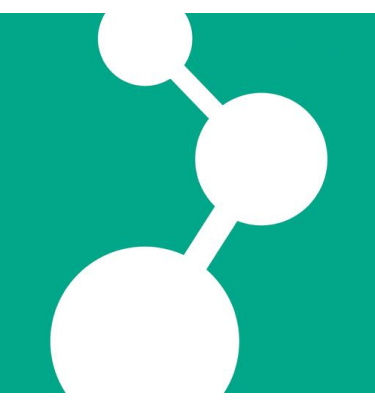




Big Data Summer, 11 Sept 2019



Variational autoencoders for dimensionality reduction and clustering of molecular dynamics data

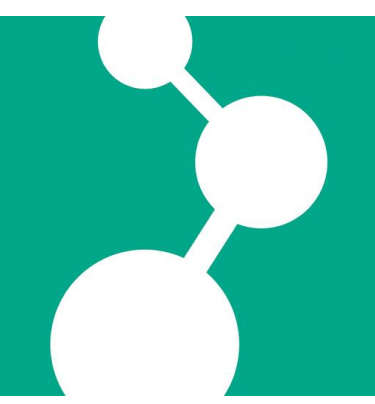
Jose

Max Planck Institute for

+ Automated detection of many-particle solvation states using hidden Markov models



Big Data Summer, 11 Sept 2019



Variational autoencoders for dimensionality reduction and clustering of molecular dynamics data

Joseph F. Rudzinski

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Prof. Kurt Kremer



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Dr. Yani Zhao



Funding

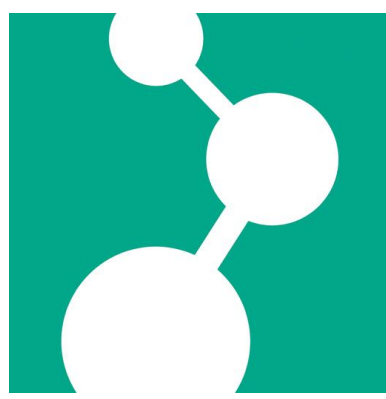
Humboldt postdoctoral fellowship



Alexander von Humboldt
Stiftung/Foundation

DFG Deutsche
Forschungsgemeinschaft



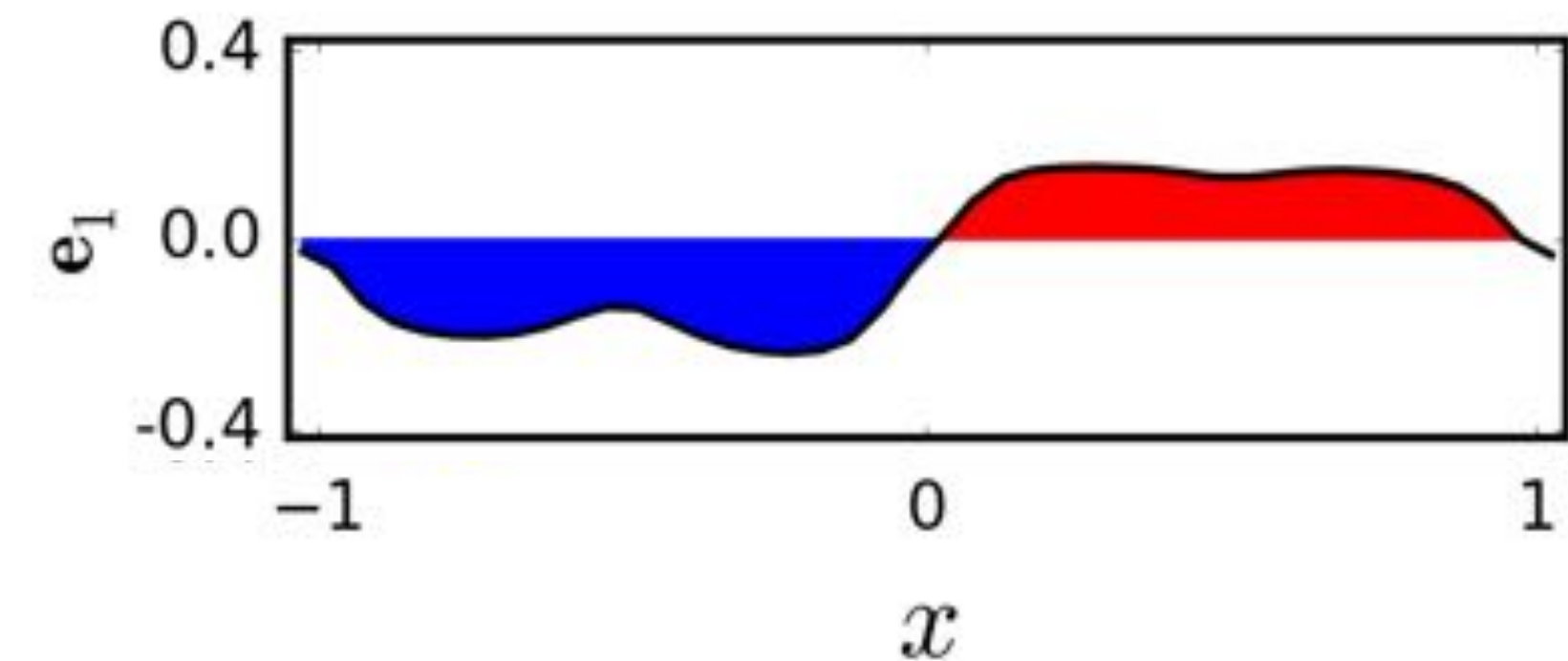
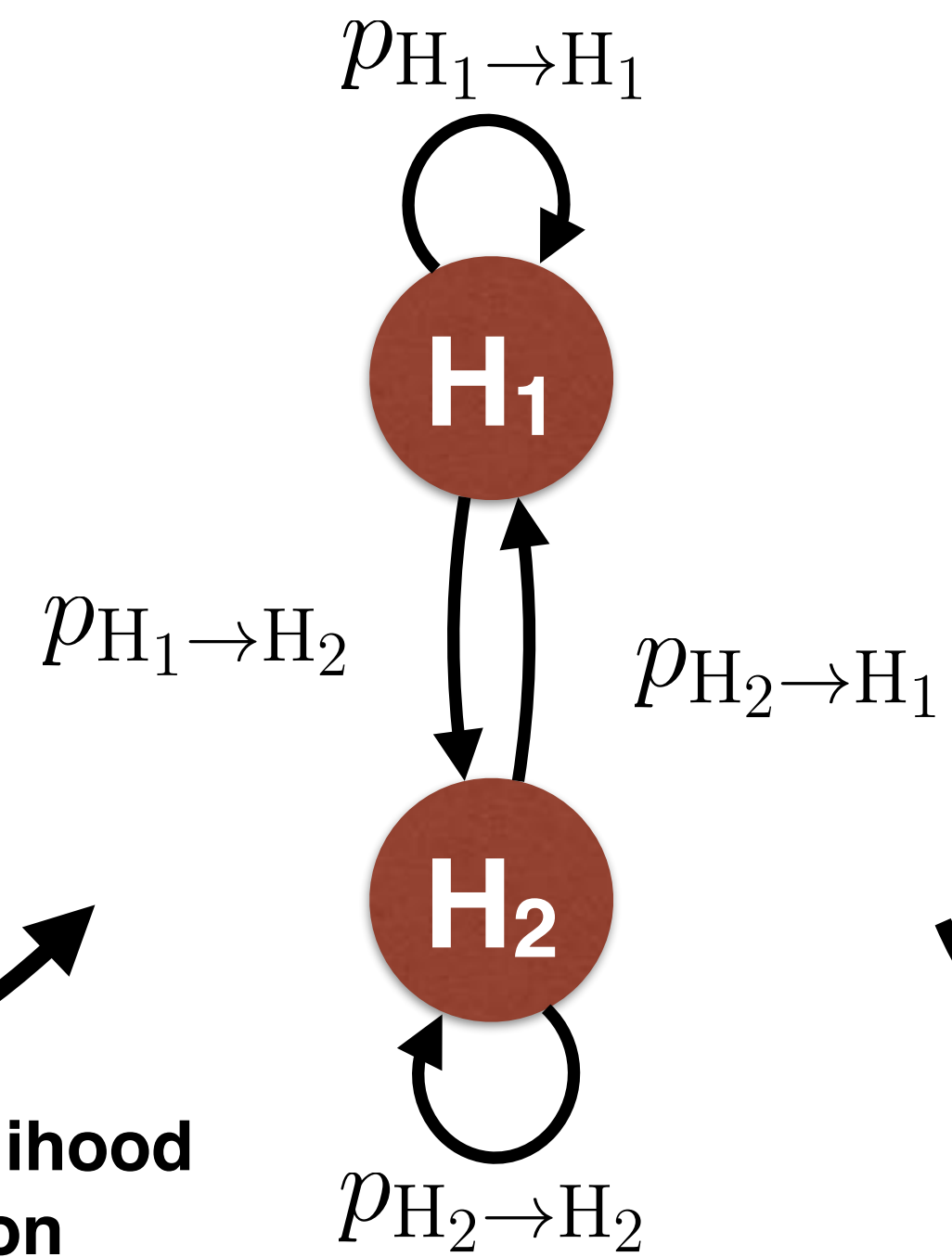
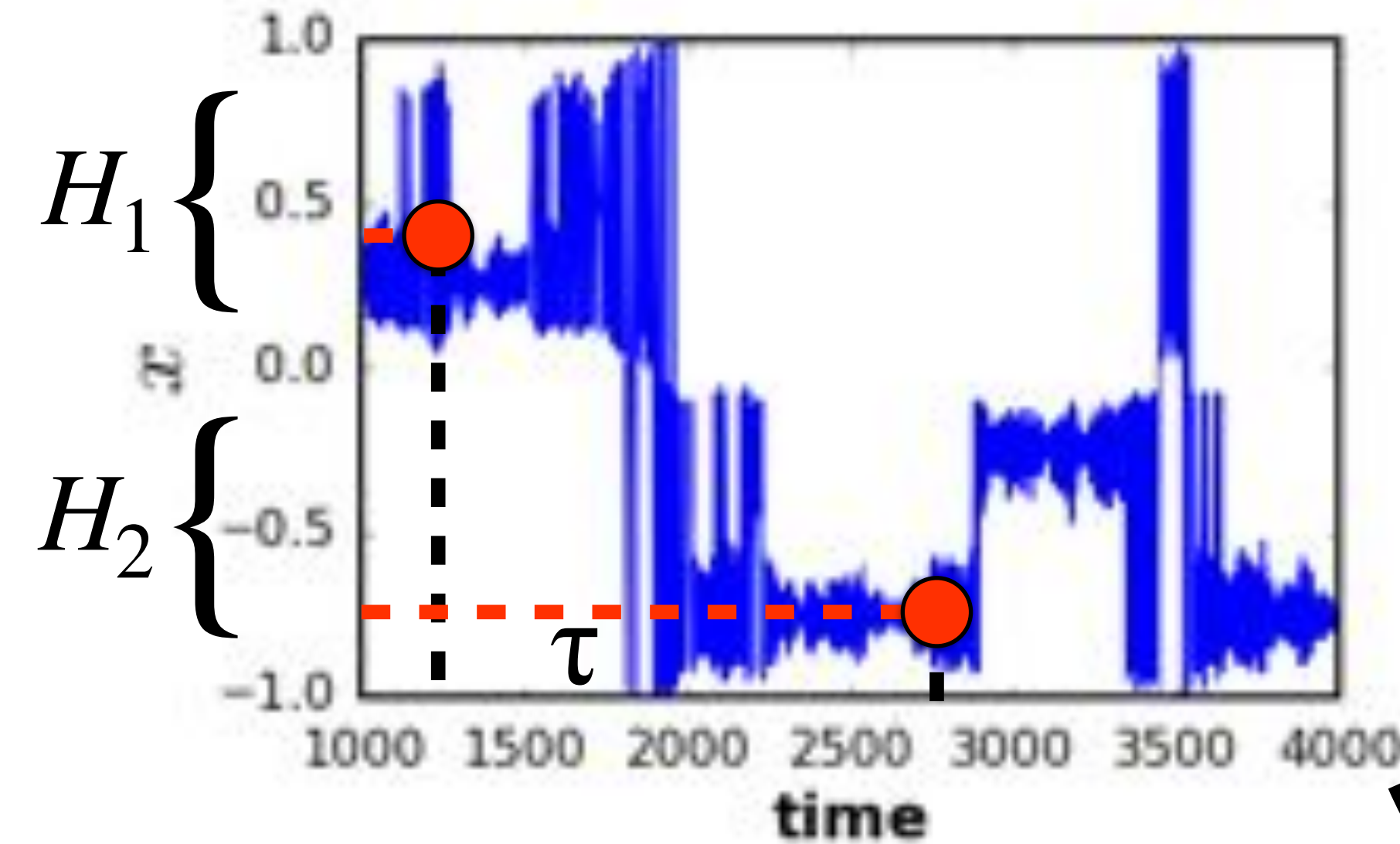


Markov State Models

micro-trajectory

transition probabilities

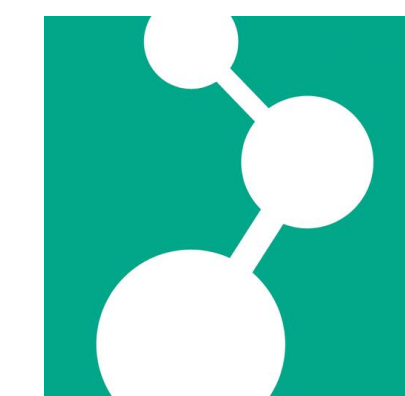
hierarchy of kinetic processes



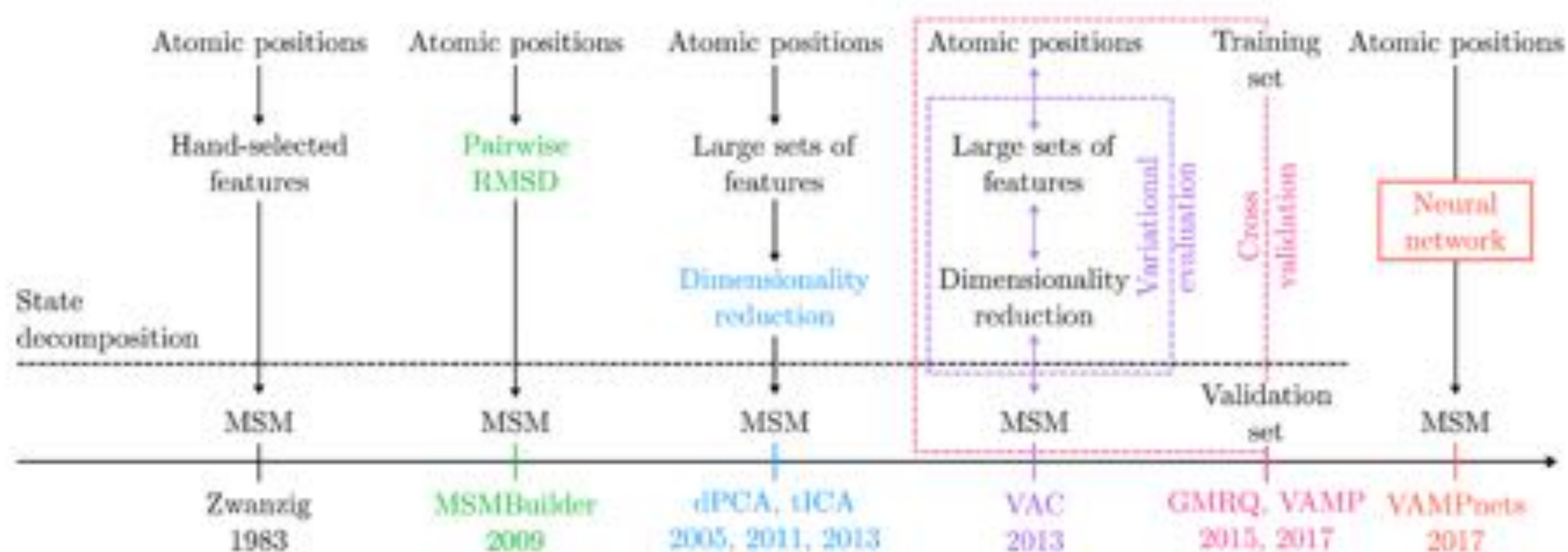
maximum likelihood optimization

diagonalize

MSMs link microtrajectories with long timescale kinetic processes

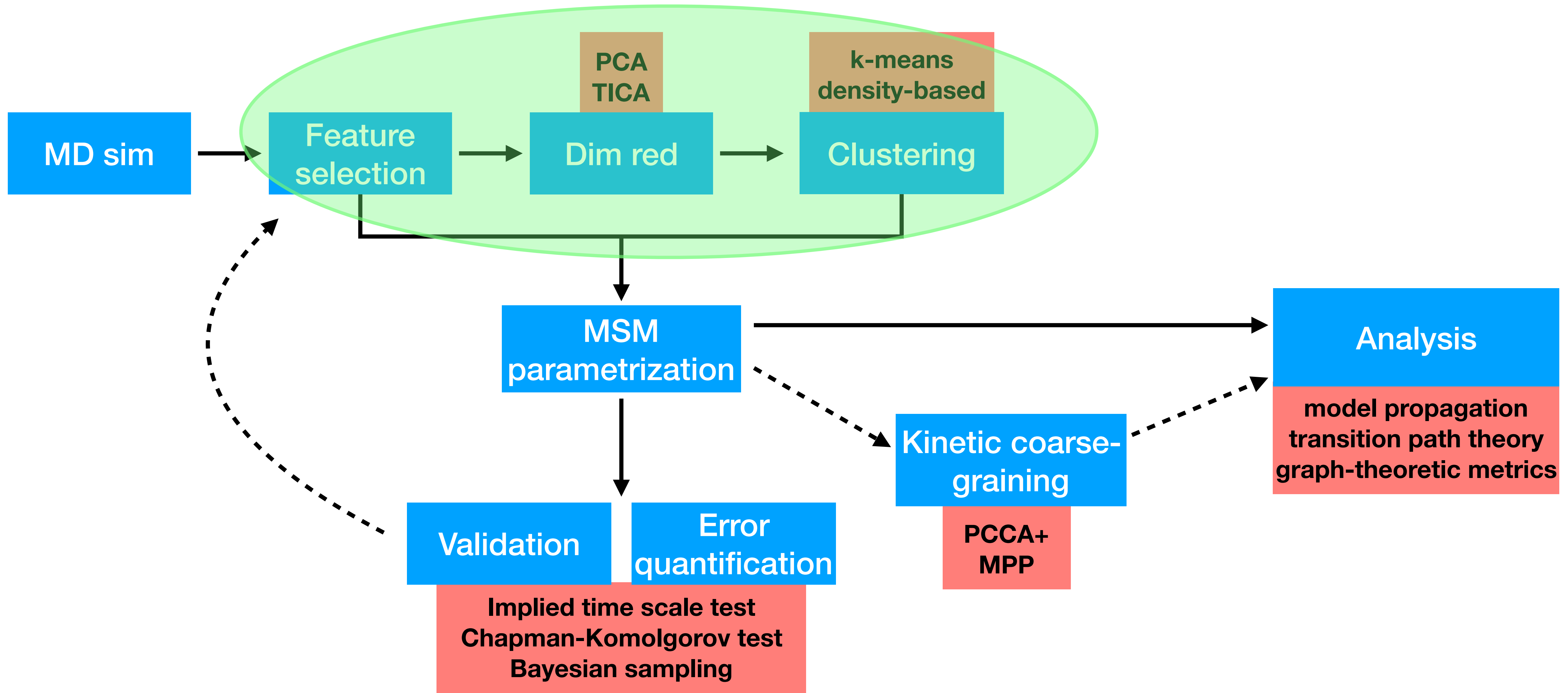
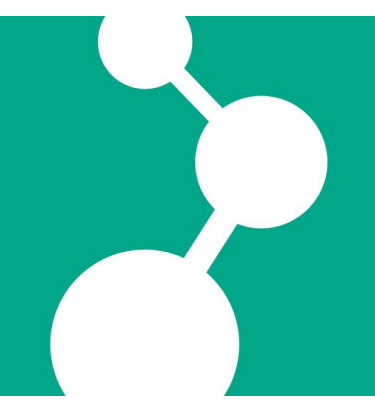


Analysis: Markov state models (MSMs)



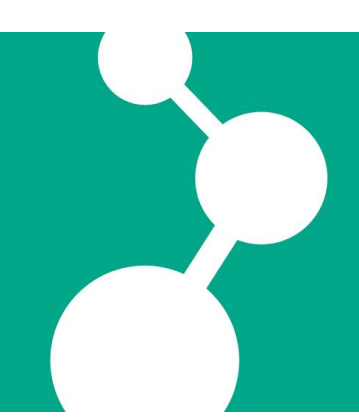


Analysis: Markov state models (MSMs)

