2, \dots$  until  $\Gamma_k \geq 1 - \epsilon$  do  
     $\mathbf{X}_k := \text{expand}(\mathbf{X}_{k-1})$ ;  
     $\mathbf{A}_k := \text{submatrix}(\mathbf{X}_k)$ ;  
     $\Gamma_k := \mathbb{1}^T \exp(t_f \mathbf{A}_k) \mathbf{p}_k(0)$ ;  
endfor  
return  $\exp(t_f \mathbf{A}_k) \mathbf{p}_k(0)$ ;
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```

Issues

- ❑ Implemented via MATLAB's *expm*
- ❑ No intermediate time points
- ❑ Repeated $e^{t_f \mathbf{A}_k} \mathbf{p}_k(0)$ with the *same* t_f

The new Krylov FSP

- Krylov-based instead of $\exp m$

$$e^{\tau \mathbf{A}} \mathbf{v} \approx \beta \mathbf{V}_{m+1} \exp(\tau \bar{\mathbf{H}}_{m+1}) \mathbf{e}_1$$

- Embedded exponentials

$$e^{t \mathbf{A}} \mathbf{v} \approx e^{\tau_k \mathbf{A}_k} \dots e^{\tau_0 \mathbf{A}_0} \mathbf{v}$$

- Inexact/relaxed matrix-vector products

Embedded exponentials

$$\mathbf{p}(t_f) \approx e^{\tau_K \mathbf{A}_K} \dots e^{\tau_0 \mathbf{A}_0} \mathbf{p}(0), t_f = \sum_{k=0}^K \tau_k$$

- Harness Expokit's step-by-step scheme.
- The \mathbf{A}_k changes between internal steps!

Integration scheme (dense output)

$$0 \equiv t_0 < t_1 < \dots < t_K < t_{K+1} \equiv t_f$$

$$\mathbf{p}(t_{k+1}) = \mathbf{p}(t_k + \tau_k) = e^{\tau_k \mathbf{A}_k} \mathbf{p}(t_k).$$

Embbded exponentials

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Krylov FSP criteria

- $\Gamma_k = \mathbb{1}^T \exp(\tau_k \mathbf{A}_k) \mathbf{p}(t_k) \geq 1 - \epsilon \frac{t_k}{t_f}$
 $\tau_k := \tau_k/2$ until the criteria is met.
- Expand the FSP

Embbded exponentials

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- ❑ Expand the FSP

Accuracy criteria

- ❑ Select τ_k for a good approximation
- ❑ Integrate without expanding the FSP

Inexact matrix-vector product

Don't use the finite state projection \mathbf{X}_k entirely. Only capture an approximation:

