# Supplemental Material: Slowest kinetic modes revealed by metabasin renormalization 

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In this Supplemental Material, we will show that the renormalization procedure developed in the paper can be applicable to the discrete-time kinetic equations, also known as (discrete-time) Markov state models [1], with small modifications.

Suppose that the kinetic state is described by the distribution of probability, $p_{i}$, for $i=1,2, \ldots, n$, where $n$ denotes the number of states, the kinetic equations are given by

$$
\begin{equation*}
p_{i}(t+1)=\sum_{j=1}^{n} t_{i j} p_{j}(t)+p_{i}(t)\left(1-\sum_{j=1}^{n} t_{j i}\right) \quad \text { for } i=1,2, \ldots, n, \tag{1}
\end{equation*}
$$

where $p_{i}(t)$ is the probability distribution of the system state $i$ at discrete times $t=0,1,2, \ldots$, and $t_{i j}$ is the transition probability from state $j$ to state $i$ for $j \neq i$, otherwise $t_{i i}=0$. With the transition probability matrix $T$ defined by $(T)_{i j}=t_{i j}+\delta_{i j}\left(1-\sum_{j^{\prime}=1}^{n} t_{j^{\prime} i}\right)$, the equations can be expressed in a matrix form: $\boldsymbol{p}(t+1)=T \boldsymbol{p}(t)$. We assume the equilibrium, $\lim _{t \rightarrow \infty} \boldsymbol{p}(t)$, to be a unique static state. Accordingly, the eigenvalues of $T$ satisfy $1=\lambda_{0}>\lambda_{1} \geqslant \cdots \geqslant \lambda_{n-1}>0[1]$. The equilibrium $\boldsymbol{p}(\infty)$ coincides with the zeroth eigenvector of $T$, and the first, second, ... eigenvectors of $T$ represent the slowest relaxation modes.

Metabasins (MBs) in transition probability matrices, are determined similarly with the use of monotonic sequences $[2,3]$. A sequence of states $i_{1} \rightarrow i_{2} \rightarrow \ldots$ is called monotonic if it consists only of most probable transitions. The monotonic sequences with the same terminal state belong to the same MB.

Similarly to the transition rate matrix $K$, the columns and rows of $T$ are rearranged in the MB ordering, $\sigma(1,1), \sigma(1,2), \ldots, \sigma(2,1), \sigma(2,2), \ldots$, and the resultant matrix is denoted by $T_{\sigma}$, where $\sigma(\ell, i)$ returns the number $j$ of state $j$ that is the $i$ th energy state in $\mathrm{MB}_{\ell}$. Then, we consider the block diagonal matrix $\operatorname{diag}\left(T_{1}, \ldots, T_{\ell}, \ldots, T_{m}\right)$ to be the unperturbed matrix, where $T_{\ell}$ is given by

$$
\left(T_{\ell}\right)_{i j}=t_{\sigma(\ell, i), \sigma(\ell, j)}+\delta_{i j} \sum_{j^{\prime}=1}^{n_{\ell}}\left(1-t_{\sigma\left(\ell, j^{\prime}\right), \sigma(\ell, i)}\right),
$$

where $n_{\ell}$ is the size of $\mathrm{MB}_{\ell}$. Note that $j$ th eigenvalues, $\lambda_{\ell, j}$, of $T_{\ell}$ satisfy $1=\lambda_{\ell, 0}>\lambda_{\ell, 1} \geqslant \cdots \geqslant \lambda_{\ell, n_{\ell}-1}>0$. To consider the intra-MB relaxation modes, we form $\Lambda_{\ell}=D_{\ell}^{-1} T_{\ell} D_{\ell}$, with the use of the diagonal matrix $D_{\ell}=$ $\operatorname{diag}\left(\sqrt{\boldsymbol{p}_{\ell, 0}}\right)$, where $\boldsymbol{p}_{\ell, 0}$ is the local equilibrium in $\mathrm{MB}_{\ell}$. It is noteworthy that $\Lambda_{\ell}$ is the symmetric matrix and can be diagonalized with an orthogonal matrix $S_{\ell}=\left[\sqrt{\boldsymbol{p}_{\ell, 0}}, \boldsymbol{v}_{\ell, 1}, \ldots, \boldsymbol{v}_{\ell, n_{\ell}-1}\right]$, where $\boldsymbol{v}_{\ell, j}$ is the $j$ th eigenvectors of $\Lambda_{\ell}$.