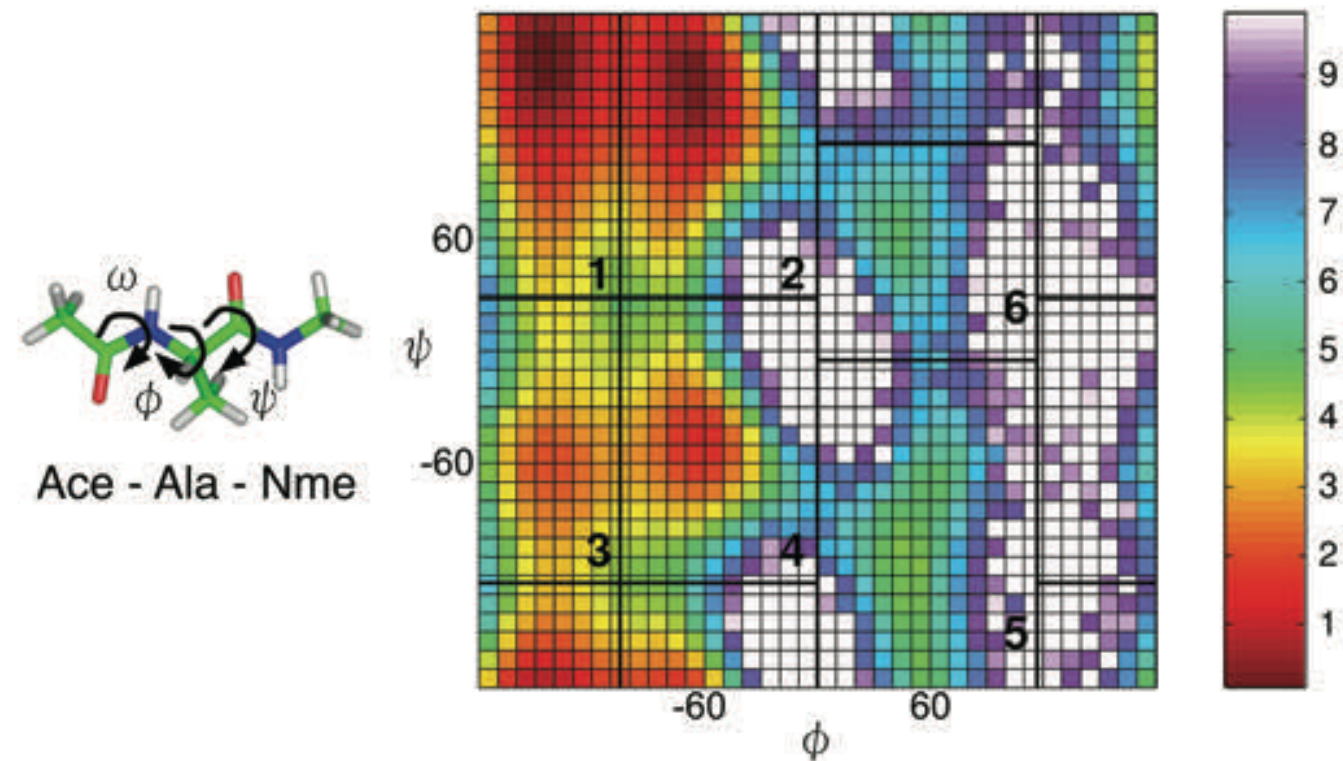




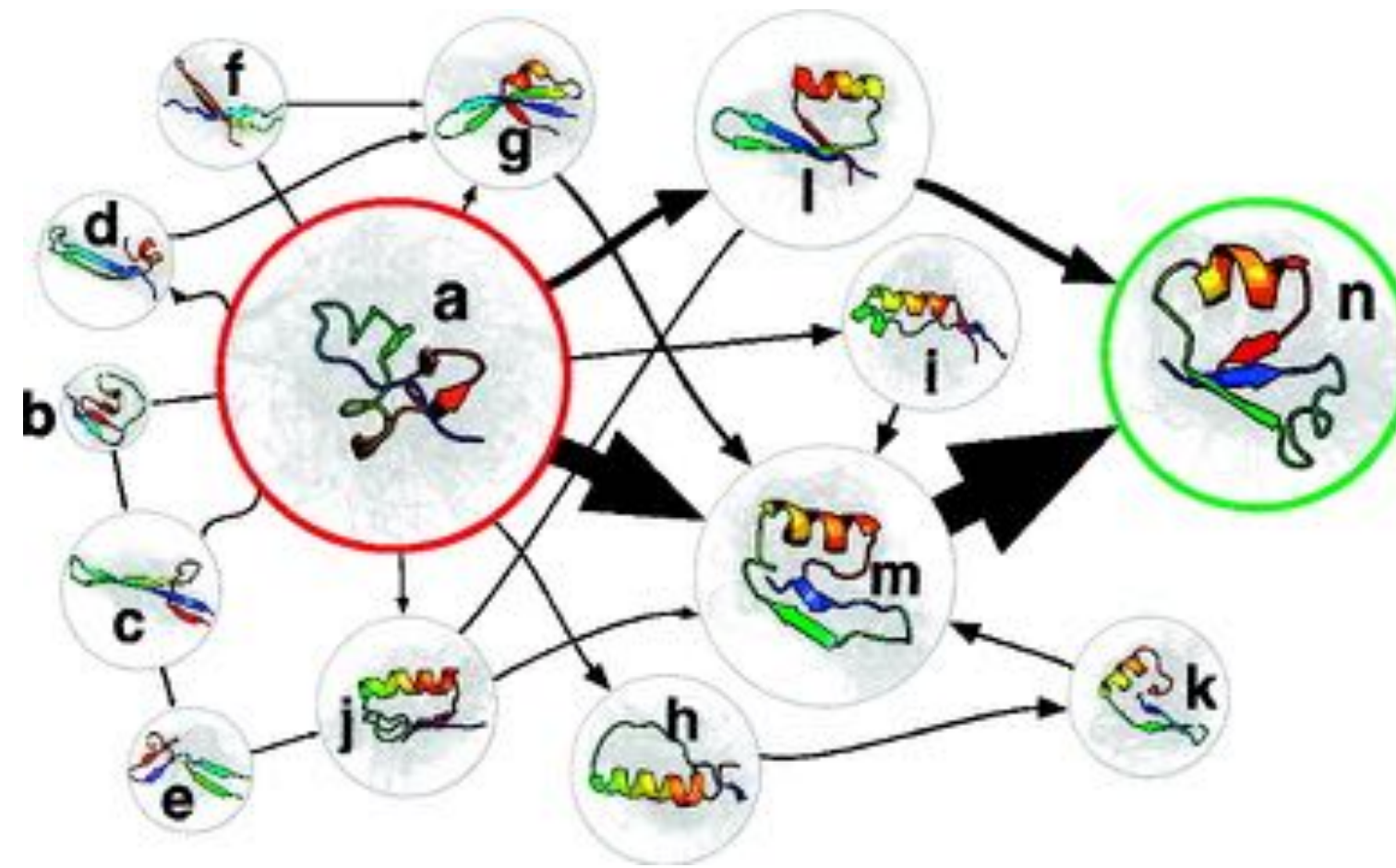
Success of MSMs for protein systems



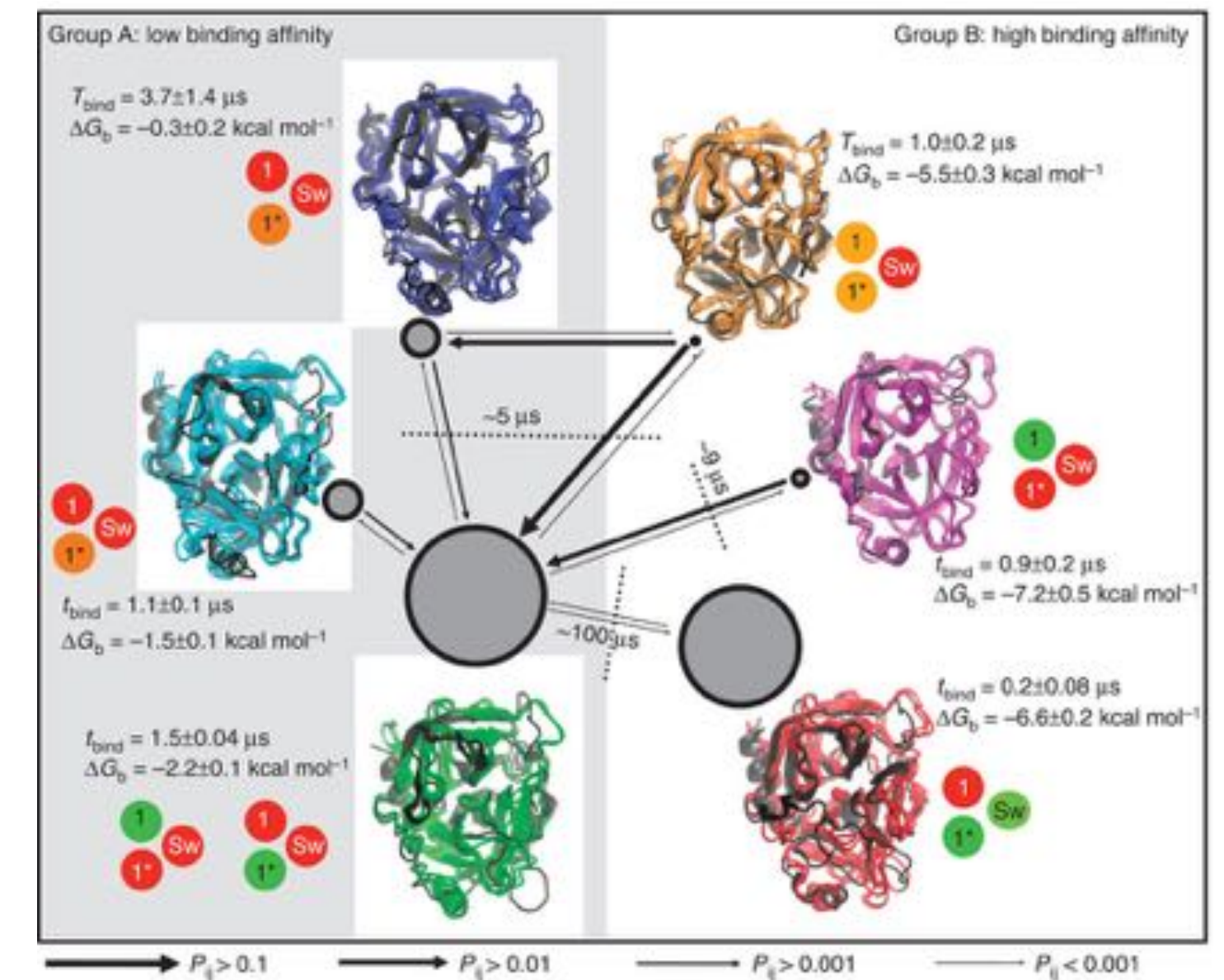
Well-characterized, low-dimensional features provide a great starting point for building kinetic models.



Chodera *et al.* *JCP* (2007)



Voelz *et al.* *JACS* (2010)



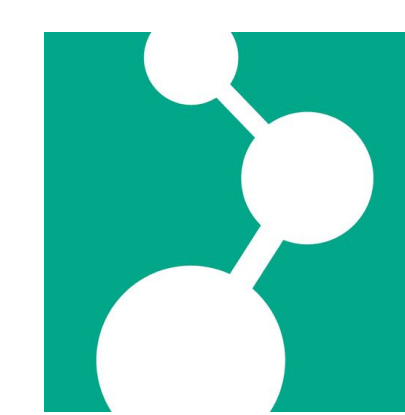
Plattner *et al.* *Nat Comm* (2015)

Pande and coworkers
Noé and coworkers
many more...

Recent Review - Husic and Pande "Markov state models: from an art to a science" *JACS* (2018)



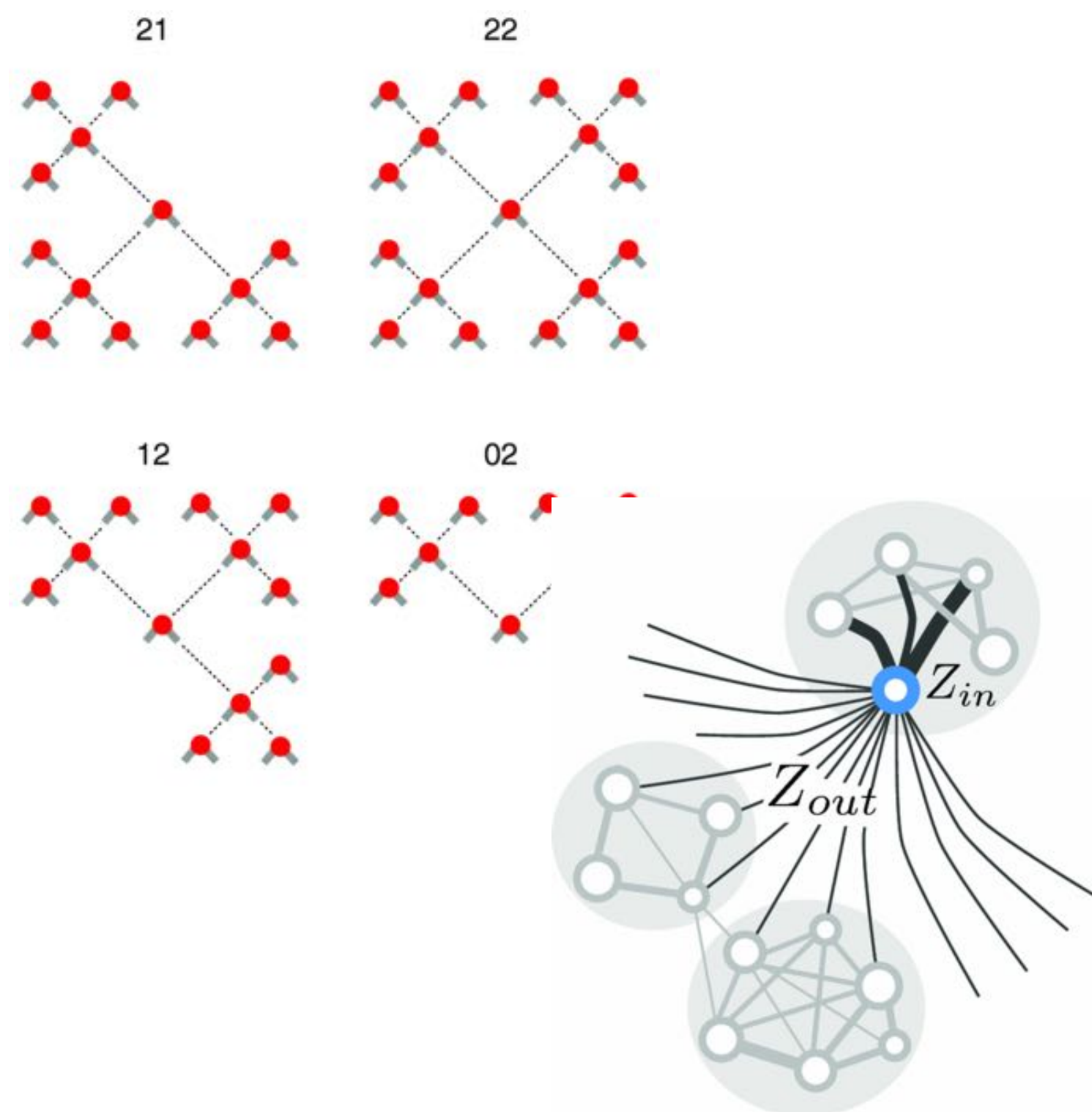
MSMs for many-particle systems



Specialized, system-dependent input features

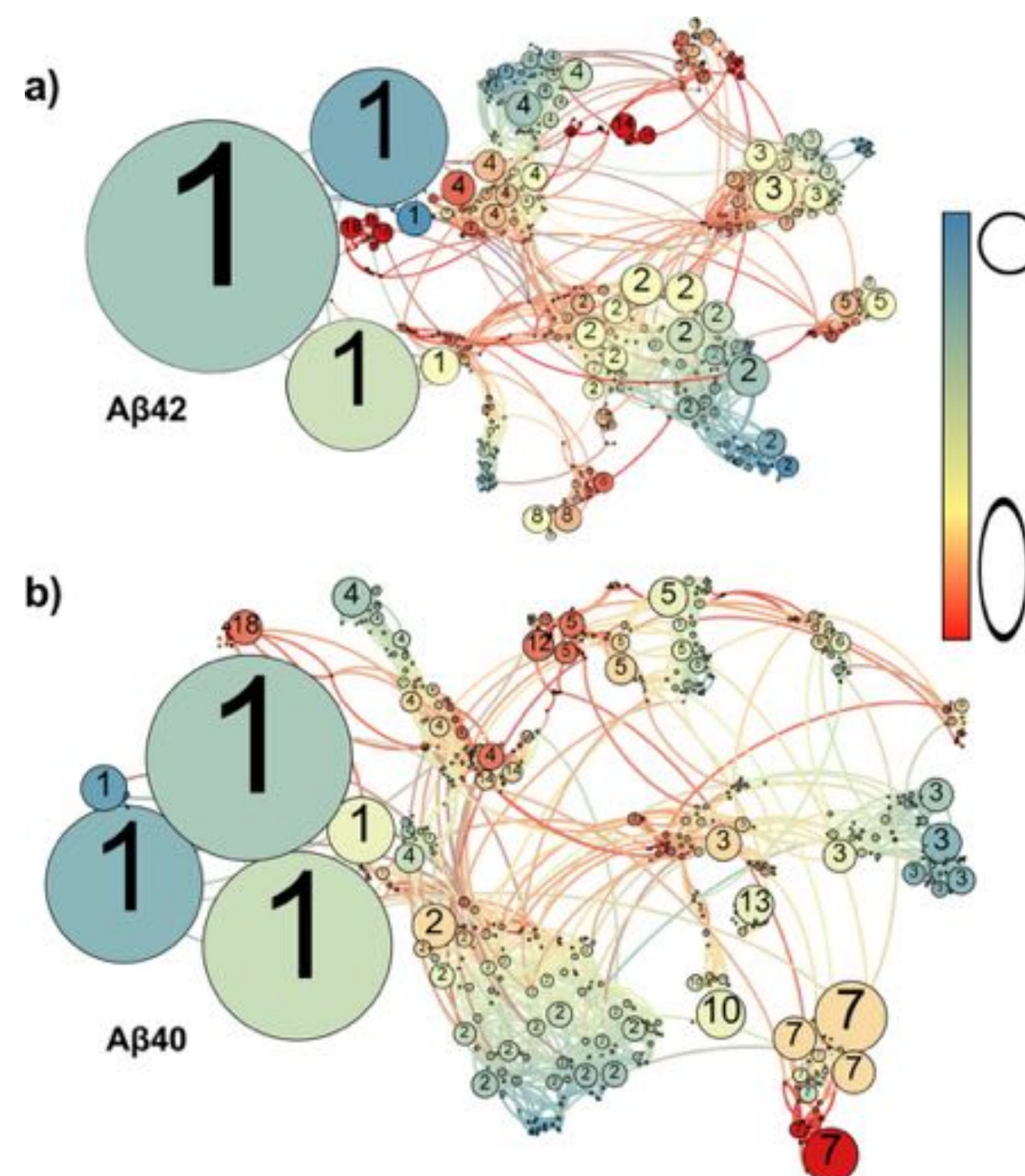
Hidden-Markov-model-based identification of states

Water hydrogen bonding dynamics



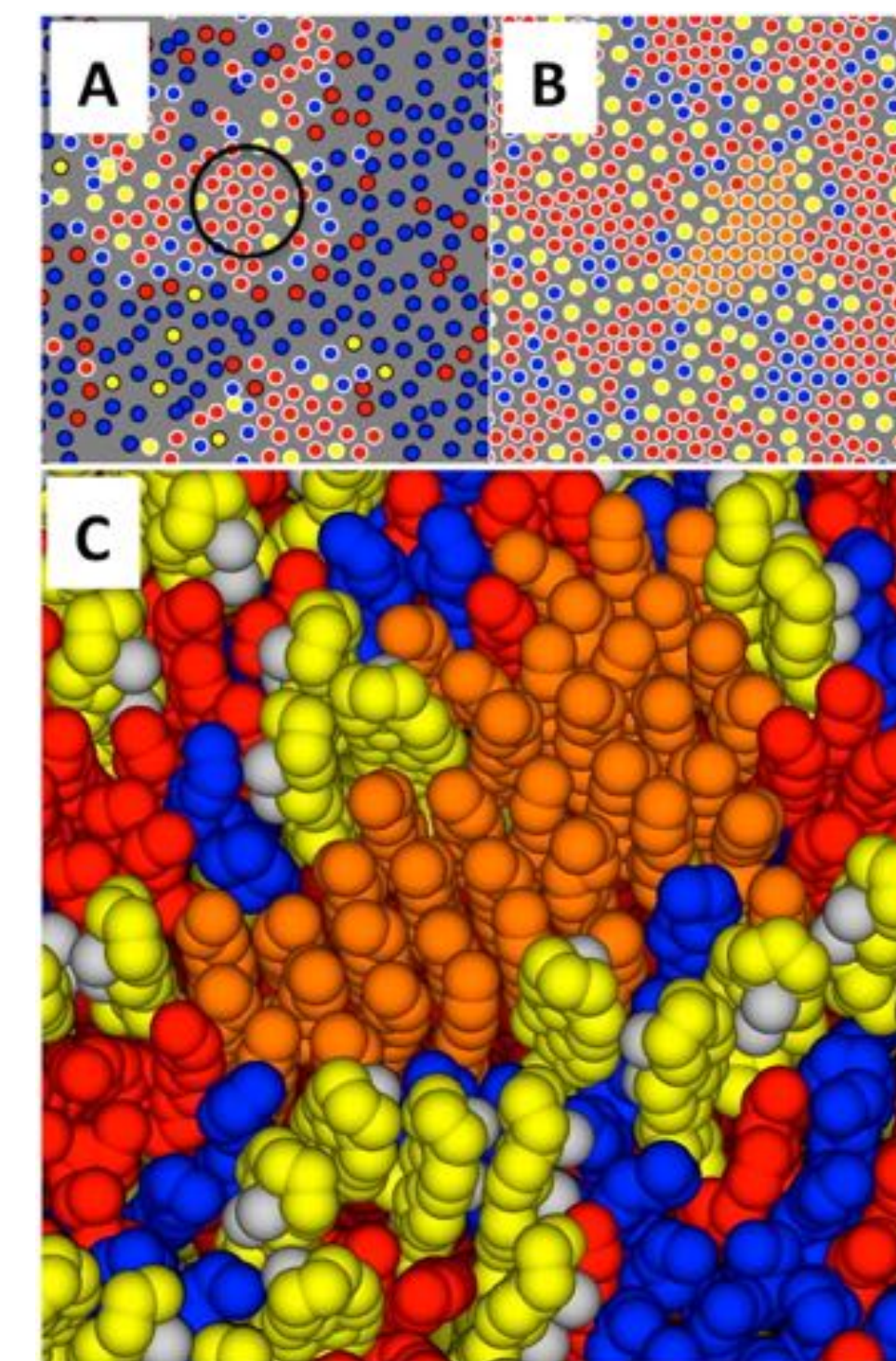
Prada-Garcia *et al.* *JCP* (2012)

Amyloid aggregation kinetics



Barz, Liao, Strodel *JACS* (2018)

