

Thrombin system reduction

Hoang M. Tran

November 14, 2014

Outline

- 1 Problem formulation
- 2 l_1 - min formulation
- 3 Data driven system reduction method
 - Correlation analysis of time series from simulated human coagulation cascade dynamics
 - System reduction using linear transformation
- 4 Update week of November 14th 2014
 - Kuramoto model reconstruction: bound on total perturbation
 - Dimension reduction

Outline

- 1 Problem formulation
- 2 l_1 - min formulation
- 3 Data driven system reduction method
 - Correlation analysis of time series from simulated human coagulation cascade dynamics
 - System reduction using linear transformation
- 4 Update week of November 14th 2014
 - Kuramoto model reconstruction: bound on total perturbation
 - Dimension reduction

Problem formulation

$$\dot{\mathbf{y}}(t) = \mathbf{f}(\mathbf{y}(t), \mathbf{k}) \quad (1)$$

where

- $\mathbf{y} = (y_1, \dots, y_n)$, $n = 34$, y_i : concentration of chemical species i
- \mathbf{k} : reaction rates, given constants
- $\mathbf{f}(\mathbf{y}(t), \mathbf{k}) = (f_1(\mathbf{y}(t), \mathbf{k}), \dots, f_m(\mathbf{y}(t), \mathbf{k}))$, $m = 42$, there are 42 rate equation
- y_{14} : concentration of Thrombin

Problem formulation

Find another system of equations in terms of new variable \mathbf{z} such that

- $\mathbf{z}(t)$ is a transformation of $\mathbf{y}(t)$
- $\dim(\mathbf{z}) \leq 4$

• Same estimation error $\|\mathbf{y}(t) - \hat{\mathbf{y}}(t)\|$ and $\|\mathbf{z}(t) - \hat{\mathbf{z}}(t)\|$

Outline

- 1 Problem formulation
- 2 l_1 - min formulation
- 3 Data driven system reduction method
 - Correlation analysis of time series from simulated human coagulation cascade dynamics
 - System reduction using linear transformation
- 4 Update week of November 14th 2014
 - Kuramoto model reconstruction: bound on total perturbation
 - Dimension reduction

l_1 - min formulation l_1 - min formulation

$$\min |C|_1 \text{ s. t. } \begin{cases} \|\dot{\mathbf{z}}(t) - \mathbf{f}(\mathbf{z}(t), C)\|_k & \leq \varepsilon_1 \forall t \in \{t_1, \dots, t_N\} \\ \|\mathbf{z}_{14} - \mathbf{f}_{14}(\mathbf{z}(t), C)\|_k & \leq \varepsilon_2 \end{cases}$$

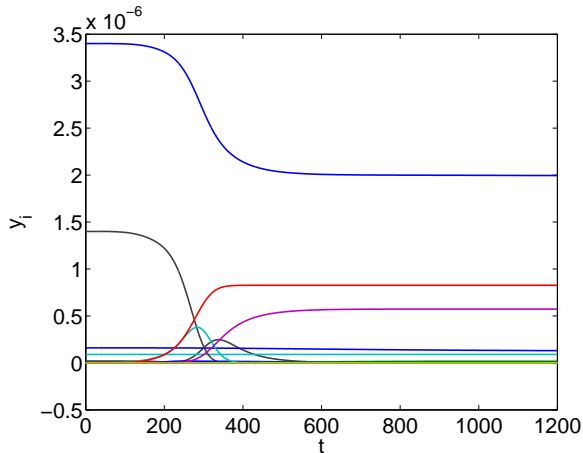
where $\varepsilon_1 > \varepsilon_2$

Note that the transformation: $\mathbf{y} \rightarrow \mathbf{z} = \mathbf{y}$.