To consider the inter-MB transitions, we form the symmetric matrix $\Lambda^{\prime}=S^{T} D^{-1} T_{\sigma} D S$, with $S=\operatorname{diag}\left(S_{1}, S_{2}, \ldots\right)$ and $D=\operatorname{diag}\left(\sqrt{\boldsymbol{p}_{\text {eq }}}\right)$, where $\boldsymbol{p}_{\text {eq }}$ is the equilibrium of $T_{\sigma}$. Moreover, to introduce the division of intra-MB relaxation modes into slow and fast modes, we set a certain threshold $\lambda_{\text {cut }}$ satisfying $1 \geqslant \lambda_{\text {cut }}>0$ : the slow relaxation modes are $1 \geqslant \lambda_{\ell, j} \geqslant \lambda_{\text {cut }}$ and the fast relaxation modes are $\lambda_{\text {cut }}>\lambda_{\ell, j} \geqslant 0$. We, then, reorder the columns and lows of $\Lambda^{\prime}$ in the slow-to-fast relaxation block order, and the resultant matrix is denoted by $\Lambda_{\text {slow-fast }}$. Note that $\Lambda_{\text {slow }}$, defined by the first $n_{\text {slow }} \times n_{\text {slow }}$ submatrix with $n_{\text {slow }}$ denoting the number of unperturbed slow relaxation modes, generally describes an approximate transitions between the slow modes. To obtain the accurate results, we need a renormalized transition matrix $\Lambda_{\text {slow }}^{\mathrm{RG}}$. To this end, we use a Jacobi rotation $\Lambda_{\text {slow-fast }} \mapsto \Lambda_{\text {slow-fast }}^{\mathrm{RG}}=G^{T} \Lambda_{\text {slow-fast }} G$ such that the resultant couplings between slow and fast modes, $\left(\Lambda_{\text {slow-fast }}^{\mathrm{RG}}\right)_{i j}$ with $i \leqslant n_{\text {slow }}<j$, are vanishing. Now, the renormalized transition matrix $\Lambda_{\text {slow }}^{\mathrm{RG}}$ is defined by the first $n_{\text {slow }}$-by- $n_{\text {slow }}$ submatrix of $\Lambda_{\text {slow-fast }}^{\mathrm{RG}}$, which reproduces the exact slowest relaxations of the kinetic equation (1).

Table I summarizes the differences between the renormalization procedures for discrete-time and for continuous-time kinetic equations, from which we clearly see that the renormalization procedure for discrete-time kinetic equations is constructed almost in the same way as that for continuous-time kinetic equations developed in the paper.

TABLE I. Differences between discrete-time and continuous-time renormalization procedures: The listed differences are absorbed by the matrix $\Lambda^{\prime}$ and the definitions of slow modes. In other words, all the procedures after specifying these are exactly the same way between for the discrete-time equation and for the continuous-time equation.

| Time | Matrix to be normalized | $\Lambda^{\prime}$ | Slow modes |
| :--- | :--- | :--- | :--- |
| Discrete time | Transition probability matrix $T$ | $S^{T} D^{-1} T_{\sigma} D S$ | $1 \geqslant \lambda_{\ell, i} \geqslant \lambda_{\text {cut }}$ |
| Continuous time | Transition rate matrix $K$ | $S^{T} D^{-1} K_{\sigma} D S$ | $0 \geqslant \lambda_{\ell, i} \geqslant \lambda_{\text {cut }}$ |

[1] An Introduction to Markov State Models and Their Application to Long Times cale Molecular Simulation, edited by G. R. Bowman, V. S. Pande, and F. Noé (Springer, New York, 2013).
[2] For another formularization for MB decompositions of discrete-time Markov state models, see K. Klemm, C. Flamm, and P. F. Stadler, Eur. Phys. J. B 63, 387 (2008).
[3] T. Okushima, T. Niiyama, K. S. Ikeda, and Y. Shimizu, Phys. Rev. E 80, 036112 (2009).