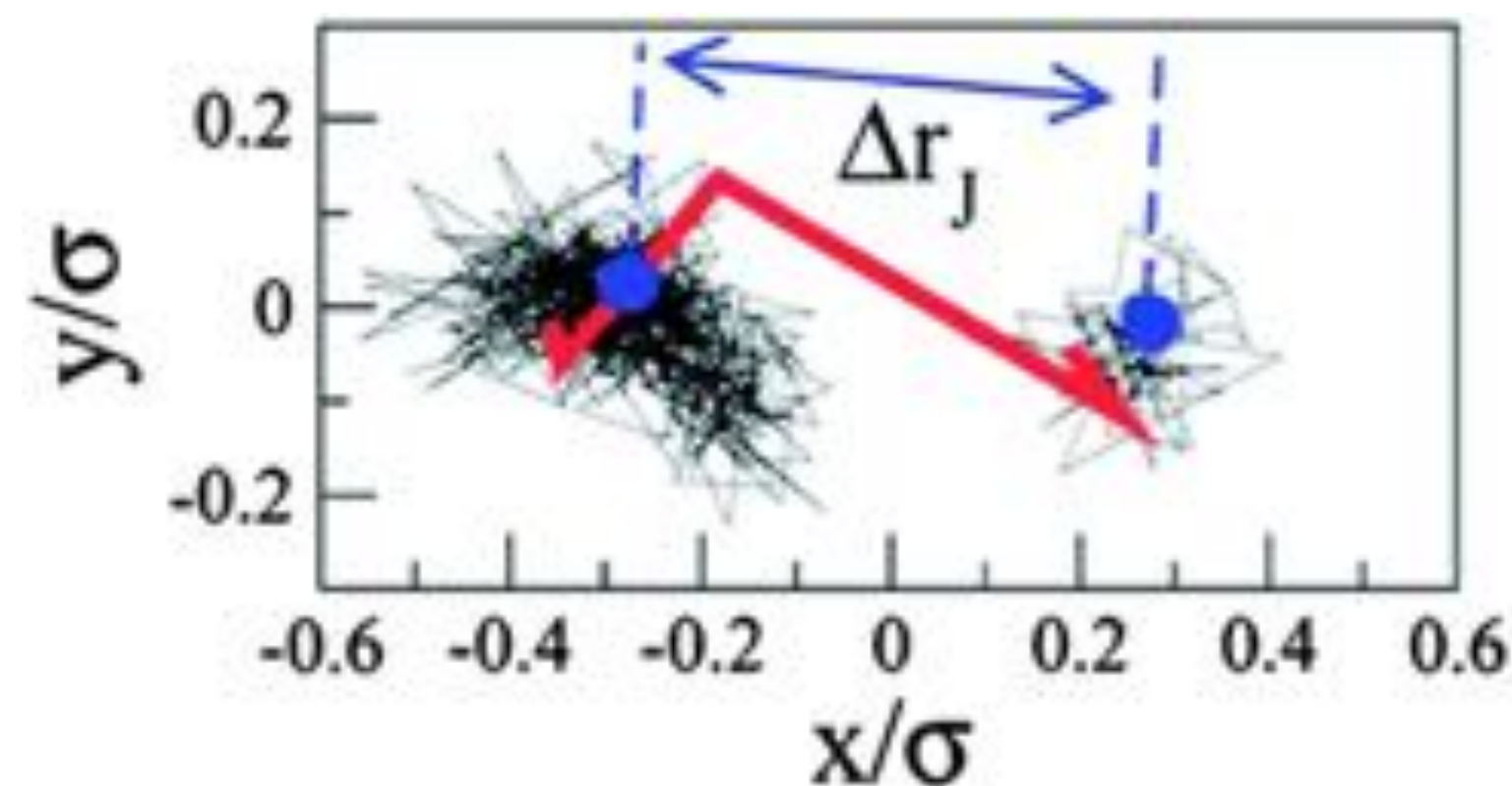
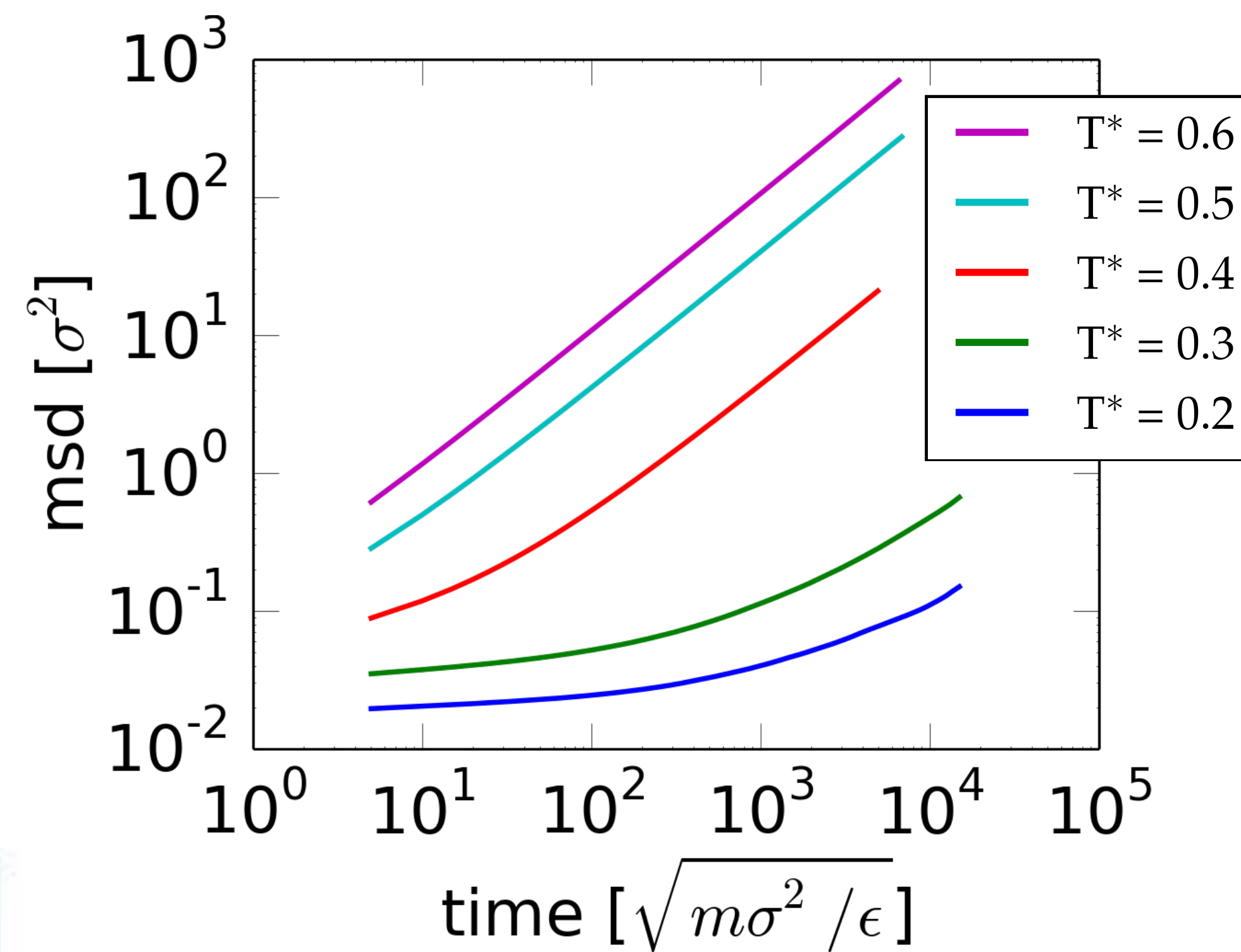
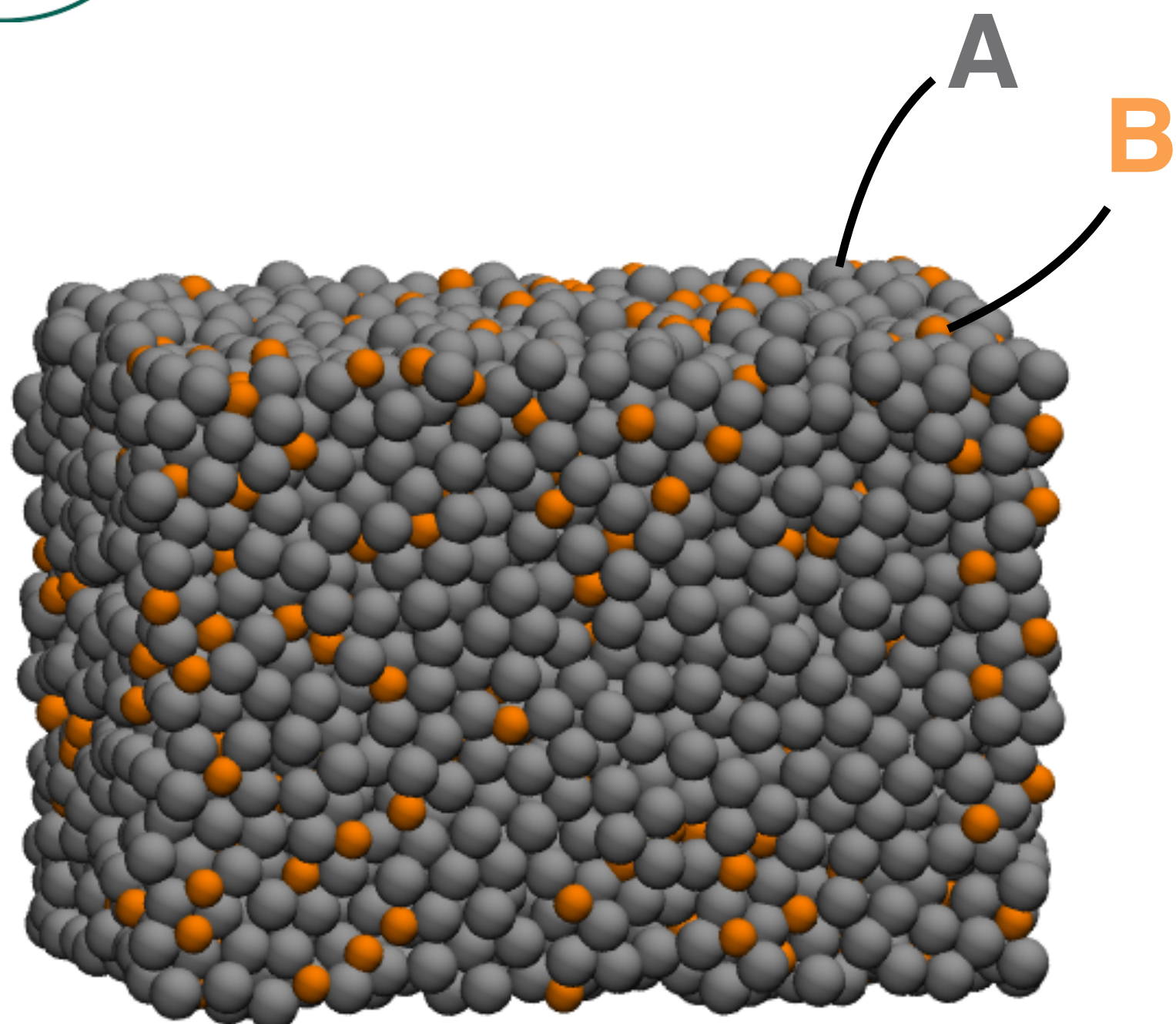
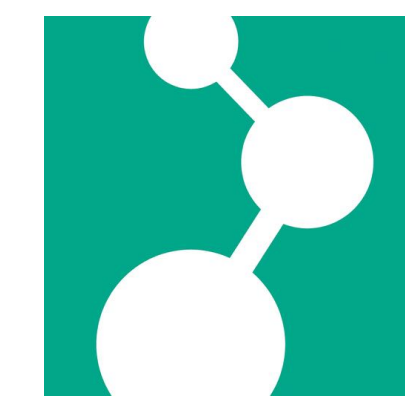
Lipid ordering in bilayers



Sodt *et al.* *JACS* (2014)

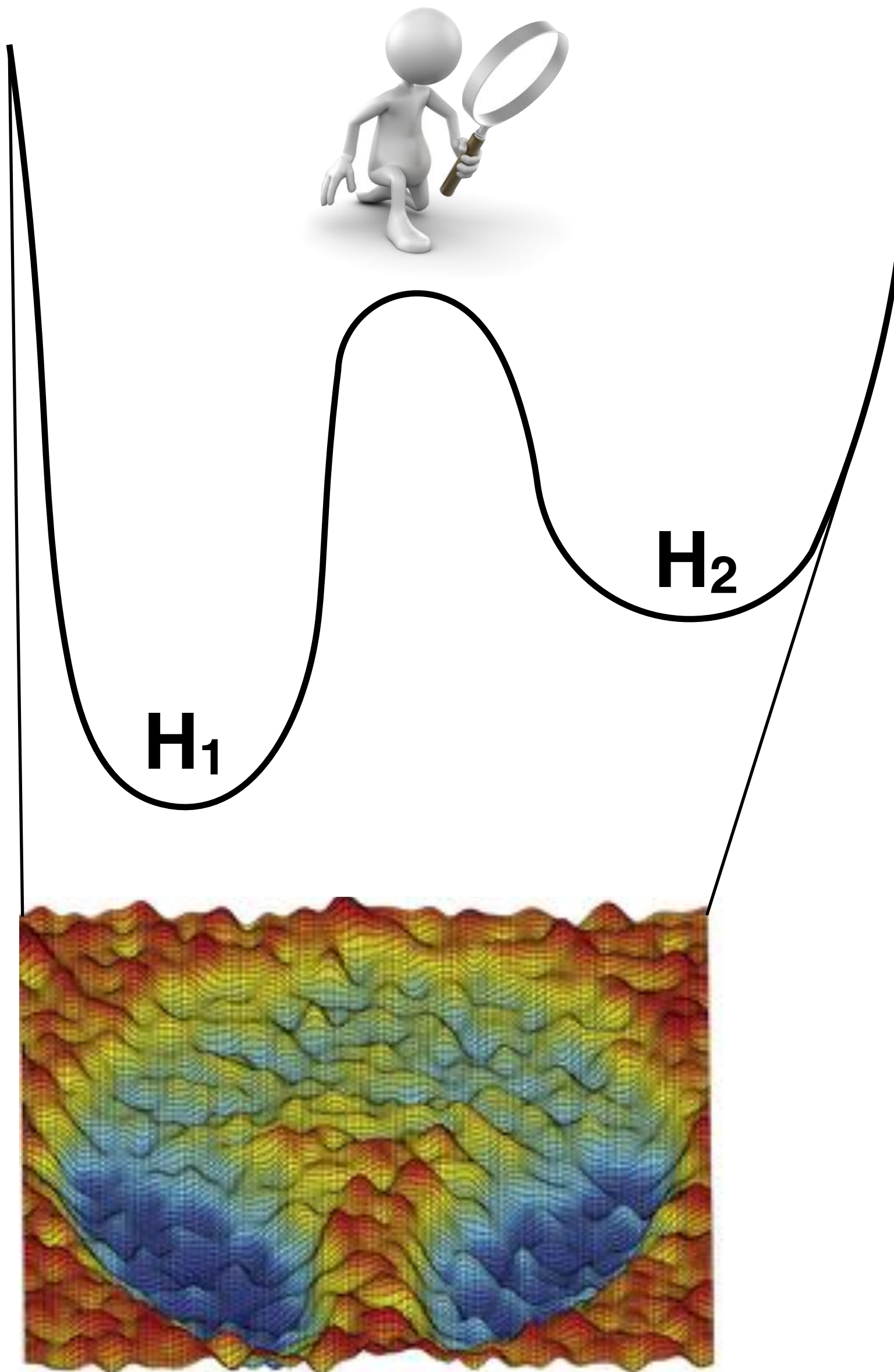
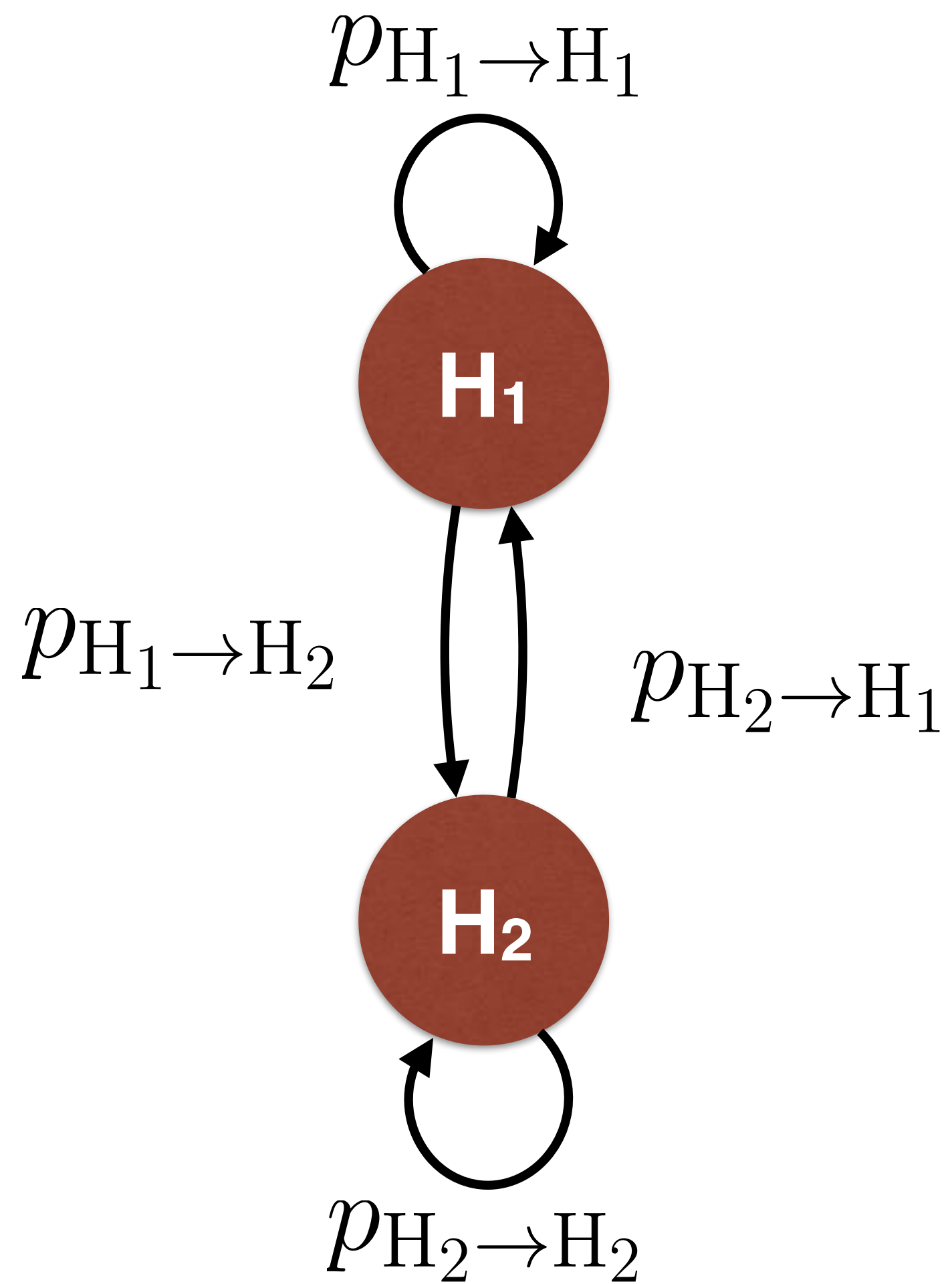
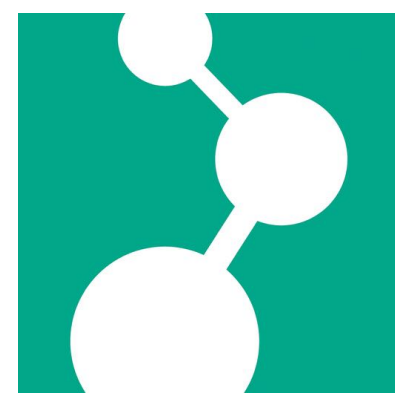


Kob-Andersen model for glassy liquids



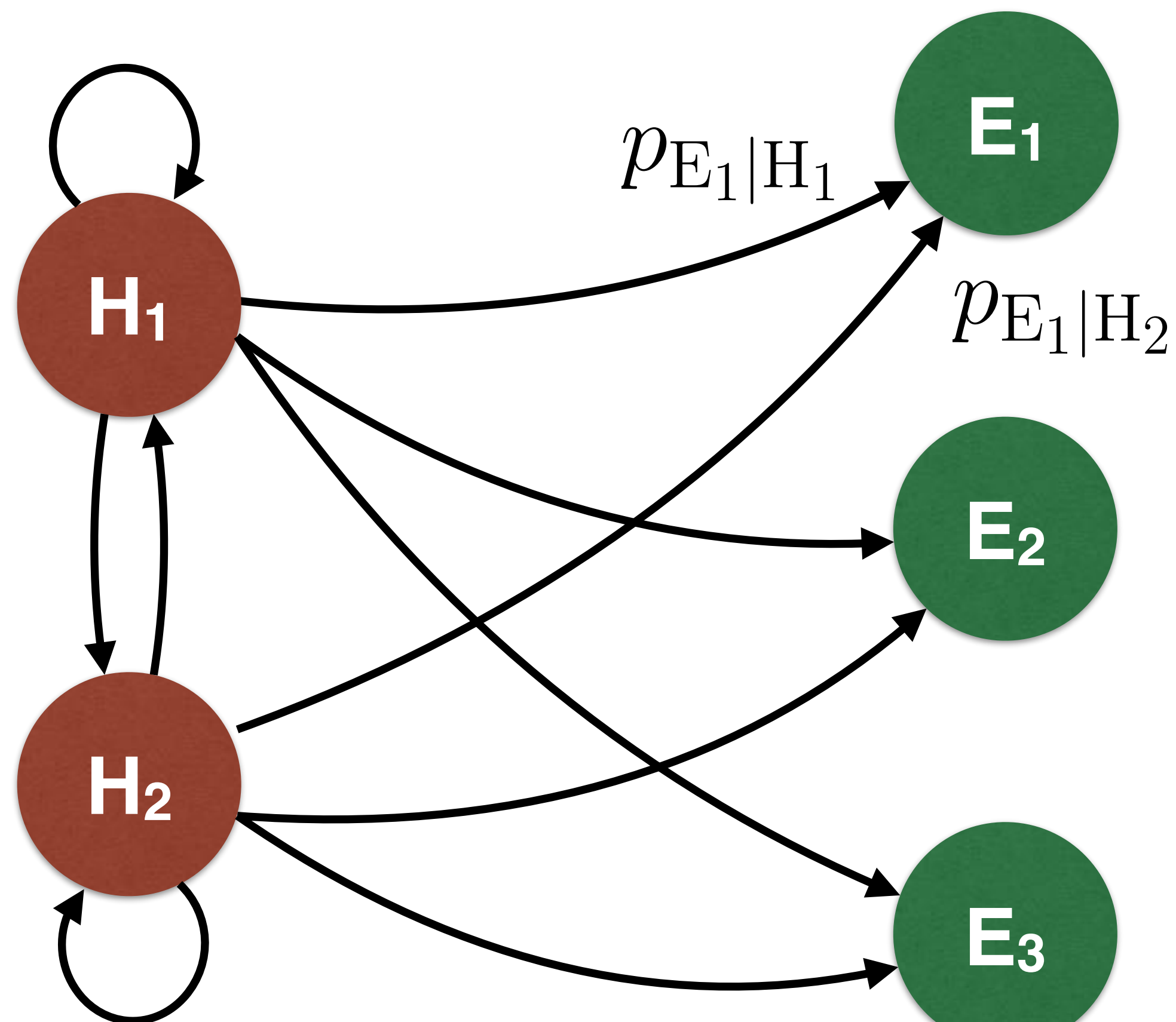
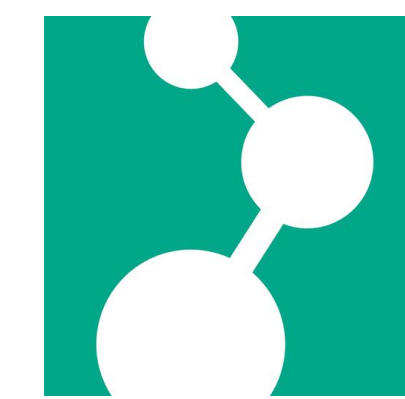