Outline

- 1 Problem formulation
- 2 l_1 - min formulation
- 3 **Data driven system reduction method**
 - Correlation analysis of time series from simulated human coagulation cascade dynamics
 - System reduction using linear transformation
- 4 Update week of November 14th 2014
 - Kuramoto model reconstruction: bound on total perturbation
 - Dimension reduction

Time series



Time series: observations

- Time series are highly correlated. 14 \rightarrow 25: correlation about 1.
- Simulated data Y when solving (1) can be approximately represented in lower dimension space

SVD the simulated data $Y_{n \times N} \xrightarrow{A} Y_{n \times N} \approx A_{n \times p} Z_{p \times N}, p \ll n$, use A as transformation to reduce the dynamical system (1)

Dimension reduction using singular value decomposition

$$\begin{aligned}
 C_{N \times n} &= S_{N \times N} \Sigma_{N \times n} (V_{n \times n})^T \\
 \tilde{C} &= Z_{N \times p} \Sigma_{p \times p} (V_{p \times n})^T \\
 &\approx C \\
 \tilde{C}^T &= (\Sigma_{p \times p} (V_{p \times n})^T)^T S_{N \times p}^T
 \end{aligned} \tag{2}$$

- N : sample size, p : reduced dimension.

Apply (2) to $C = Y_{n \times N}^T$, denote $Z_{p \times N} = S_{N \times p}^T$,

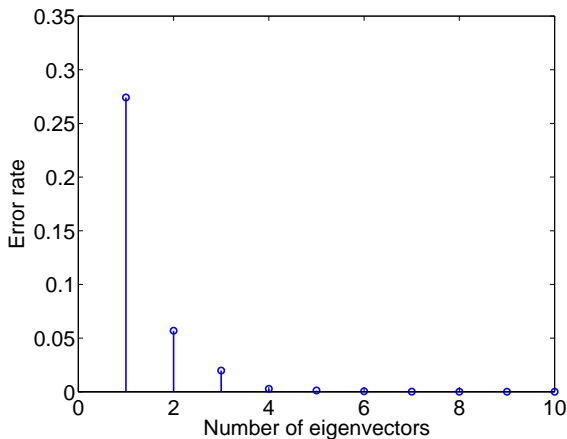
$$A = (\Sigma_{p \times p} (V_{p \times n})^T)^T = V_{p \times n} \Sigma_{p \times p}^T$$

$$\tilde{Y}_{n \times N} = A Z_{p \times N} \tag{3}$$

$$y(t) = A z(t) \tag{4}$$

$$z(t) = A^{-1} y(t) \tag{5}$$

Dimension reduction using singular value decomposition



Error rate: $\frac{\|\tilde{Y}^T - Y\|_F}{\|Y\|_F}$

Dimension reduction using linear transformations: singular value decomposition

$$\begin{aligned}
 \mathbf{y}(t) &\approx \mathbf{A}\mathbf{z}(t) \\
 \mathbf{z}(t) &\approx \mathbf{A}^{-1}\mathbf{y}(t) \\
 \frac{d\mathbf{z}(t)}{dt} &= \mathbf{A}^{-1}\frac{d\mathbf{y}(t)}{dt} \\
 \frac{d\mathbf{z}(t)}{dt} &= \mathbf{A}^{-1}\mathbf{f}(\mathbf{y}(t), \mathbf{k}) \\
 \frac{d\mathbf{z}(t)}{dt} &\approx \mathbf{A}^{-1}\mathbf{f}(\mathbf{A}\mathbf{z}(t), \mathbf{k})
 \end{aligned} \tag{6}$$

- \mathbf{A}^{-1} : may need to use pseudo inverse.
- From $\mathbf{z}(t)$ we can get Thrombin and other components by $\mathbf{y}(t) = \mathbf{A}\mathbf{z}(t)$.

(6) - equation for the reduced system

Validation procedure

- 1 Solve the original system numerically to get $Y_{N \times n}$,
 $N = 1200, n = 34$, **not 29**.
- 2 Do SVD $Y_{N \times n} = Z_{N \times N} \Sigma_{N \times n} (V_{n \times n})^T$, $A = (\Sigma_{p \times p} (V_{p \times n})^T)^T$,
 $\tilde{Y}^{svd} = (\Sigma_{p \times p} (V_{p \times n})^T)^T Z_{N \times p}^T$
- 3 Solve $\dot{z}(t) \approx A^{-1} f(Az(t), k)$ to get $(Z_{N \times p}^{reduced})^T$
- 4 Compute $(\tilde{Y}^{reconstruct})^T = A(Z_{N \times p}^{reduced})^T$
- 5 Compute error rate using $\tilde{Y}^{reconstruct}$, $\tilde{Y}_{N \times n}^{svd}$