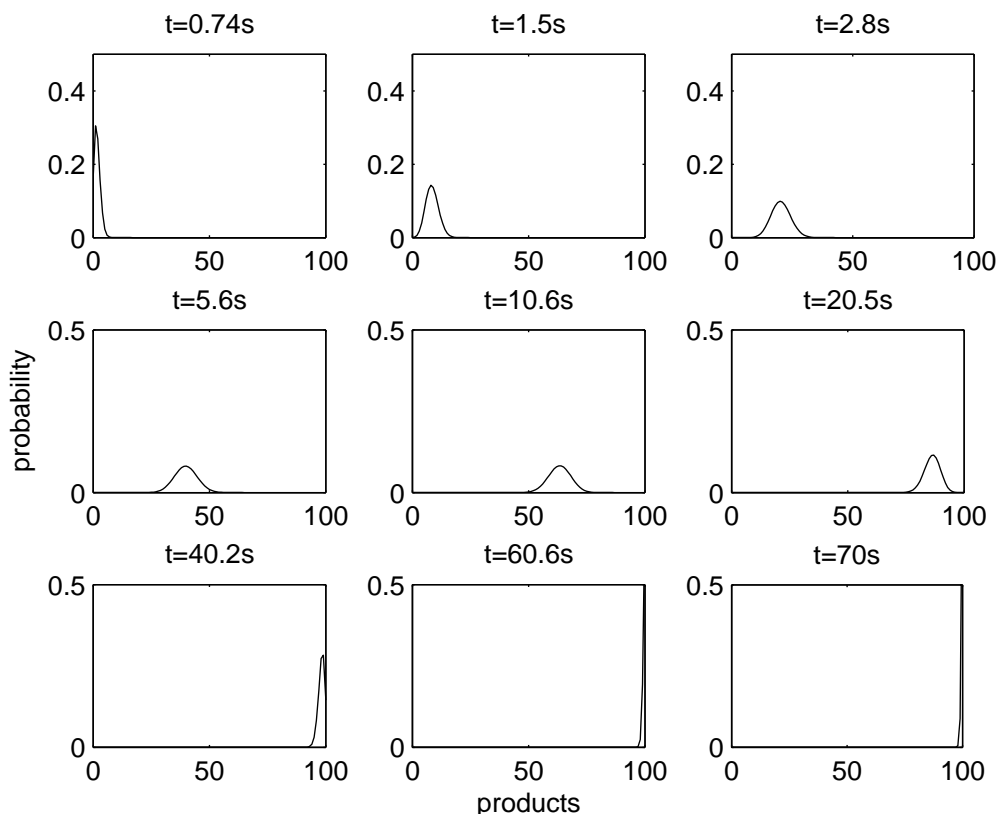
$$\begin{aligned}\mathbf{A}_k \mathbf{v} &= \begin{bmatrix} \mathbf{a}_1^{(k)} & \mathbf{a}_2^{(k)} & \dots \end{bmatrix} \begin{bmatrix} v_1 \\ v_2 \\ \vdots \end{bmatrix} \\ &\approx \sum_{i \in \hat{\mathbf{X}}_k} v_i \mathbf{a}_i^{(k)} \\ &= \hat{\mathbf{A}}_k \mathbf{v}.\end{aligned}$$

Inexact matrix-vector product

Do not do the entire finite state projection \mathbf{X}_k .

Capture only an approximation:

$$\begin{aligned}\mathbf{A}_k \mathbf{v} &= \begin{bmatrix} \mathbf{a}_1^{(k)} & \mathbf{a}_2^{(k)} & \dots \end{bmatrix} \begin{bmatrix} v_1 \\ v_2 \\ \vdots \end{bmatrix} \\ &\approx \sum_{i \in \hat{\mathbf{X}}_k} v_i \mathbf{a}_i^{(k)} \\ &= \hat{\mathbf{A}}_k \mathbf{v}.\end{aligned}$$



Krylov FSP tracks the support.