MSMs on observable states

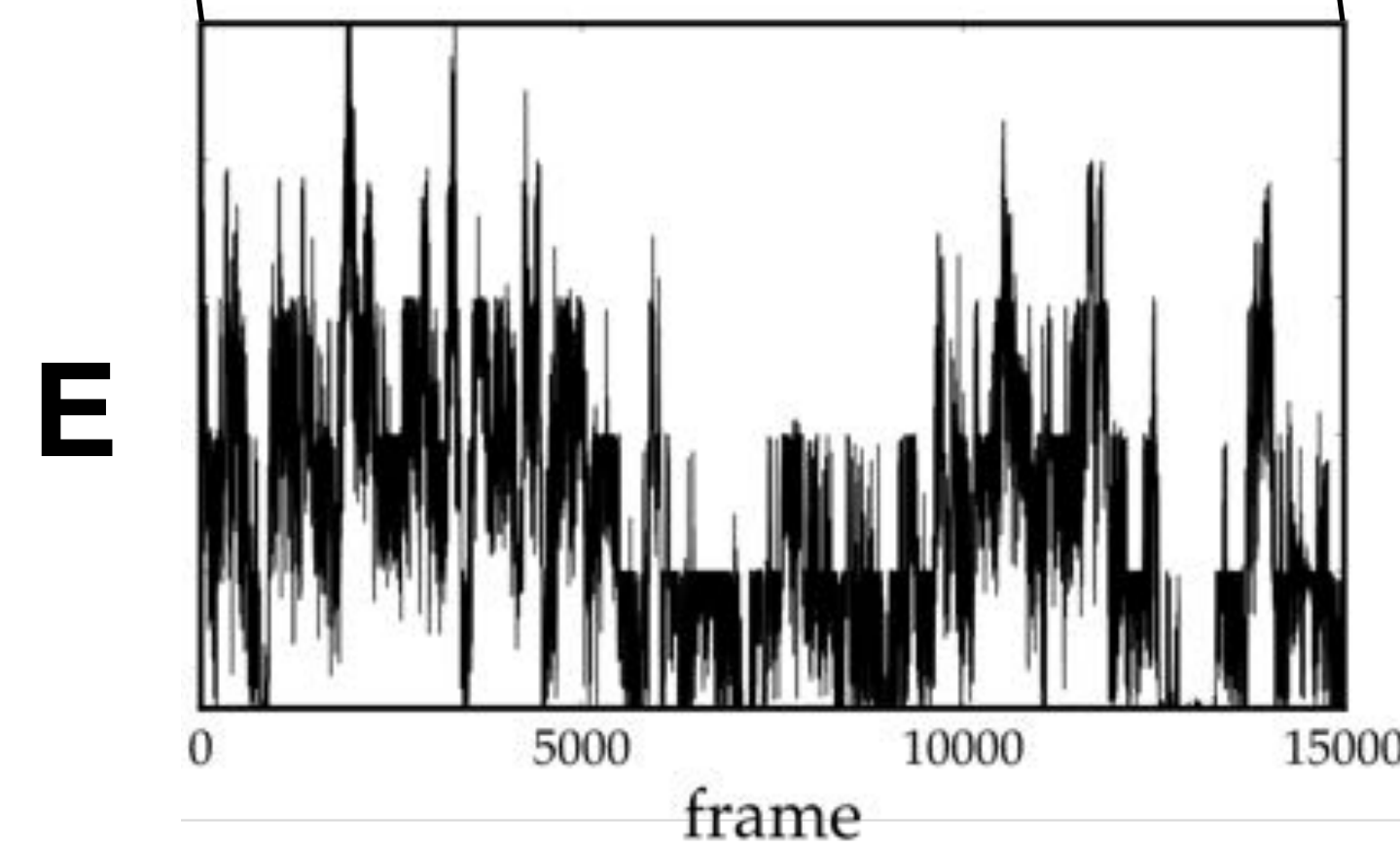
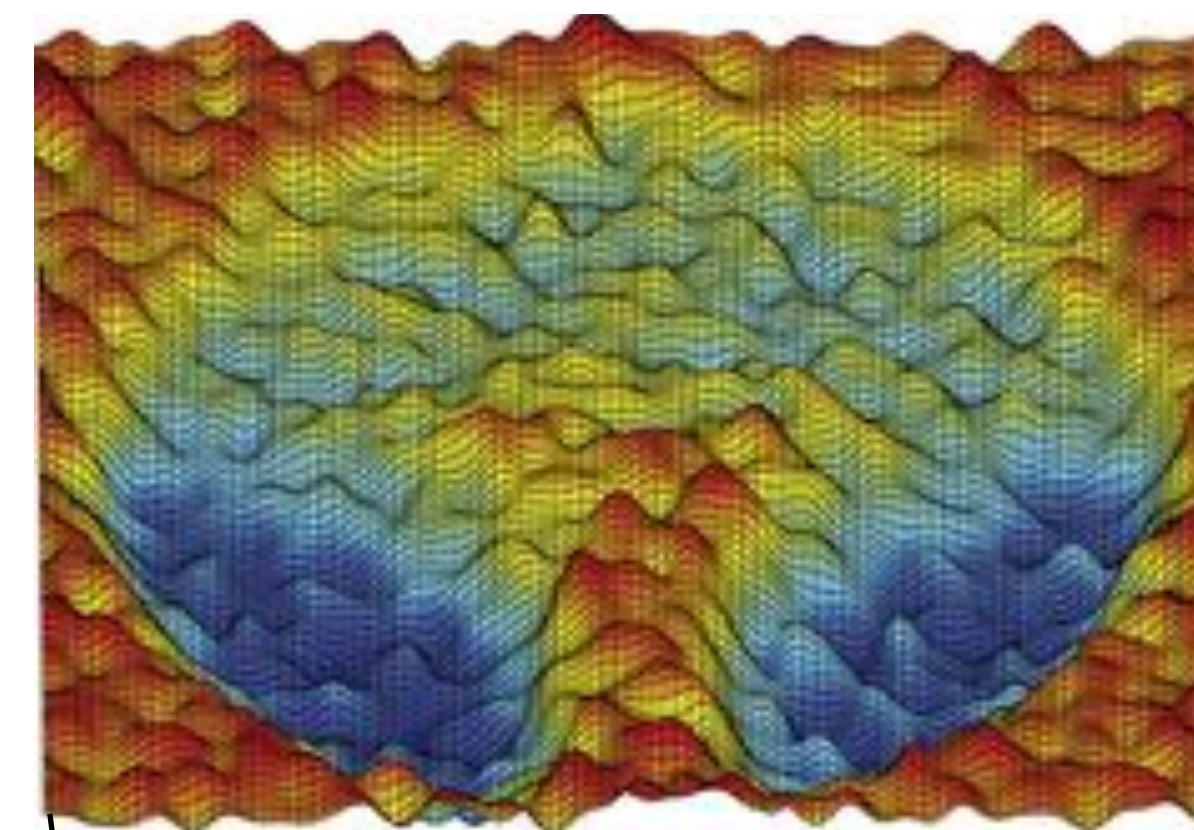




MSMs on hidden states

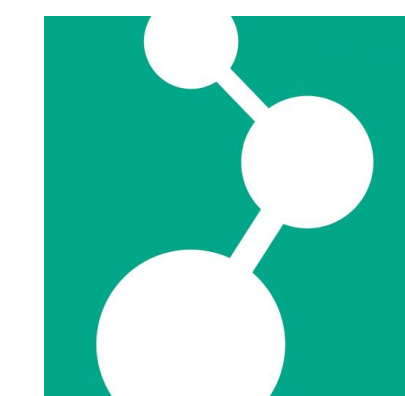


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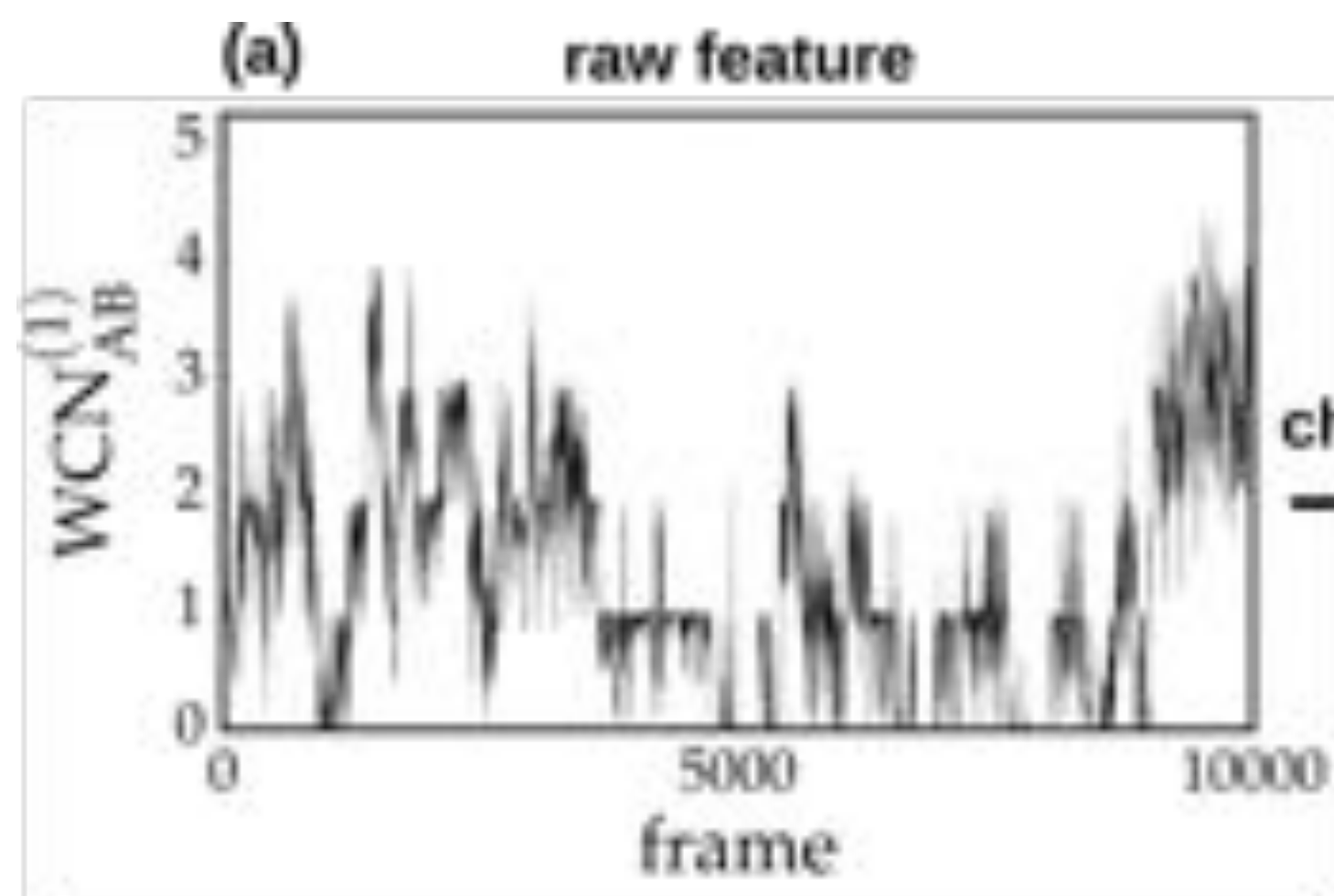
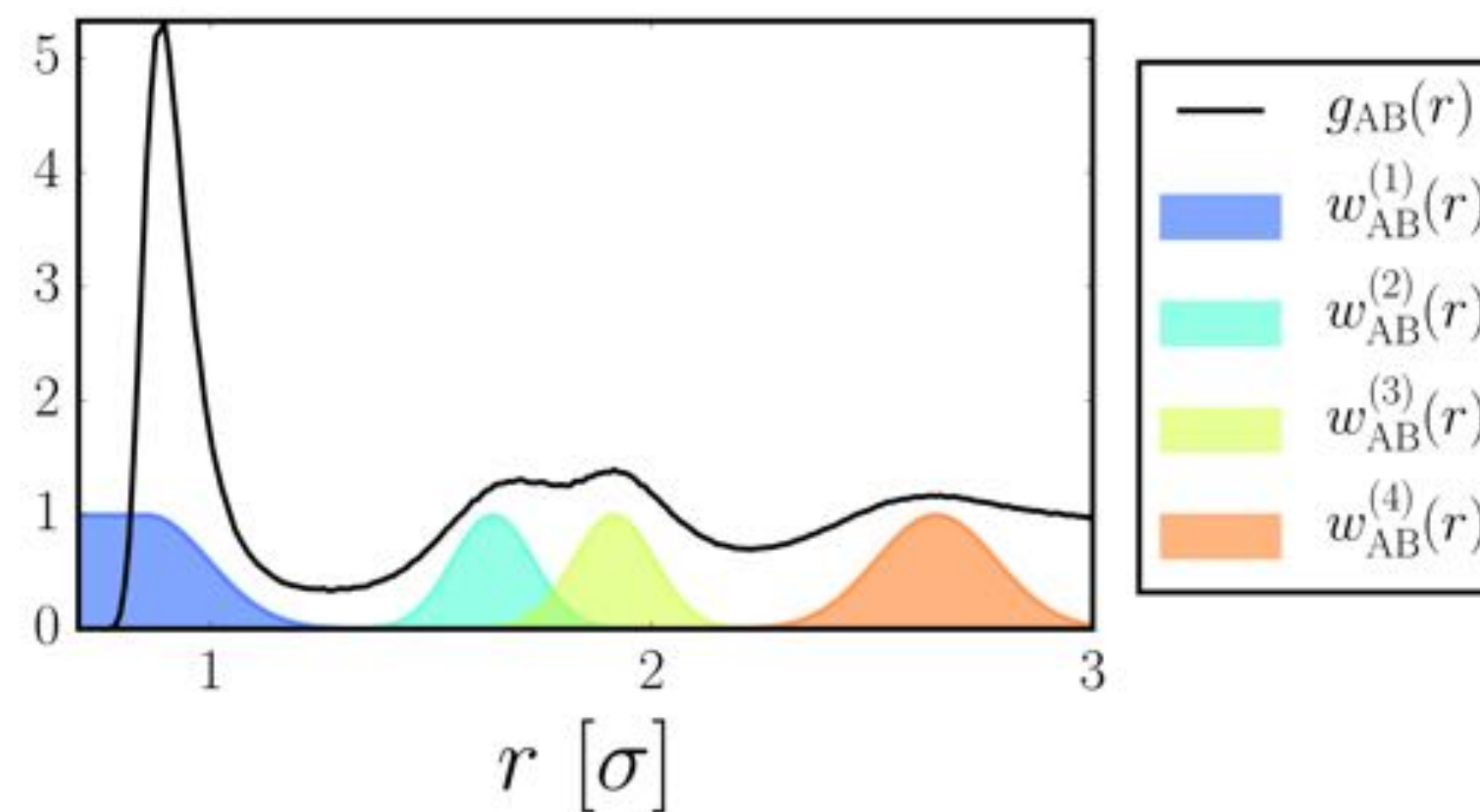
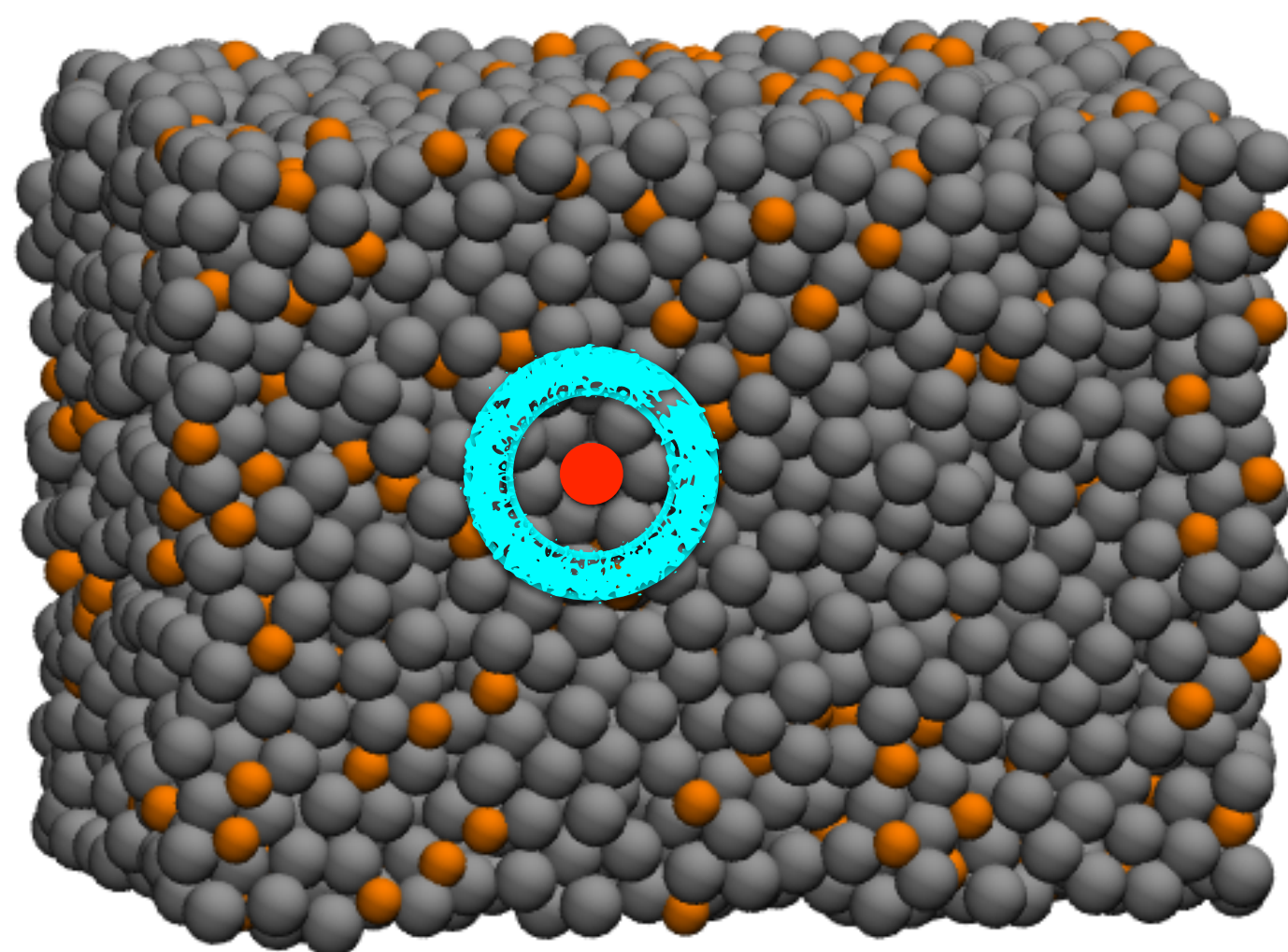




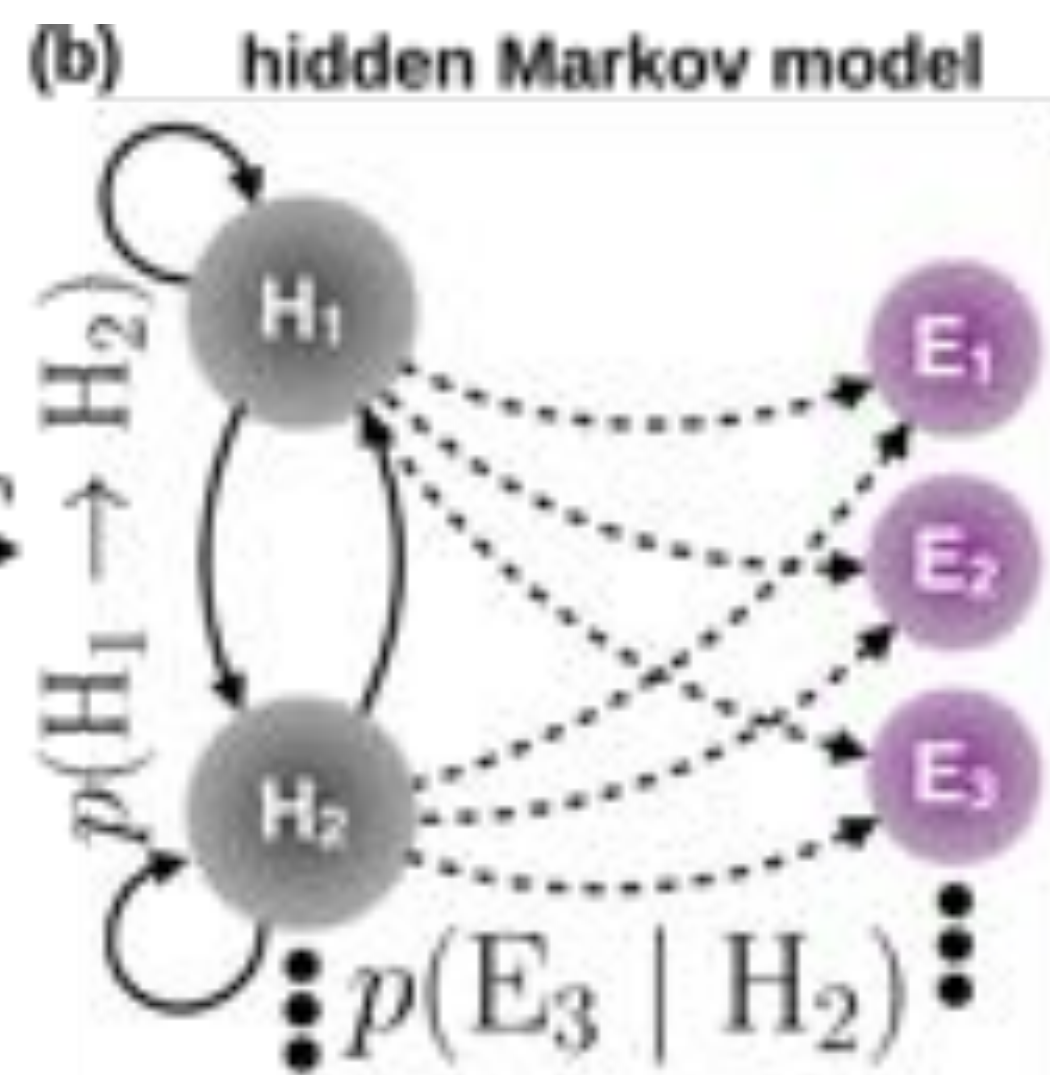
Automated detection of many-particle solvation states