Result - observations

(1)	1	2	3	4	5	6	7	8	9	10
(2)	-	stiff	-	stiff	stiff	-	stiff	stiff	stiff	stiff
(3)	0.86	5.78	0.17	18	108	0.06	0.07	6.68	5.57	1.8

Row (1) - number of eigenvectors, p or $\dim(\mathbf{z})$

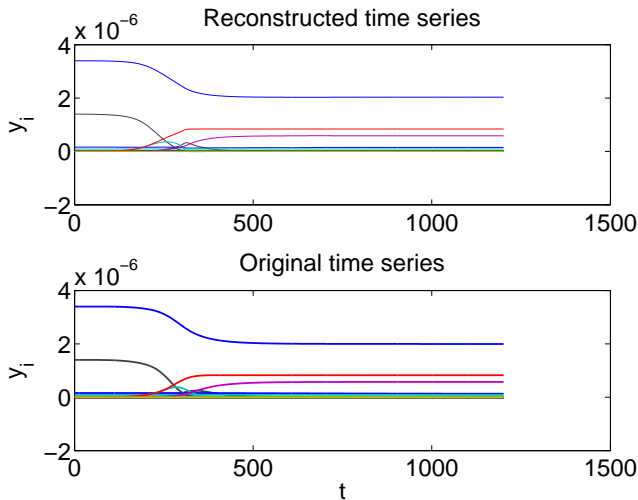
Row (2) - nonstiffness of the reconstructed system

$$\dot{\mathbf{z}}(t) \approx \mathbf{A}^{-1} \mathbf{f}(\mathbf{A}\mathbf{z}(t), \mathbf{k})$$

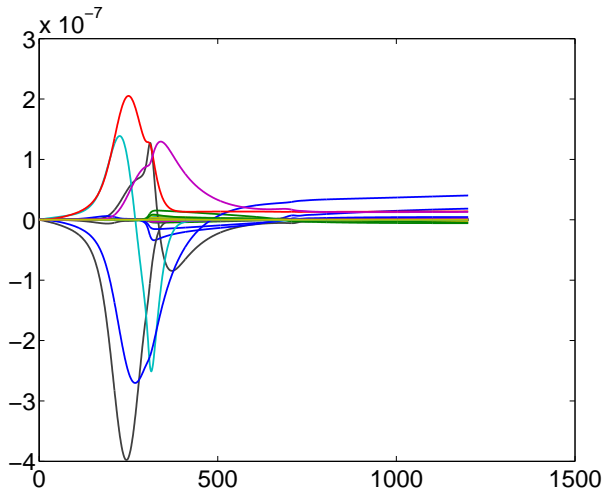
Rows (3): error rate computed using $\tilde{\mathbf{Y}}^{reconstruc}$

- Training data length does matter
- The new system might become very stiff
- Accuracy does not linearly depends on training data length
- $p = 3$ or $p = 6$