Krylov FSP Algorithm

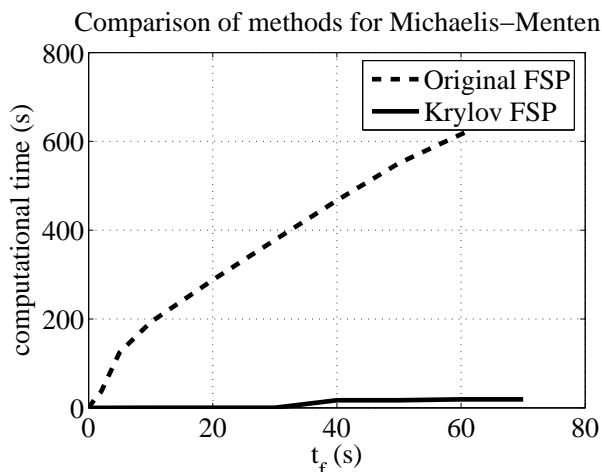
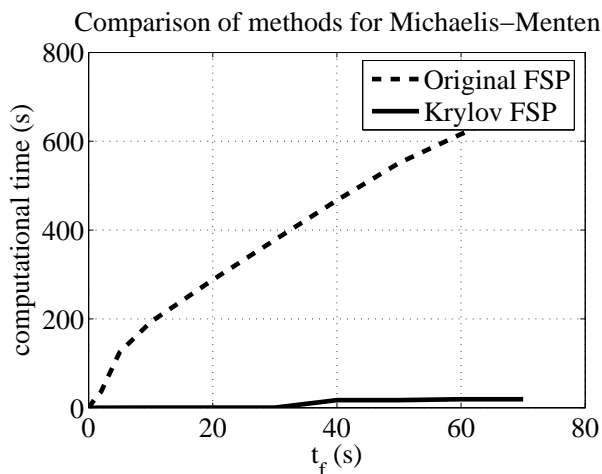
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ALGORITHM: KrylovFSP( $\mathbf{A}, \mathbf{p}_0(0), t_f, \epsilon, \mathbf{X}_0, m, tol$ )
 $t_0 := 0$ ;  $\hat{\mathbf{X}}_0 := \mathbf{X}_0$ ;
 $\mathbf{A}_0 := \text{submatrix}(\mathbf{X}_0)$ ;  $\hat{\mathbf{A}}_0 := \mathbf{A}_0$ ;
expandFSP := FALSE;
for  $k := 0, 1, 2, \dots$  until  $t_k = t_f$  do
   $[\mathbf{V}_{m+1}, \overline{\mathbf{H}}_{m+1}] := \text{Arnoldi}(\hat{\mathbf{A}}_k, \mathbf{p}_k(t_k), m)$ ;
  repeat
     $\tau_k := \text{step-size}$ ;
     $\mathbf{p}_k(t_k) := \beta \mathbf{V}_{m+1} \exp(\tau_k \overline{\mathbf{H}}_{m+1}) \mathbf{e}_1$ ;
    err := numerical-error-estimate;
  until err  $\leq 1.2 tol$ ;
   $\Gamma_k := \mathbb{1}^T \mathbf{p}_k(t_k)$ ;
  while  $\Gamma_k < 1 - \epsilon \frac{t_k + \tau_k}{t_f}$  do
    expandFSP := TRUE;
     $\tau_k := \frac{1}{2} \tau_k$ ;
     $\mathbf{p}_k(t_k) := \beta \mathbf{V}_{m+1} \exp(\tau_k \overline{\mathbf{H}}_{m+1}) \mathbf{e}_1$ ;
     $\Gamma_k := \mathbb{1}^T \mathbf{p}_k(t_k)$ ;
  endwhile
  if expandFSP
     $\mathbf{X}_{k+1} := \text{expand}(\mathbf{X}_k)$ ;  $\hat{\mathbf{X}}_{k+1} := \text{select}(\mathbf{X}_{k+1})$ ;
     $\mathbf{A}_{k+1} := \text{submatrix}(\mathbf{X}_{k+1})$ ;  $\hat{\mathbf{A}}_{k+1} := \text{submatrix}(\hat{\mathbf{X}}_{k+1})$ ;
    expandFSP := FALSE;
  endif
   $t_{k+1} := t_k + \tau_k$ ;
endfor
return  $\mathbf{p}(t_k)$ 

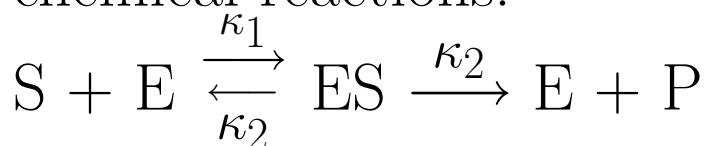
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Numerical Results

Tests on a SGI Altix (with $tol = 10^{-5}$)



The Michaelis-Menten enzyme kinetics involve four species, substrates (S), enzymes (E), enzyme-substrate complexes (ES) and products (P), interacting through three chemical reactions:



Conclusion

- ❑ Krylov FSP is *much* faster, albeit sometimes less accurate.
- ❑ It is better at memory management.
- ❑ Dense output is readily available.
- ❑ Best suited for problems where the support
 - ★ spreads slowly
 - ★ confined.
- ❑ Large t_f is a challenge for all methods.