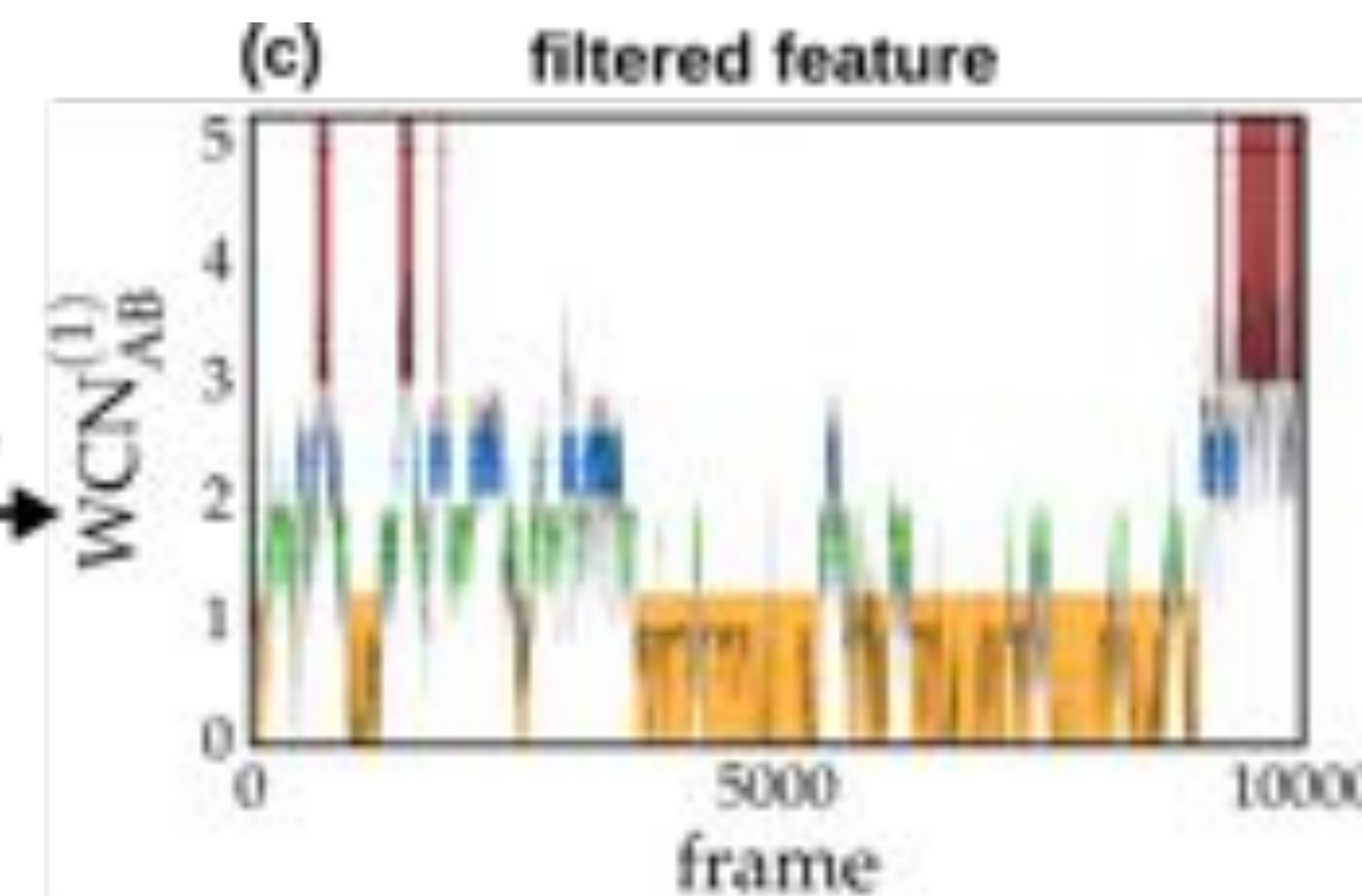
Coordination numbers as input features



choose N_h

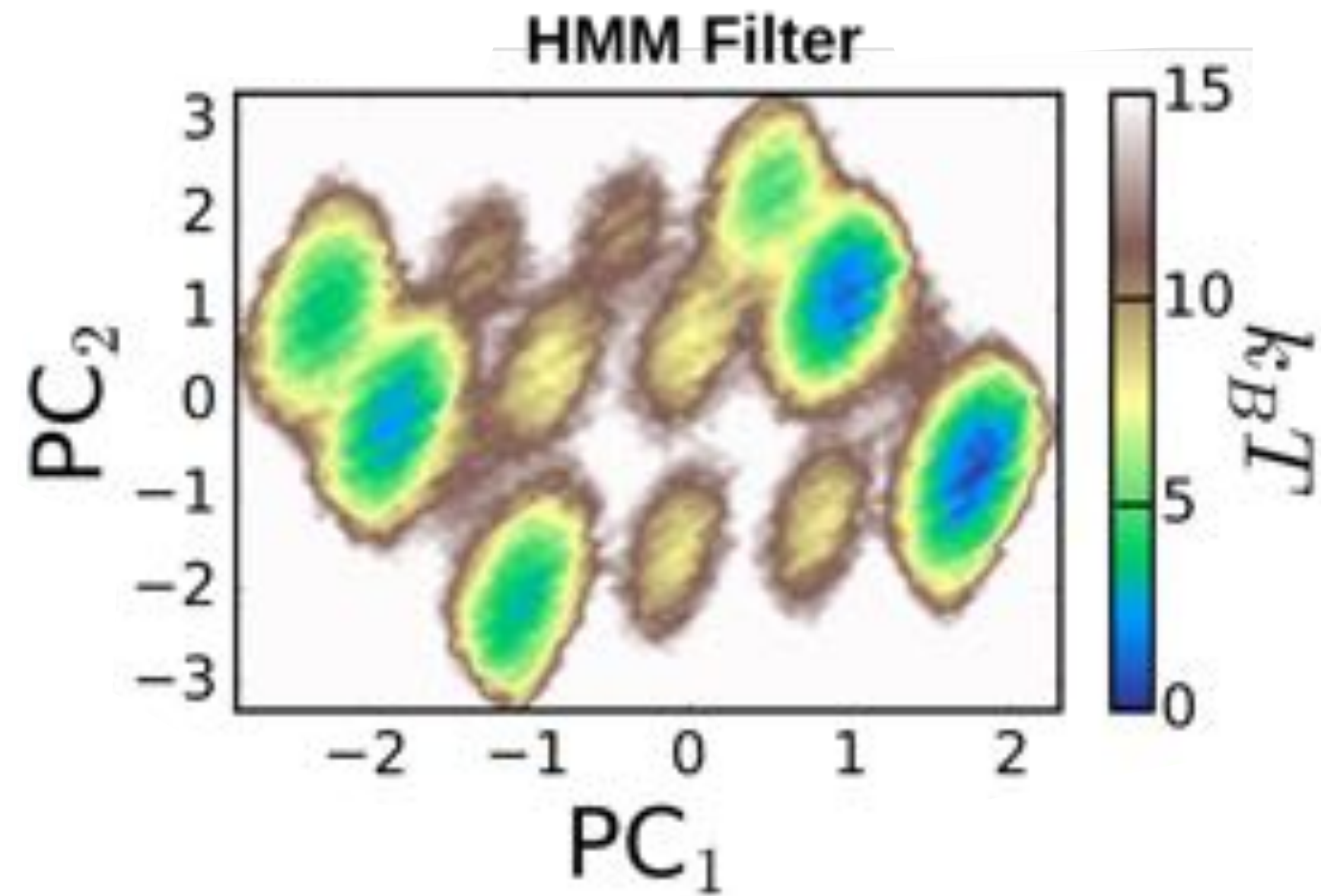
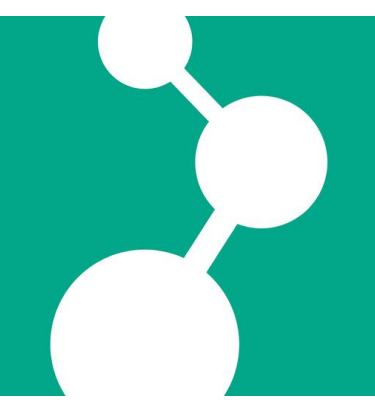


decode



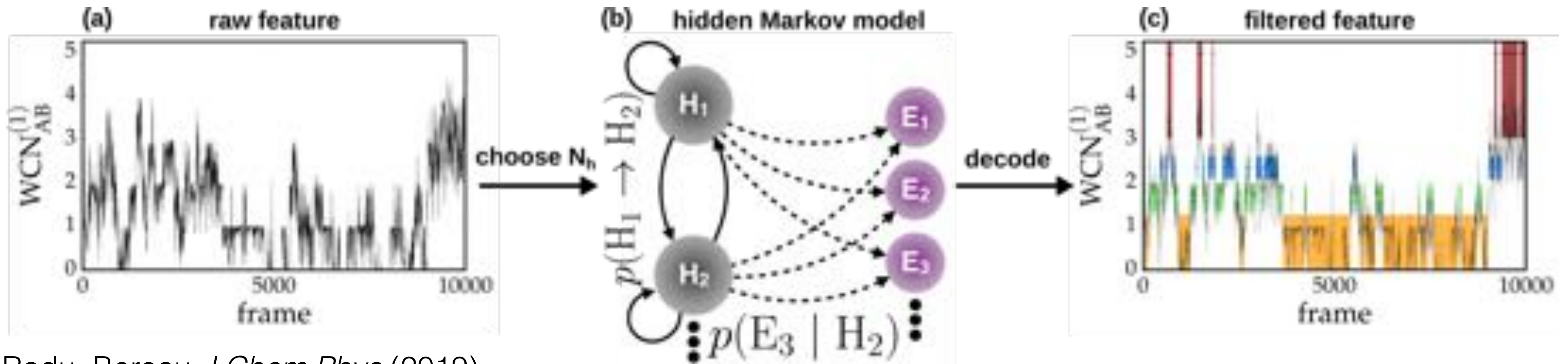


Automated detection of many-particle solvation states



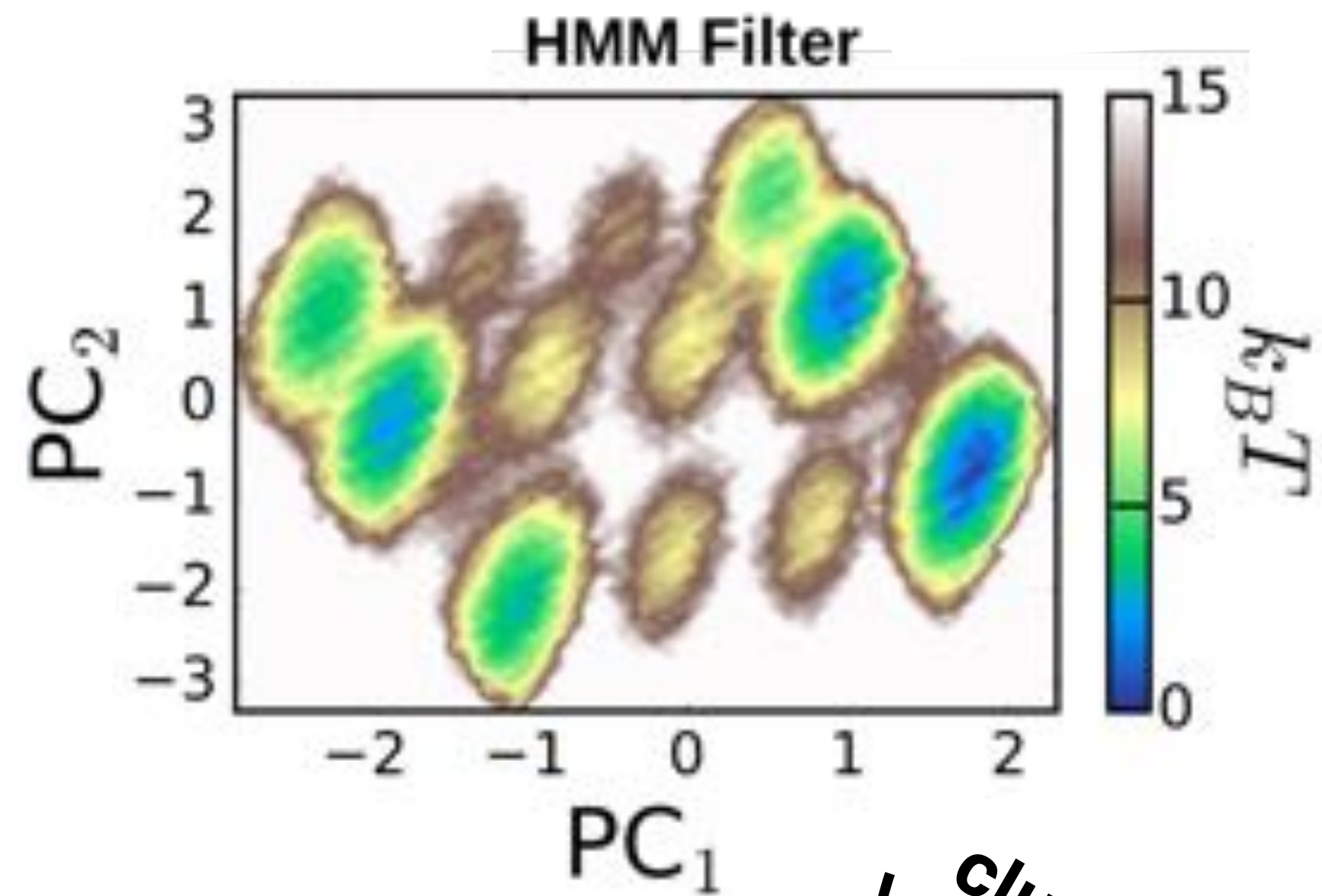
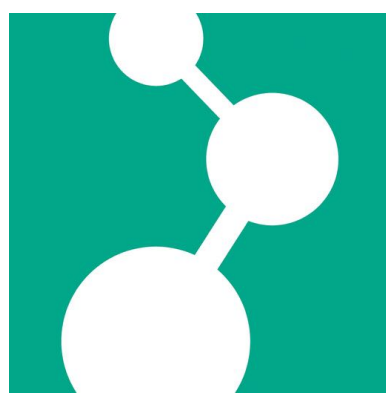
Structured free-energy landscape describing distinct solvation states

dimensionality reduction



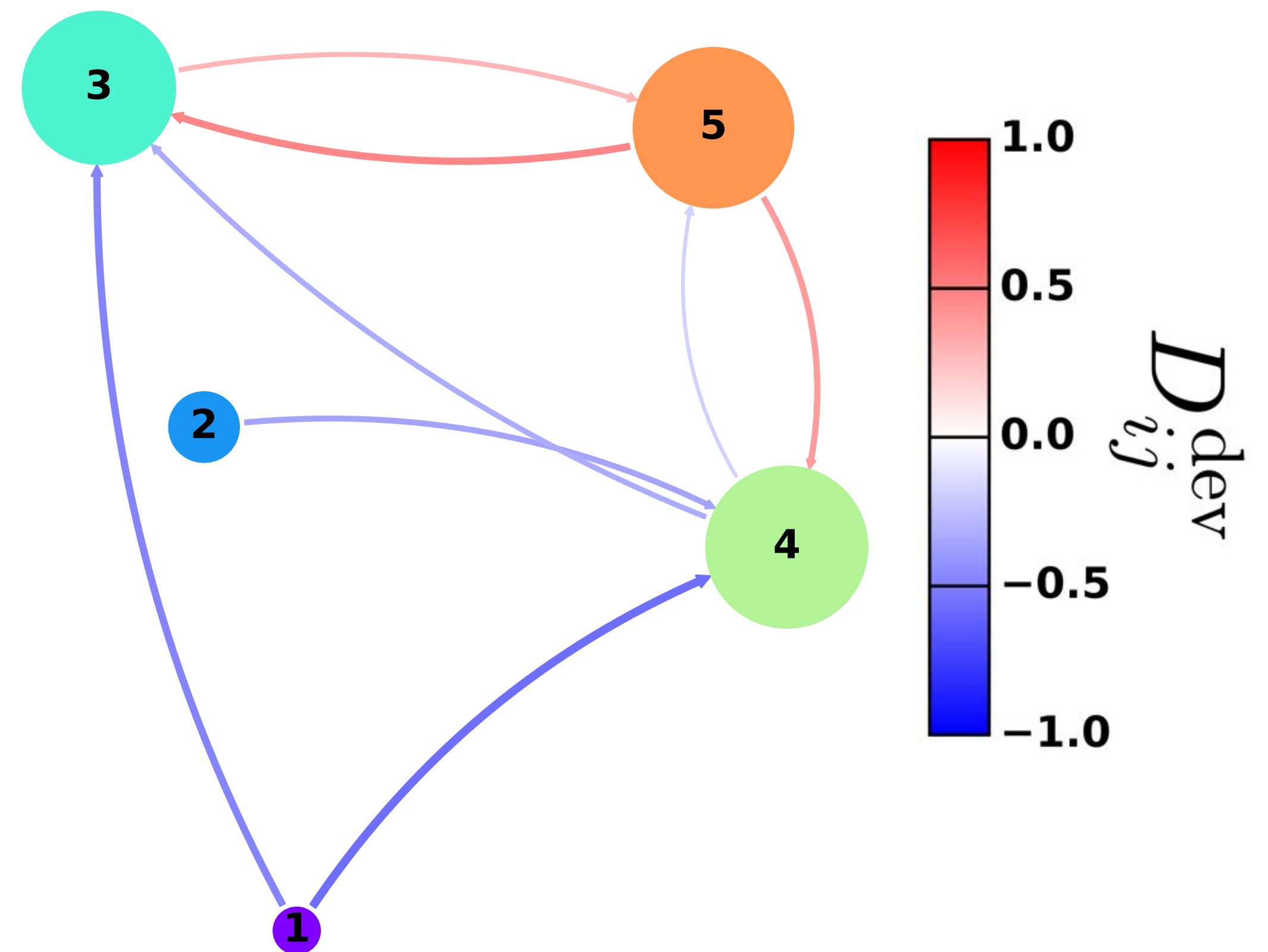


Automated detection of many-particle solvation states



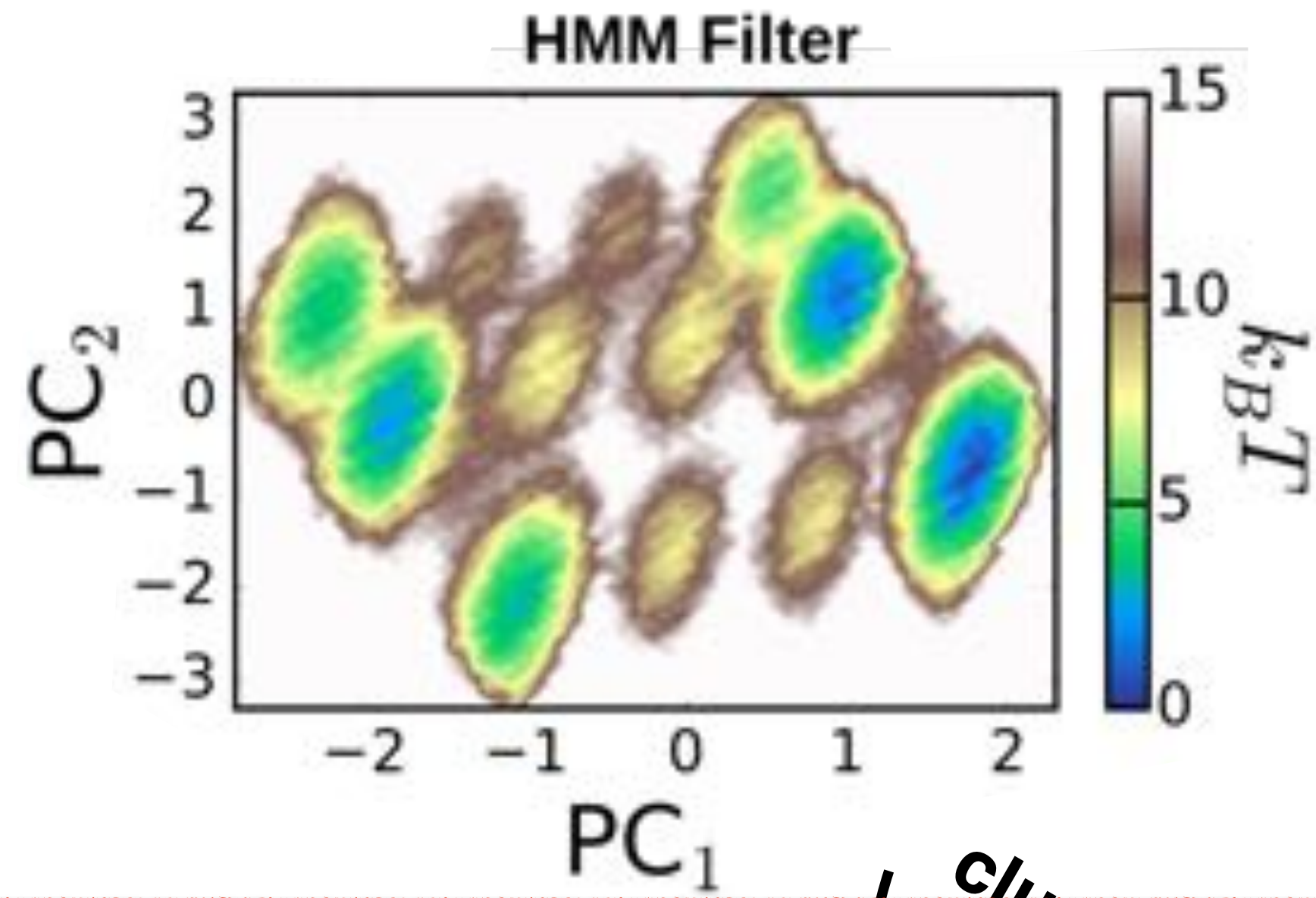
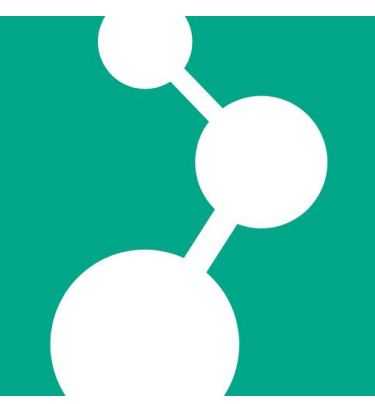
clustering + MSM
construction