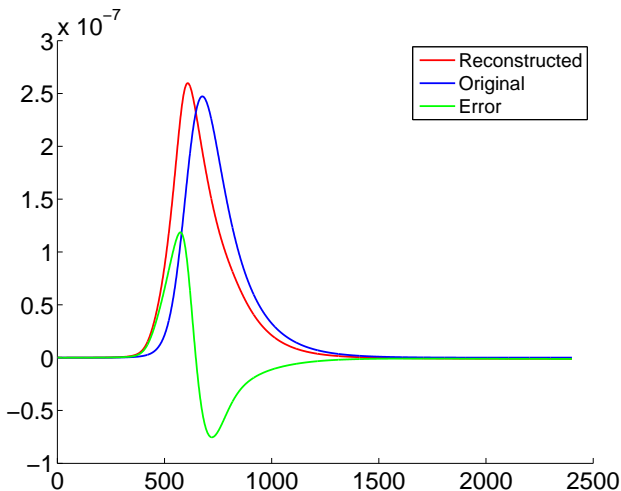
Time series comparison $p = 6$



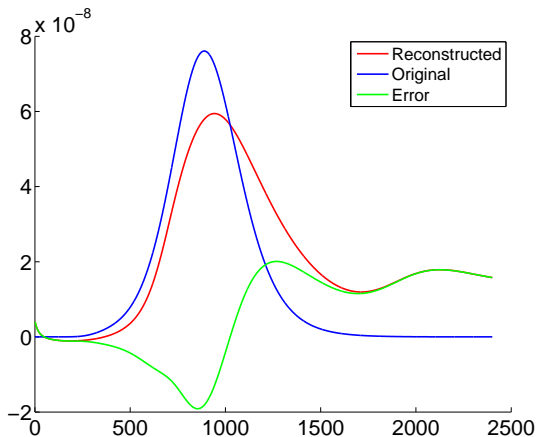
Time series comparison - error



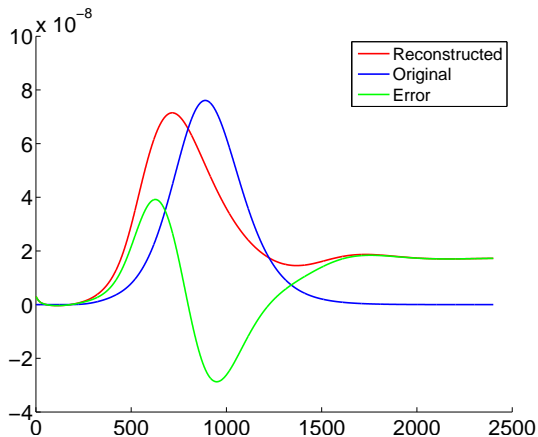
Thrombin time series



Thrombin time series - using the one initial condition, use another initial condition for validation



Thrombin time series - using same initial conditions



Result - some observations

- Using higher dimension space to represent new variable does not guarantee smaller error
- The approximated system might become very stiff
- A depends on initial condition in the original system

⇒ need a mathematical formulation and perturbation analysis

Outline

- 1 Problem formulation
- 2 l_1 - min formulation
- 3 Data driven system reduction method
 - Correlation analysis of time series from simulated human coagulation cascade dynamics
 - System reduction using linear transformation
- 4 Update week of November 14th 2014
 - Kuramoto model reconstruction: bound on total perturbation
 - Dimension reduction

Kuramoto model reconstruction: Bound on total perturbation

Theorem

For matrices R, R^0 if $\|R - I\|_F + \|\Delta R\|_F < 1$ we have:

$$\|R^{-1} - (R^0)^{-1}\|_F \leq \|\Delta R\|_F \frac{1}{(1 - \|R - I\|_F - \|\Delta R\|_F)^2}$$

where $\Delta R = R - R^0$.

Proof.

Apply the Theorem 6.2.30 from Johnson's book for:

$f(t) = \sum_{k=0}^{\infty} (-1)^k t^k = \frac{1}{1+t}$, $f_{abs}(t) = \sum_{k=0}^{\infty} t^k = \frac{1}{1-t}$, we have

$f'_{abs}(t) = \frac{1}{(1-t)^2}$, $f(R - I) = (I + R - I)^{-1} = R^{-1}$,

$f(R^0 - I) = (R^0)^{-1}$

$$\begin{aligned} \|R^{-1} - (R^0)^{-1}\|_F &= \|f(R - I) - f(R^0 - I)\|_F \\ &\leq \|R - R^0\|_F f'_{abs}(\|R - I\|_F + \|(R - R^0)\|_F) \end{aligned}$$

Kuramoto model reconstruction: Bound on total perturbation

Theorem

For the network untangling method, $P^0 = S^0 R^0$, the total perturbation $P - S^0 R = -(\Delta S)R$ is bounded by

$$(\|P\|_F + \|\Delta P\|_F) \|\Delta R\|_F \frac{1}{(1 - \|R - I\|_F - \|\Delta R\|_F)^2} \|R\|_F + \|\Delta P\|_F$$

Proof.