network description of diffusion

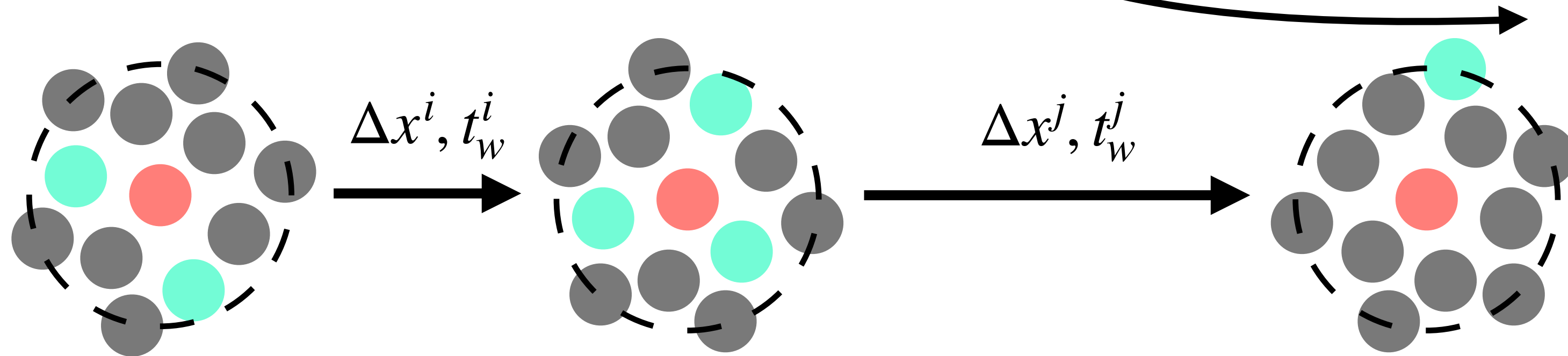




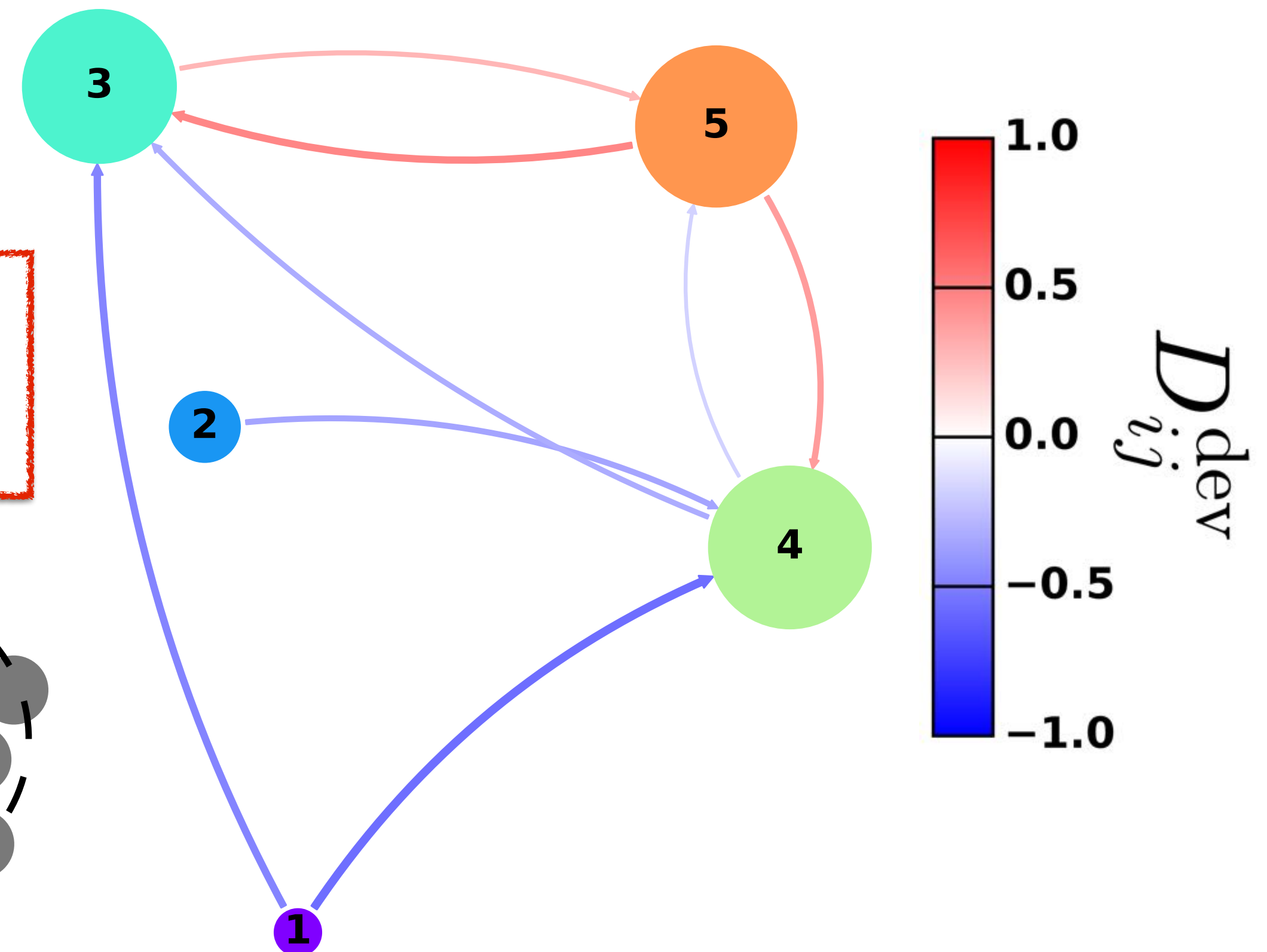
Automated detection of many-particle solvation states



- Mechanistic insight into solvation shell dynamics
- Quantification of dynamic heterogeneity

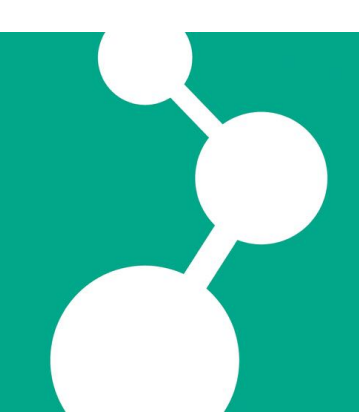


network description of diffusion

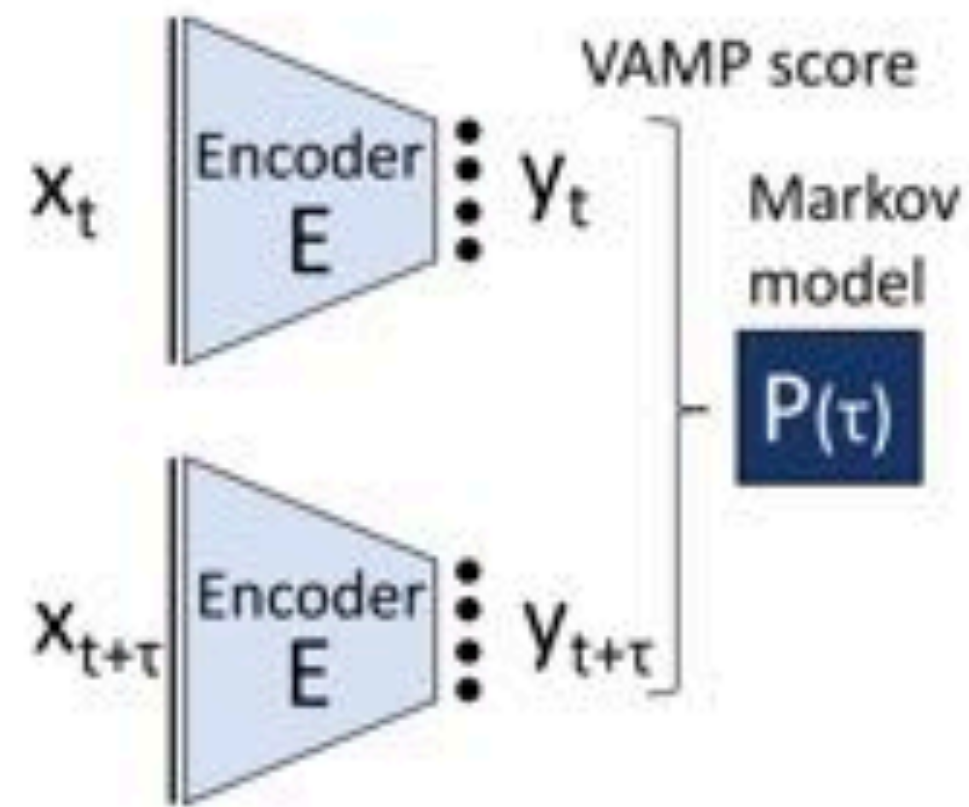




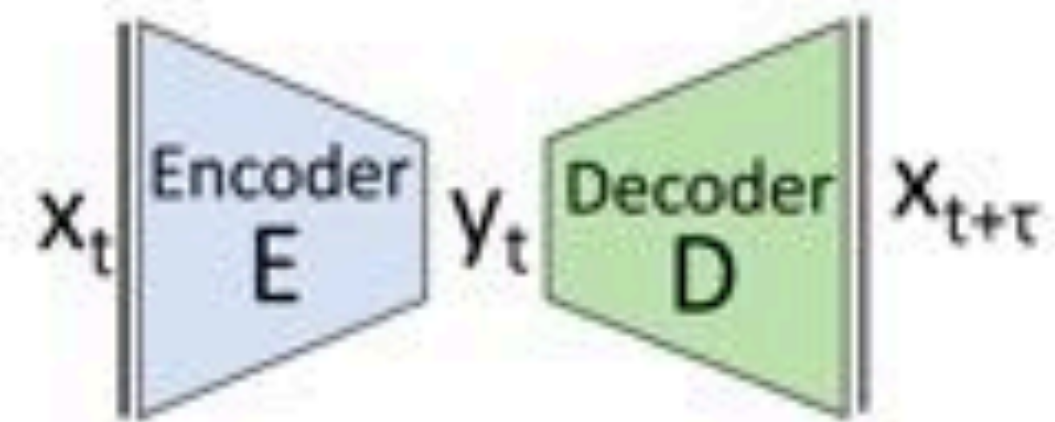
VAEs for MD analysis



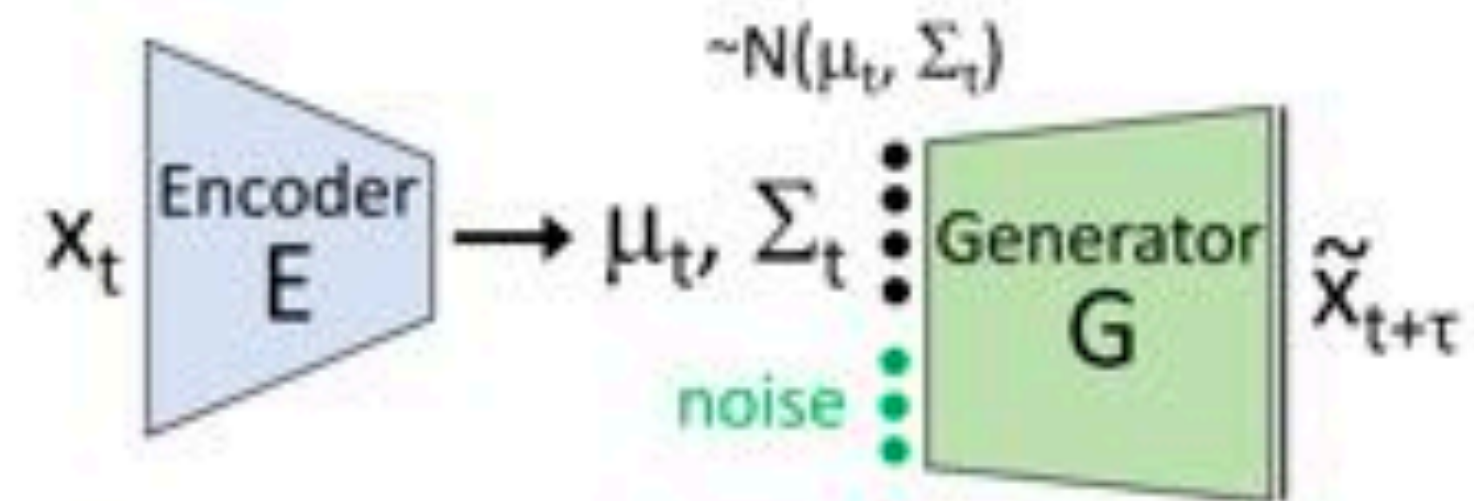
a) VAMPnet



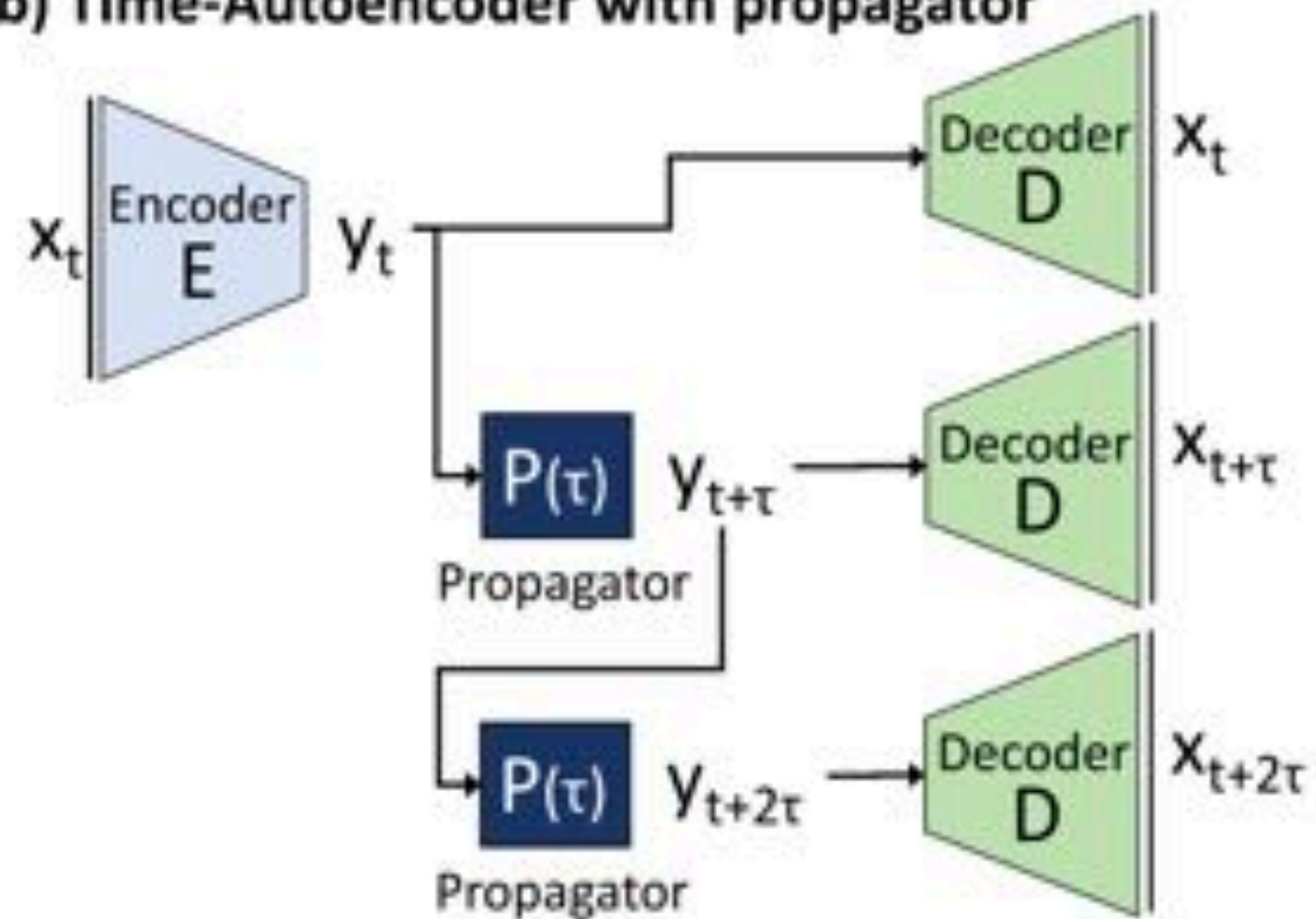
c) Time-Autoencoder (TAE)



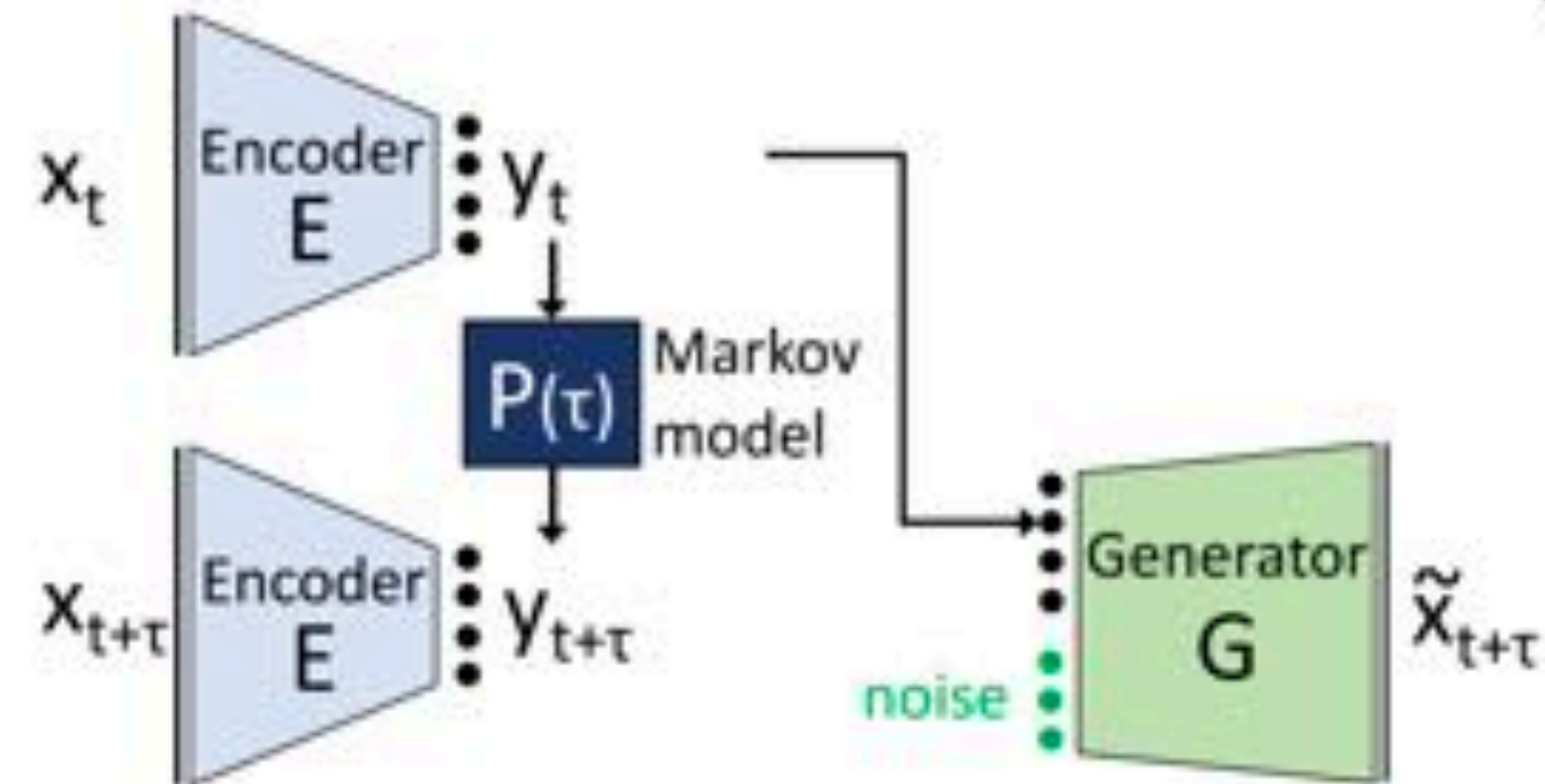
d) Variational time-Encoder



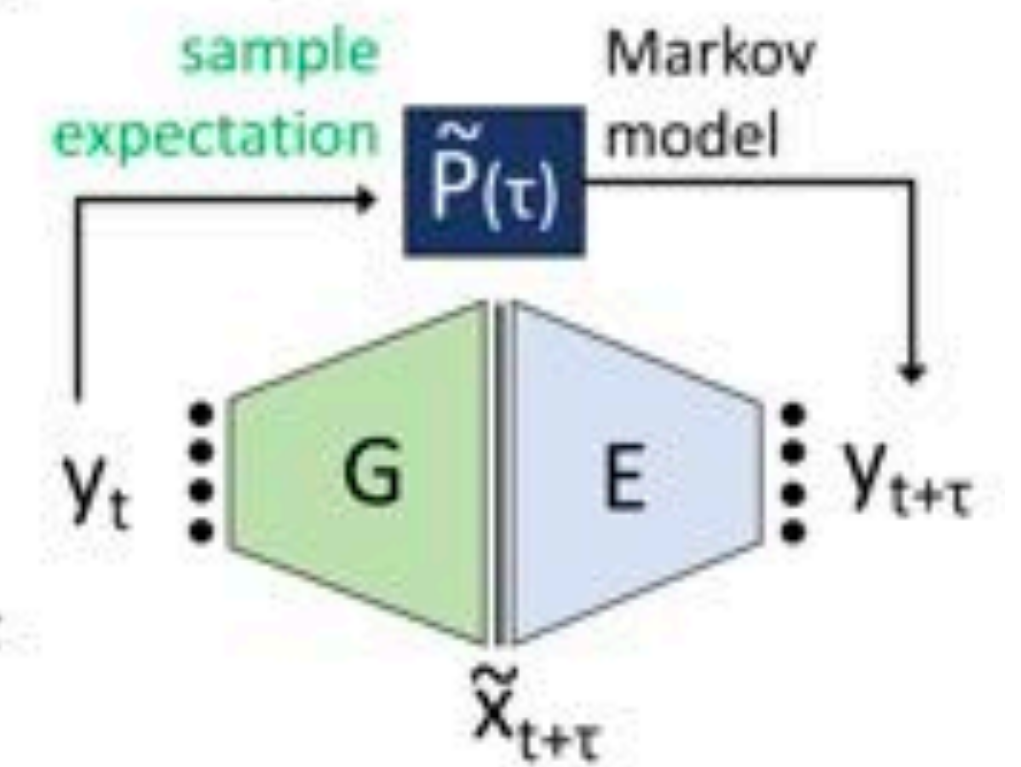
b) Time-Autoencoder with propagator



e) Deep Generative MSM

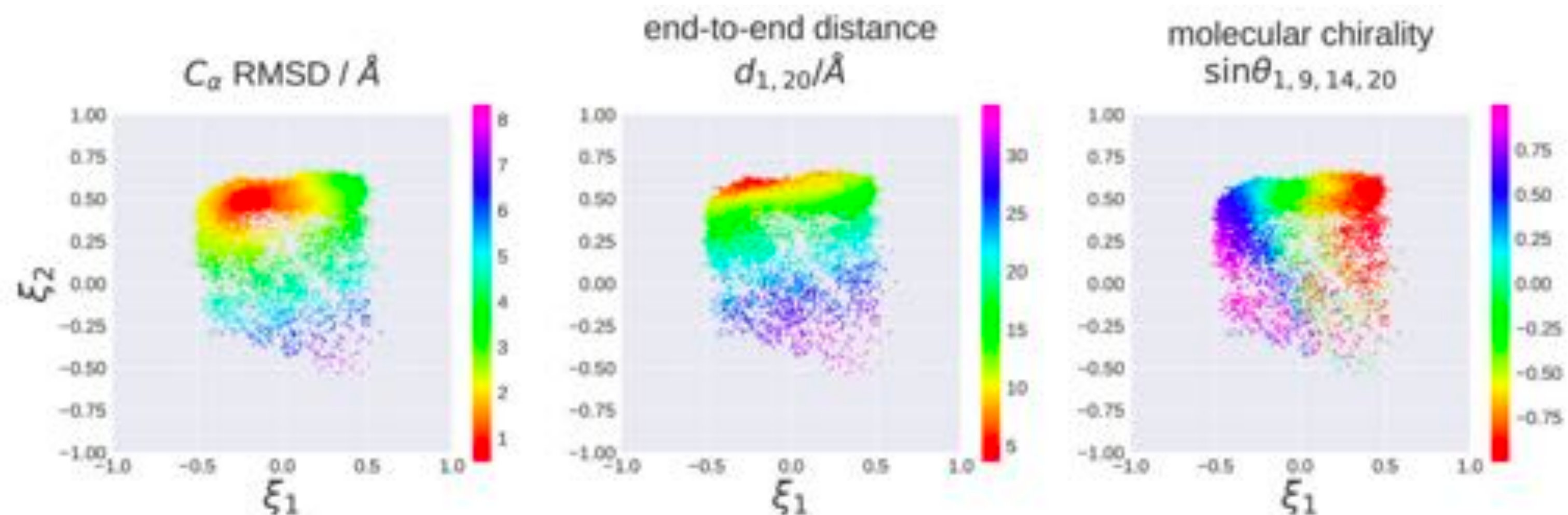
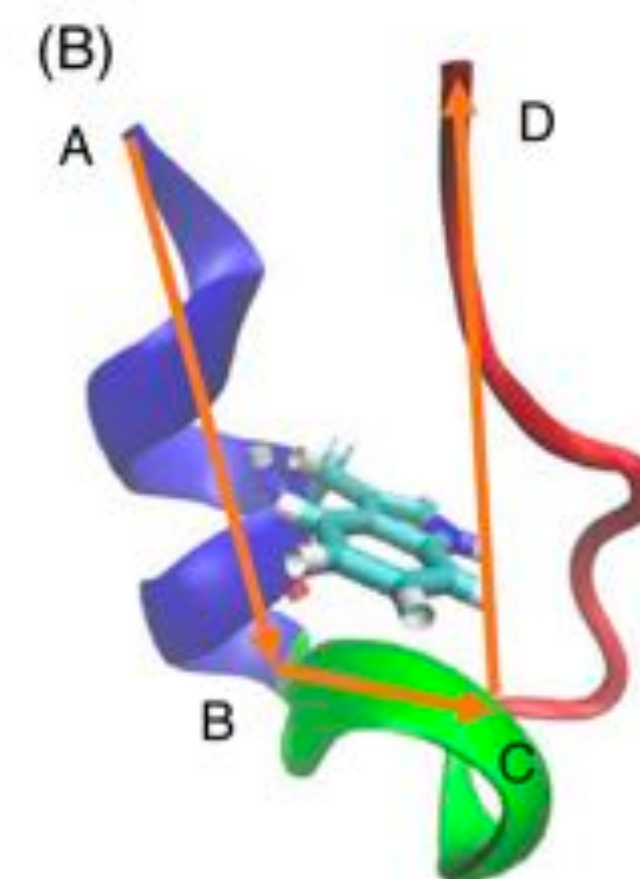
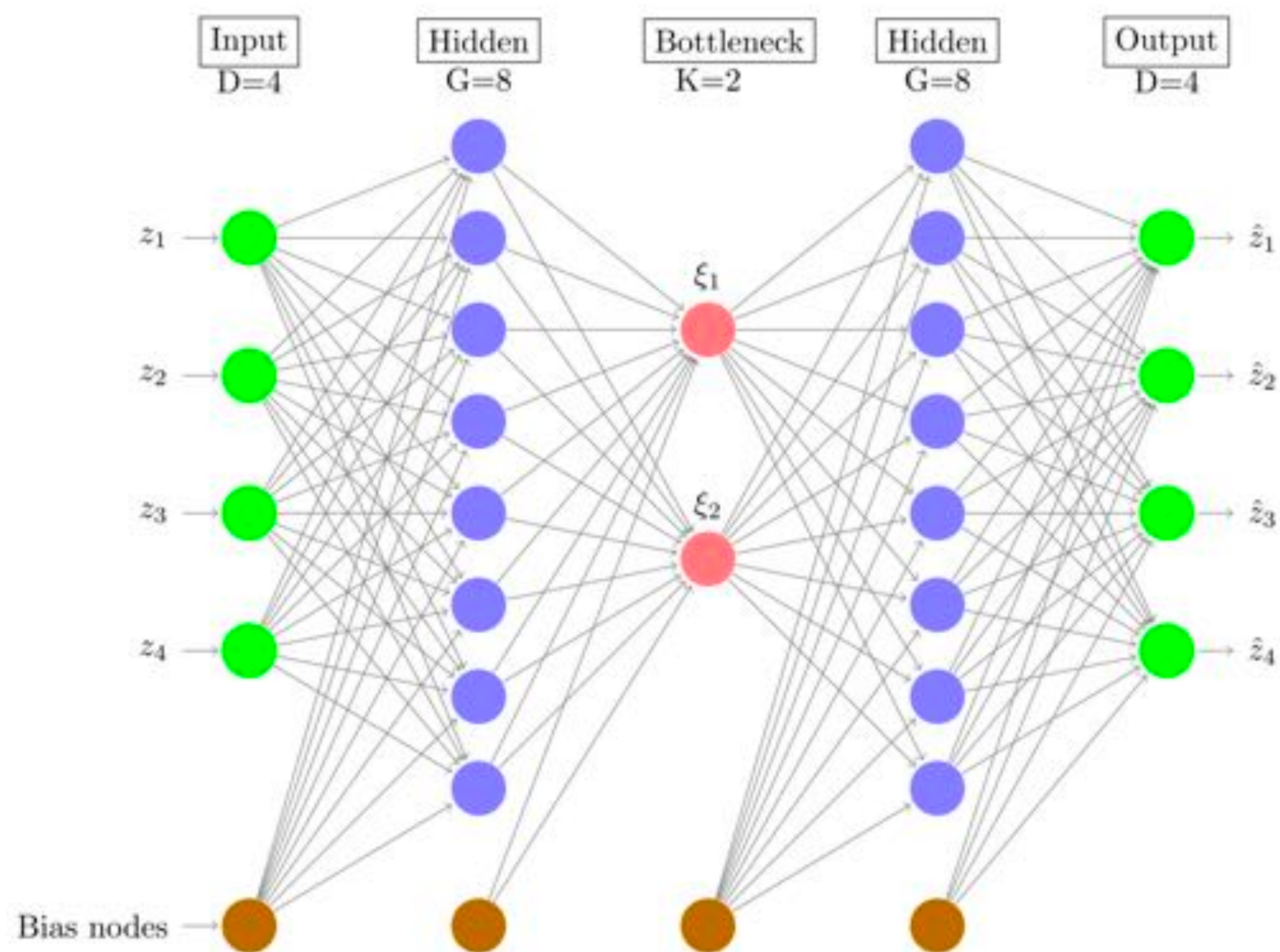
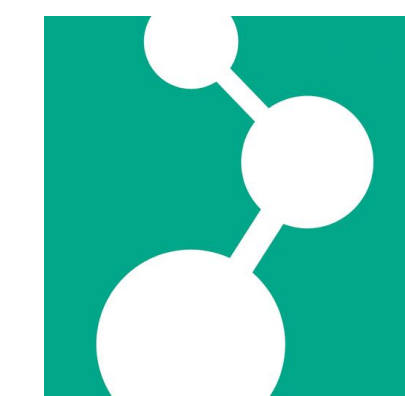


f) Rewiring Trick





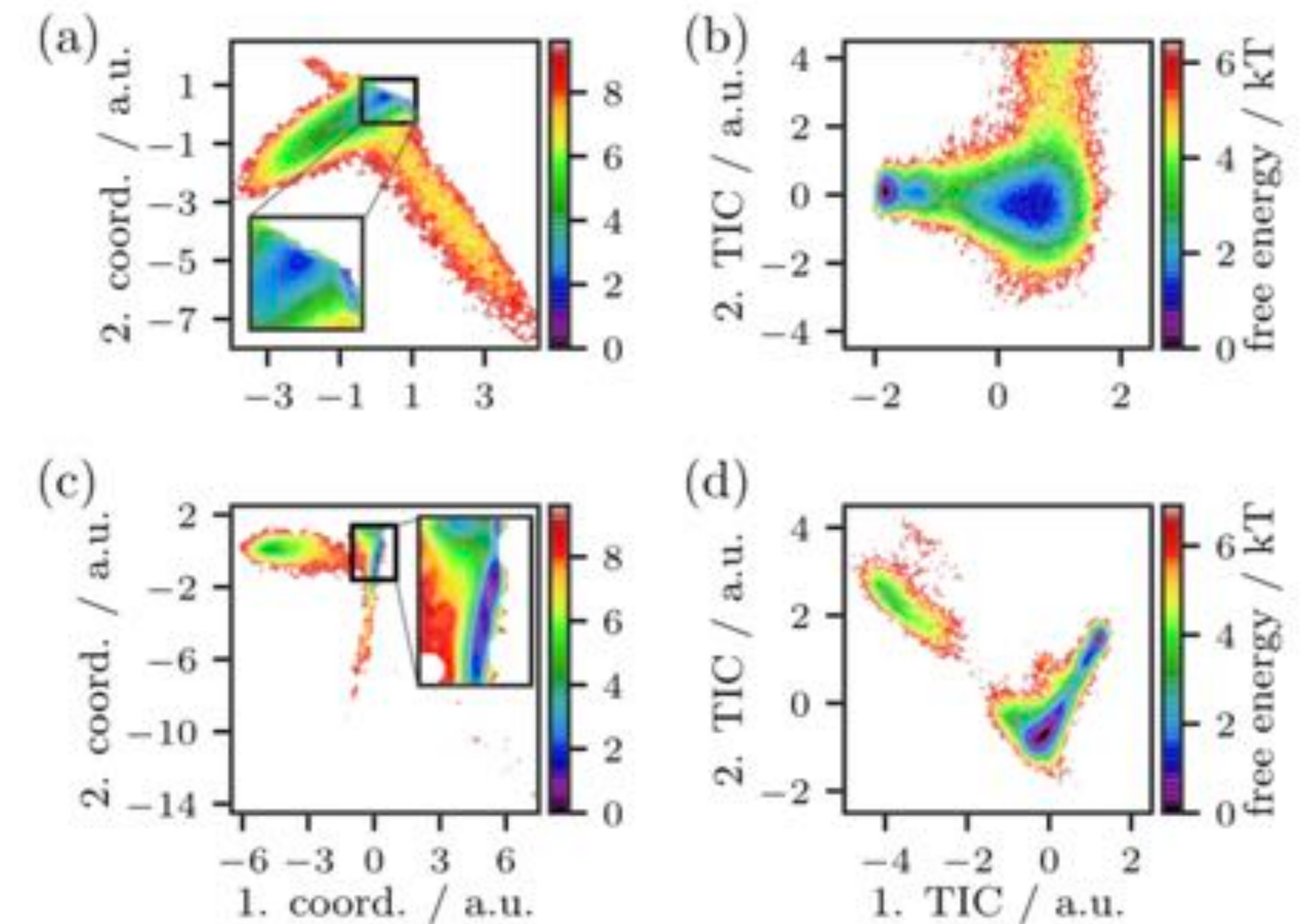
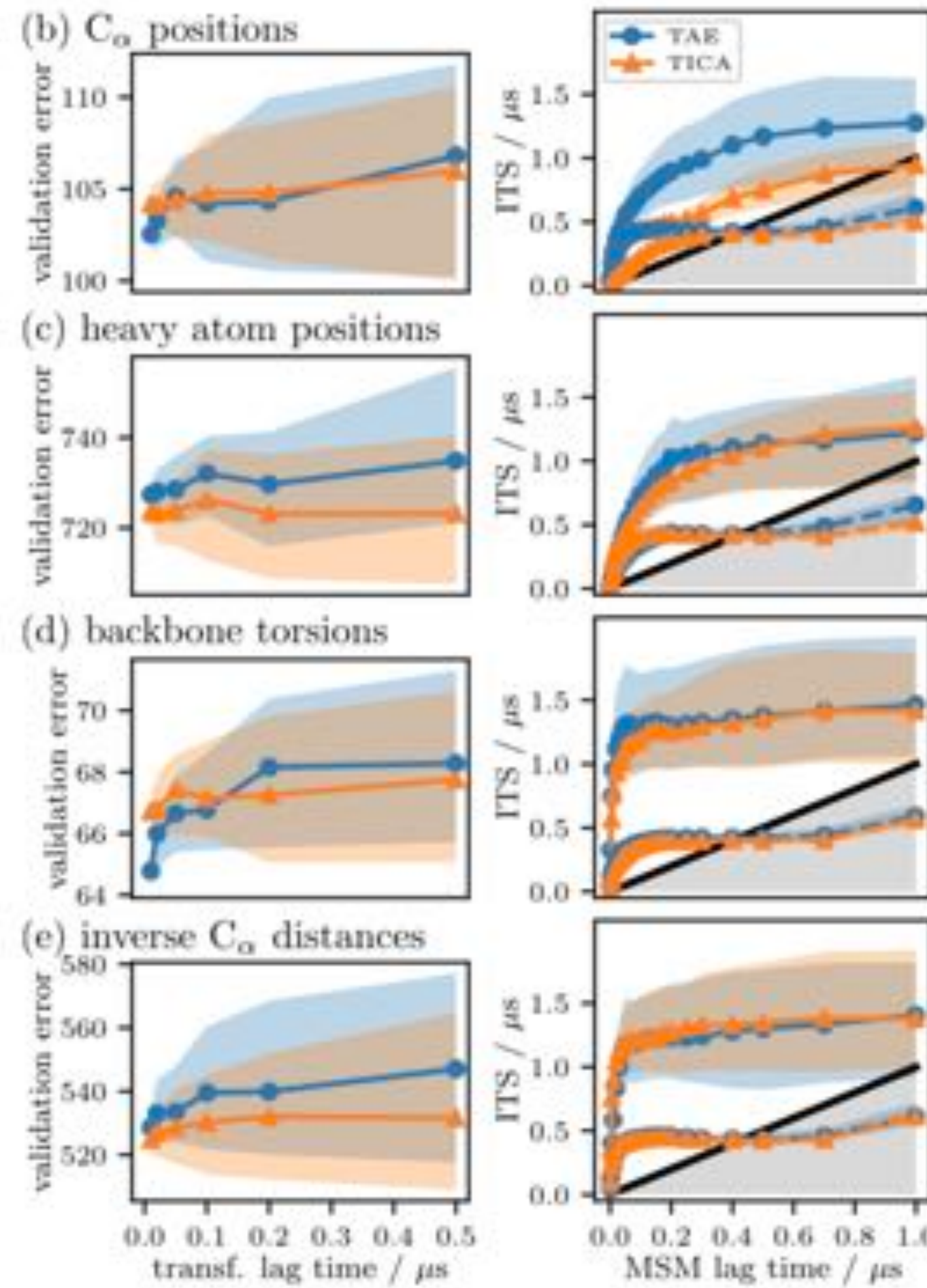
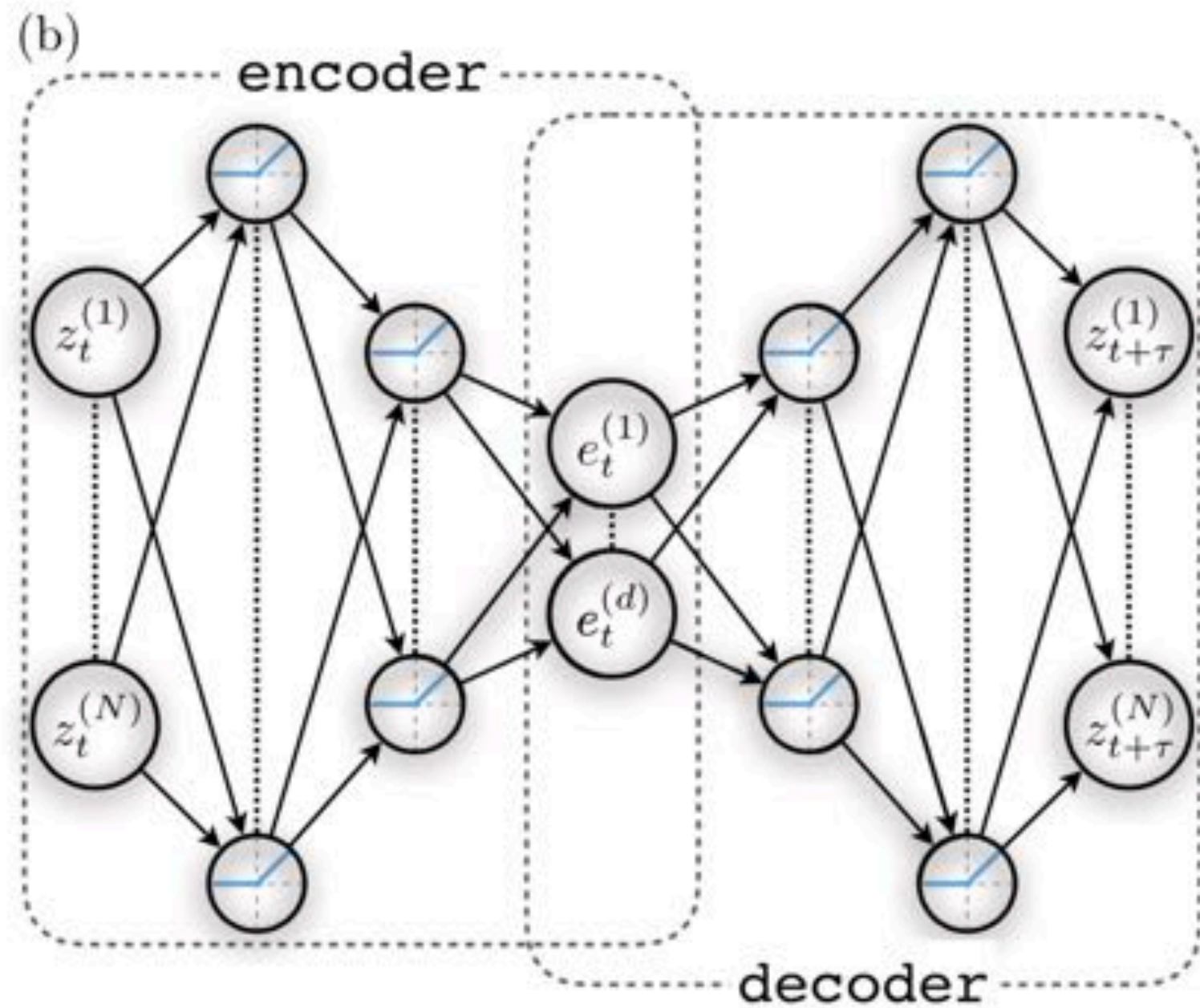
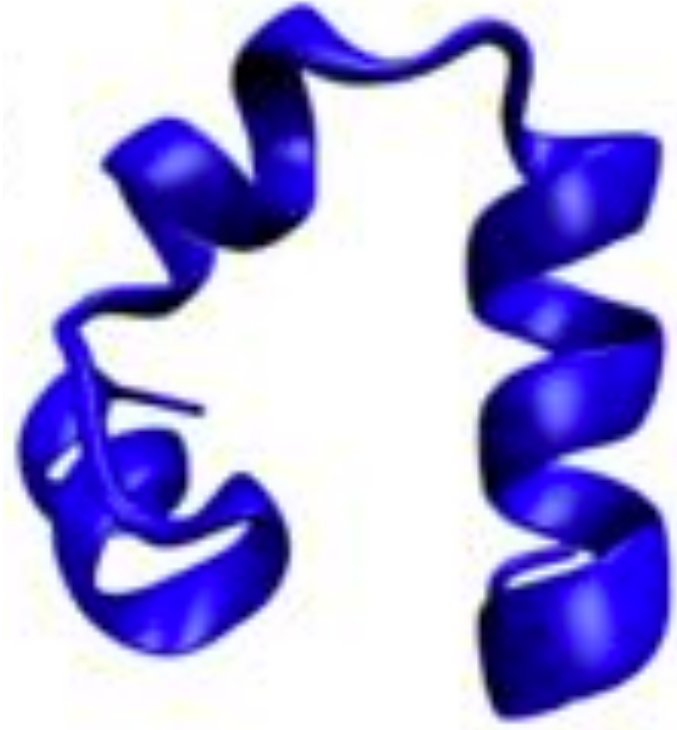
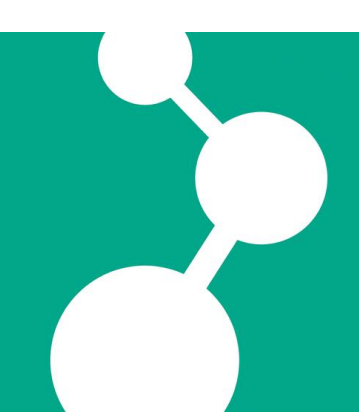
VAEs for collective variable discovery



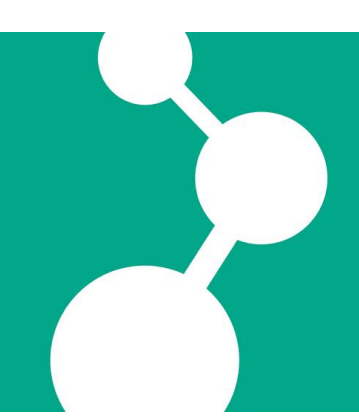
Chen, Ferguson *J Comp Chem* (2018) “Molecular enhanced sampling with autoencoders: On-the-fly collective variable discovery and accelerated free energy landscape exploration”