$$P^0 = S^0 R^0 \Rightarrow S^0 = P^0 (R^0)^{-1} \quad (7)$$

$$S = PR^{-1} \Rightarrow \Delta S = PR^{-1} - P^0 (R^0)^{-1} \quad (8)$$

$$\Delta S = (P^0 + \Delta P)R^{-1} - P^0 (R^0)^{-1} \quad (9)$$

$$= P^0 (R^{-1} - (R^0)^{-1}) + \Delta P R^{-1} \quad (10)$$

$$= (P - \Delta P) (R^{-1} - (R^0)^{-1}) + \Delta P R^{-1} \quad (11)$$

$$(\Delta S)R = (P - \Delta P) (R^{-1} - (R^0)^{-1}) R + \Delta P \quad (12)$$

Problem with SVD based model reduction

SVD the simulated data $Y_{n \times N} \xrightarrow{A} Y_{n \times N} \approx A_{n \times p} Z_{p \times N}, p \ll n$, use A as transformation to reduce the dynamical system (1)

- $Y_{N \times n} = Z_{N \times N} \Sigma_{N \times n} (V_{n \times n})^T$, $A = (\Sigma_{p \times p} (V_{p \times n})^T)^T$,
 $y(t) = Az(t) + \varepsilon_1(t)$
- $\dot{z}(t) \approx A^{-1} f(Az(t), k)$ to get $(Z_{N \times p}^{reduced})^T$
- Compute $(\tilde{Y}^{reconstruct})^T = A(Z_{N \times p}^{reduced})^T$

Problem: Not robust to the number of component used: stiffness issue, big error rate.

Dimension reduction using linear transformations

A, B are linear transformations that we need to define

$$\mathbf{y}(t) = A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t)$$

$$\mathbf{z}(t) = B\mathbf{y}(t) - \boldsymbol{\varepsilon}_2(t)$$

$$\dot{\mathbf{z}}(t) = B\dot{\mathbf{y}}(t) - \dot{\boldsymbol{\varepsilon}}_2(t)$$

$$= B\mathbf{f}(\mathbf{y}(t), \mathbf{k}) - \dot{\boldsymbol{\varepsilon}}_2(t)$$

$$= B\mathbf{f}(A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t), \mathbf{k}) - \dot{\boldsymbol{\varepsilon}}_2(t)$$

$$\dot{\mathbf{z}}(t) - B\mathbf{f}(A\mathbf{z}(t), \mathbf{k}) = B\mathbf{f}(A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t), \mathbf{k}) - \dot{\boldsymbol{\varepsilon}}_2(t) - B\mathbf{f}(A\mathbf{z}(t), \mathbf{k})$$

$$= B[\mathbf{f}(A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t), \mathbf{k}) - \mathbf{f}(A\mathbf{z}(t), \mathbf{k})] - \dot{\boldsymbol{\varepsilon}}_2(t)$$

$$\approx B \frac{J(\mathbf{f})}{J(\mathbf{y})}(A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t), \mathbf{k})\boldsymbol{\varepsilon}_1(t) - \dot{\boldsymbol{\varepsilon}}_2(t) + o(\|\boldsymbol{\varepsilon}_1(t)\|)$$

(linear approximation at $A\mathbf{z}(t) + \boldsymbol{\varepsilon}_1(t)$, not $A\mathbf{z}(t)$)

$$\dot{\mathbf{z}}(t) - B\mathbf{f}(A\mathbf{z}(t), \mathbf{k}) \approx \mathbf{e}(t)$$

$$\mathbf{e}(t) = B \frac{J(\mathbf{f})}{J(\mathbf{y})}(\mathbf{y}(t), \mathbf{k}) \varepsilon_1(t) - \dot{\varepsilon}_2(t) + o(\|\varepsilon_1(t)\|)$$
(13)

We need to find A and B such that

- $A\mathbf{z}(t) \approx \mathbf{y}(t)$. A is estimated from SVD of simulated data on $\mathbf{y}(t)$: $Y_{N \times n} = Z_{N \times N} \Sigma_{N \times n} (V_{n \times n})^T$, $A = (\Sigma_{p \times p} (V_{p \times n})^T)^T$,
- $\|\mathbf{e}(t)\|$ is as small as possible.
 - We can assume that $\varepsilon_2(t) = 0$
 - Current approach: $B = \text{pinv}(A)$ (Moore-Penrose pseudoinverse). Some row maybe very big.
 - Another option is choosing $B_{N \times p}$ such that $\|B \frac{J(\mathbf{f})}{J(\mathbf{y})}(\mathbf{y}(t), \mathbf{k})\|$ is minimized. Need to make sure that none of the rows of B is very big.