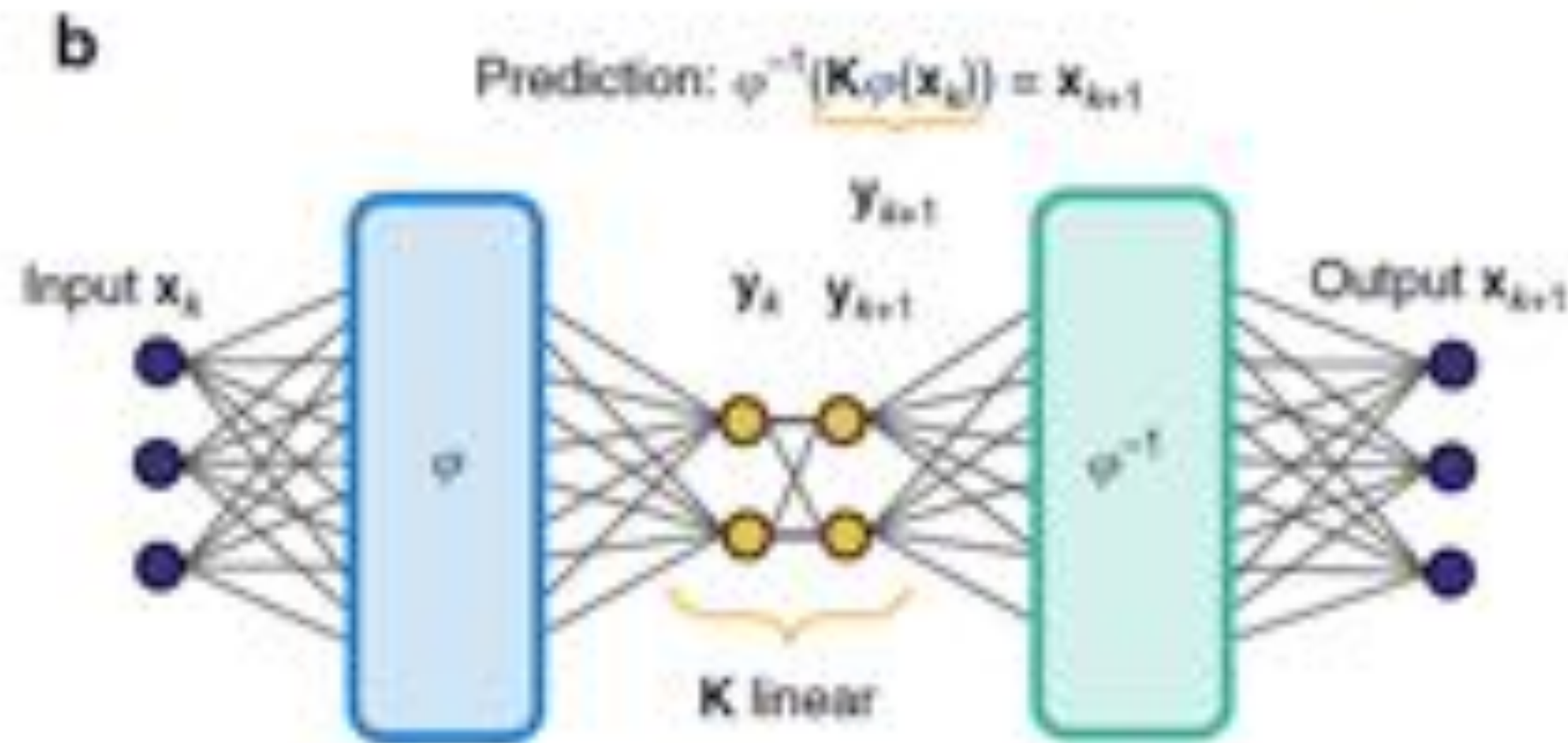
Time-lagged autoencoders



Christoph Wehmeyer and Frank Noe *JCP* (2018) “Time-lagged autoencoders: Deep learning of slow collective variables for molecular kinetics”



Loss functions



$$\mathcal{L} = \alpha_1 (\mathcal{L}_{\text{recon}} + \mathcal{L}_{\text{pred}}) + \mathcal{L}_{\text{lin}} + \alpha_2 \mathcal{L}_{\infty} + \alpha_3 \|\mathbf{W}\|_2^2$$

$$\mathcal{L}_{\text{recon}} = \|x_1 - \varphi^{-1}(\varphi(x_1))\|_{\text{MSE}}$$

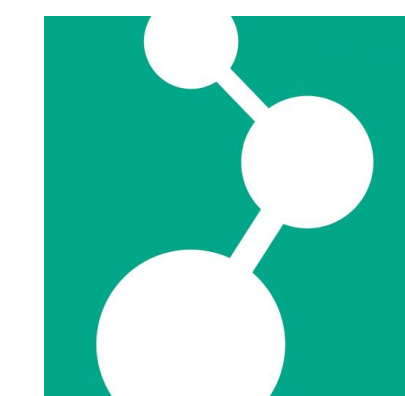
$$\mathcal{L}_{\text{pred}} = \frac{1}{S_p} \sum_{m=1}^S \|x_{m+1} - \varphi^{-1}(K^m \varphi(x_1))\|_{\text{MSE}}$$

$$\mathcal{L}_{\text{lin}} = \frac{1}{T-1} \sum_{m=1}^{T-1} \|\varphi(x_{m+1}) - K^m \varphi(x_1)\|_{\text{MSE}}$$

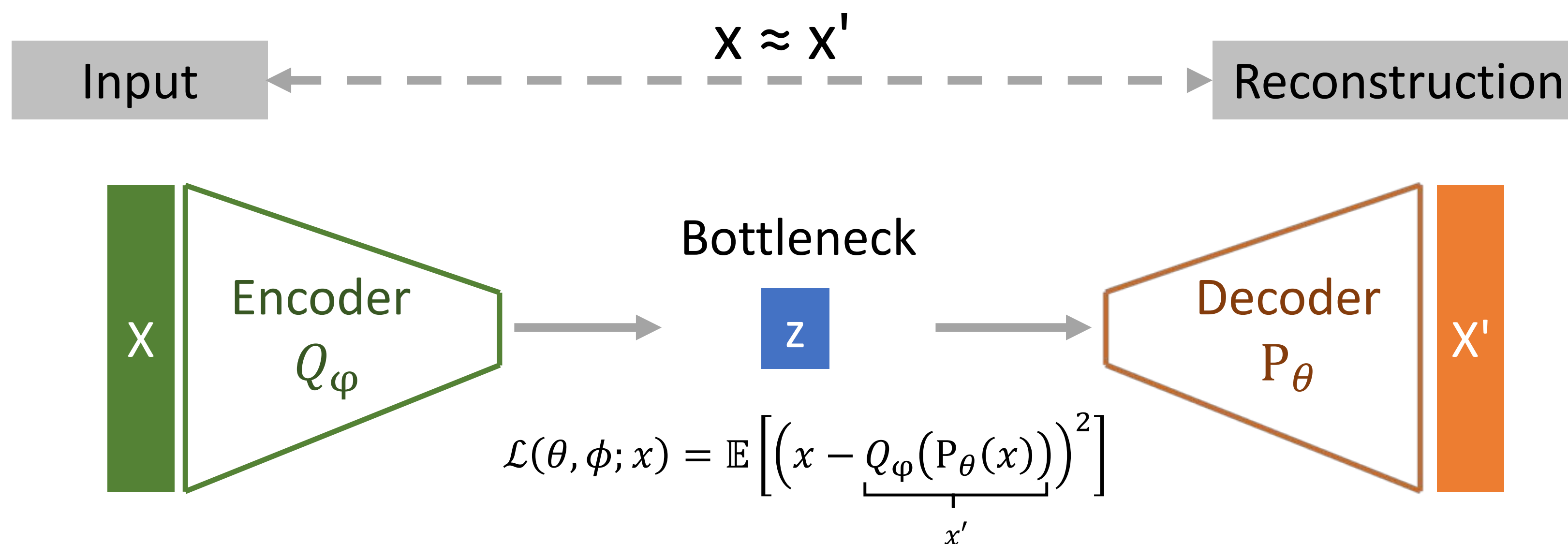
$$\mathcal{L}_{\infty} = \|x_1 - \varphi^{-1}(\varphi(x_1))\|_{\infty} + \|x_2 - \varphi^{-1}(K\varphi(x_1))\|_{\infty}$$



Interpretable embeddings for molecular kinetics using deep learning



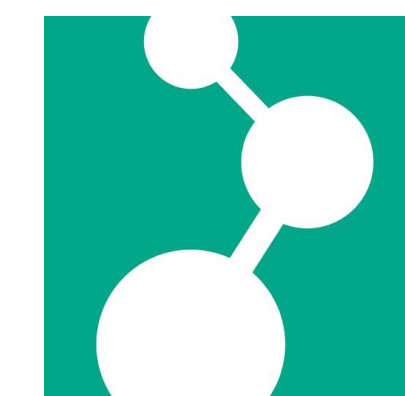
Standard Autoencoder



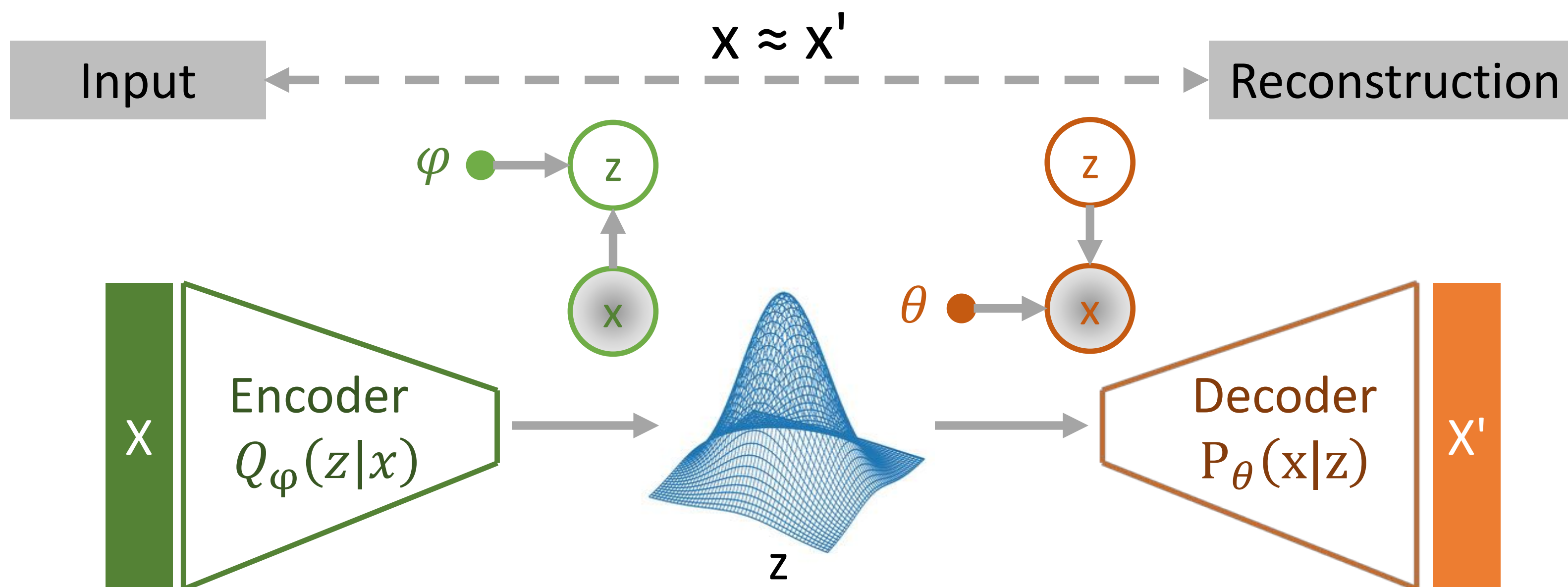
Yasemin Bozkurt Varolgunes



Interpretable embeddings for molecular kinetics using deep learning



Variational Autoencoder with Unimodal Gaussian Prior



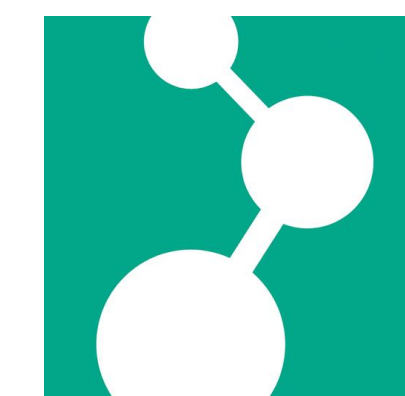
$$\mathcal{L}(\theta, \phi; x) = \mathbb{E}_{Q_\phi(x,z)} [\log P_\theta(x,z) - \log Q_\phi(z|x)]$$
$$\mathcal{L}(\theta, \phi; x) = \underbrace{\mathbb{E}_{Q_\phi(z|x)} [\log P_\theta(x|z)]}_{\text{Reconstruction}} - \underbrace{KL(Q_\phi(z|x) || P(z))}_{\text{Regularization}}$$



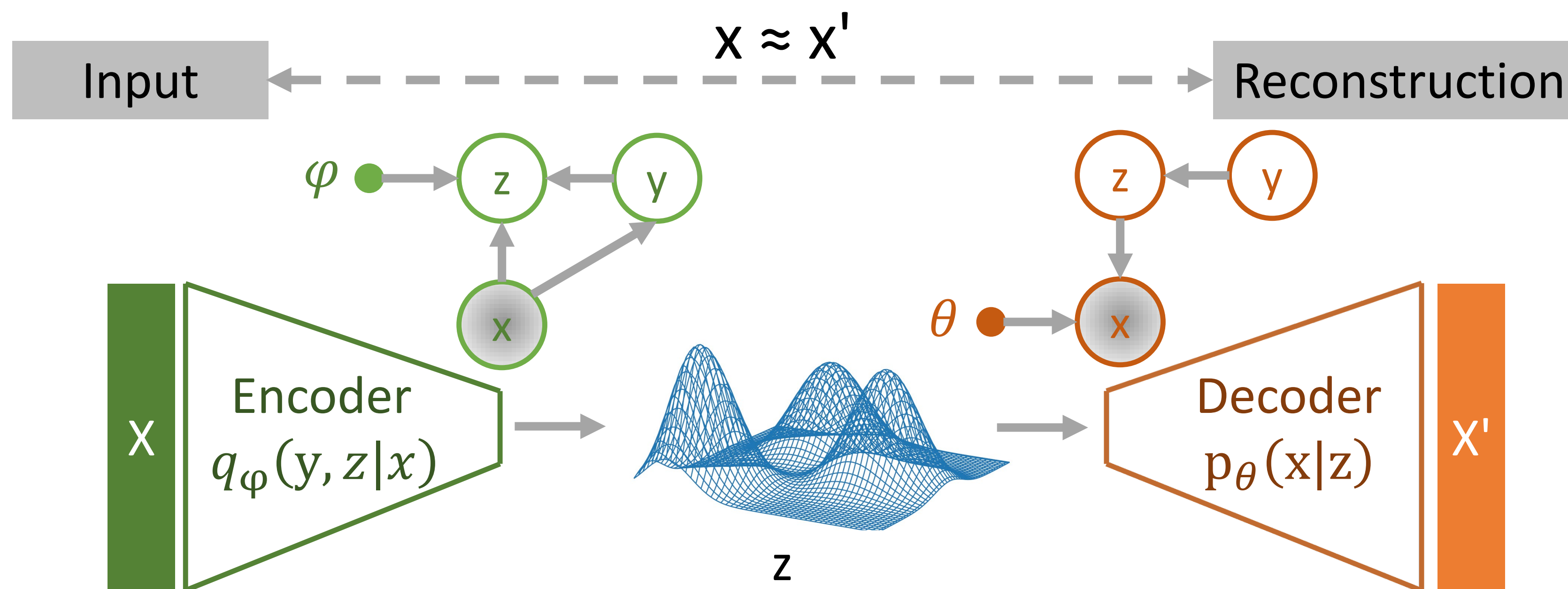
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Interpretable embeddings for molecular kinetics using deep learning



Gaussian Mixture Variational Autoencoder



$$\mathcal{L}(\theta, \phi; x) = \mathbb{E}_{q_\phi(y, z|x)} [\log p_\theta(x, y, z) - \log q_\phi(y, z|x)]$$

$$\mathcal{L}(\theta, \phi; x) = \mathbb{E}_{q_\phi(y, z|x)} \left[\underbrace{\log \frac{p_\theta(y)}{q_\phi(y|x)}}_{\text{clustering}} + \underbrace{\log \frac{p_\theta(z|y)}{q_\phi(z|x, y)}}_{\text{regularization}} + \underbrace{\log p_\theta(x|z)}_{\text{reconstruction}} \right]$$

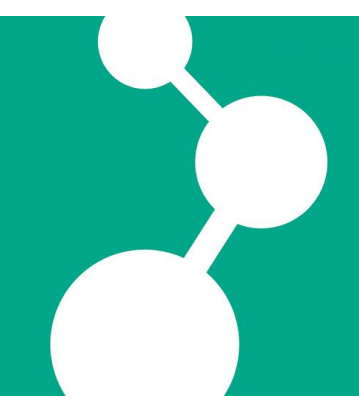
- ▶ Dimensionality reduction + Clustering
- ▶ Synthetic trajectory generation



Yasemin Bozkurt Varolgunes

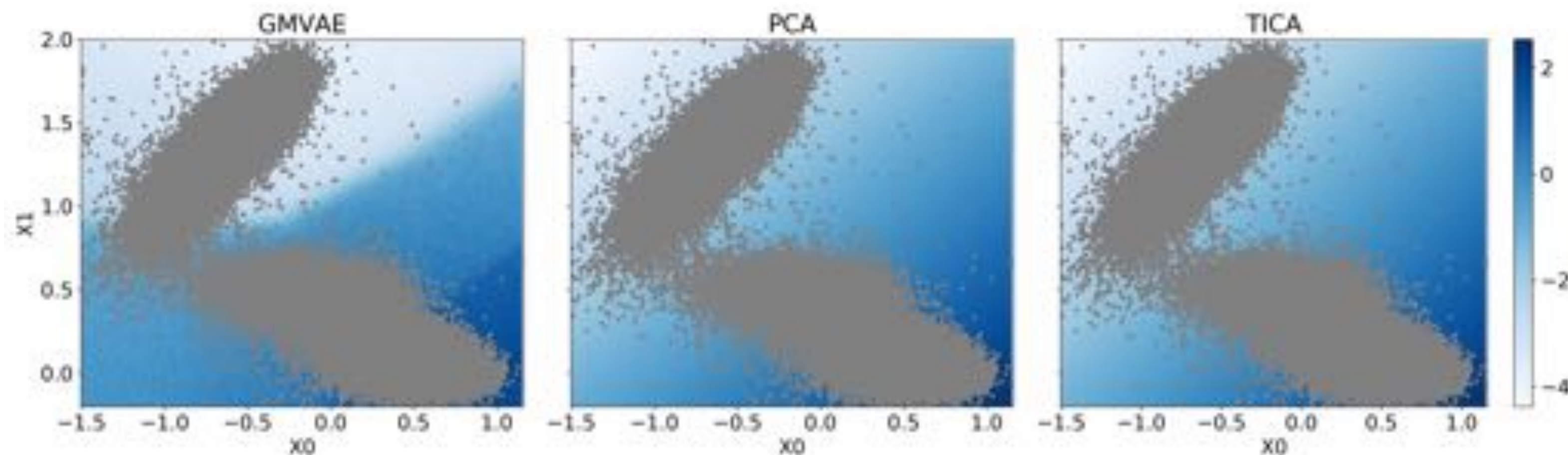
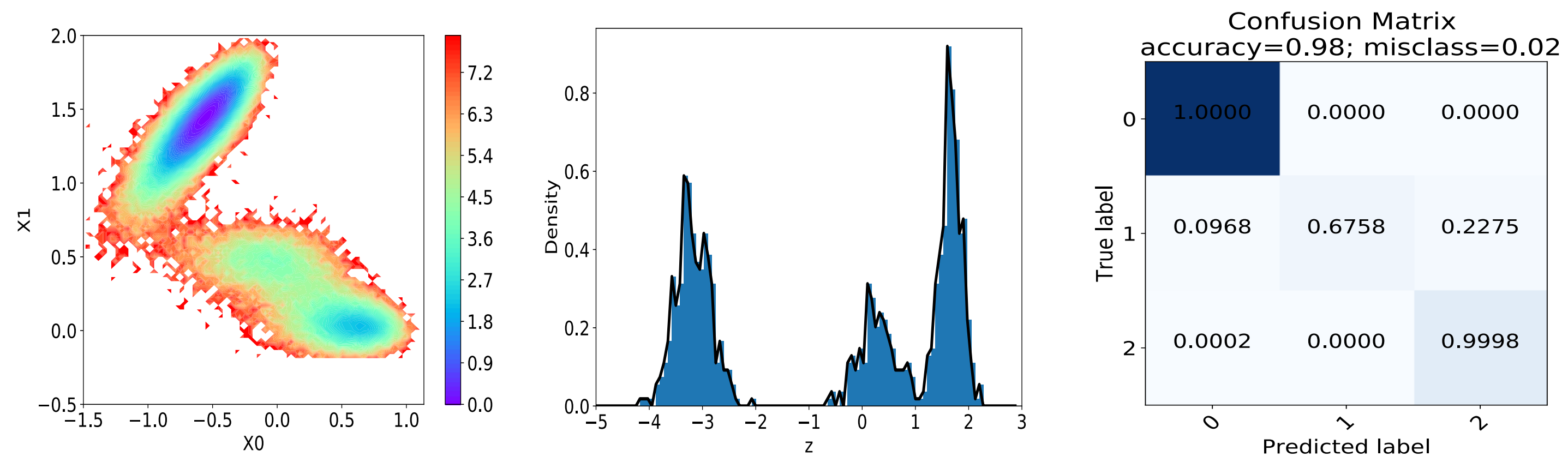


Interpretable embeddings for molecular kinetics using deep learning



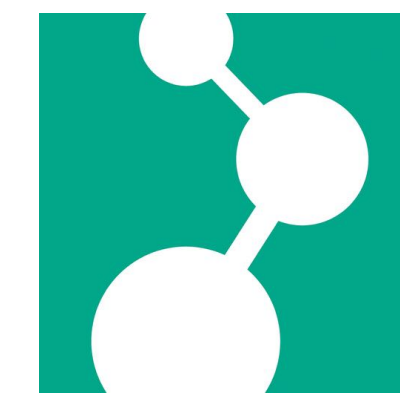
Two dimensional Muller-Brown potential

$$V(x) = \sum_{j=1}^4 A_j \exp[a_j(x_1 - X_j)^2 + b_j(x_1 - X_j)(x_2 - Y_j) + c_j(x_2 - Y_j)^2]$$

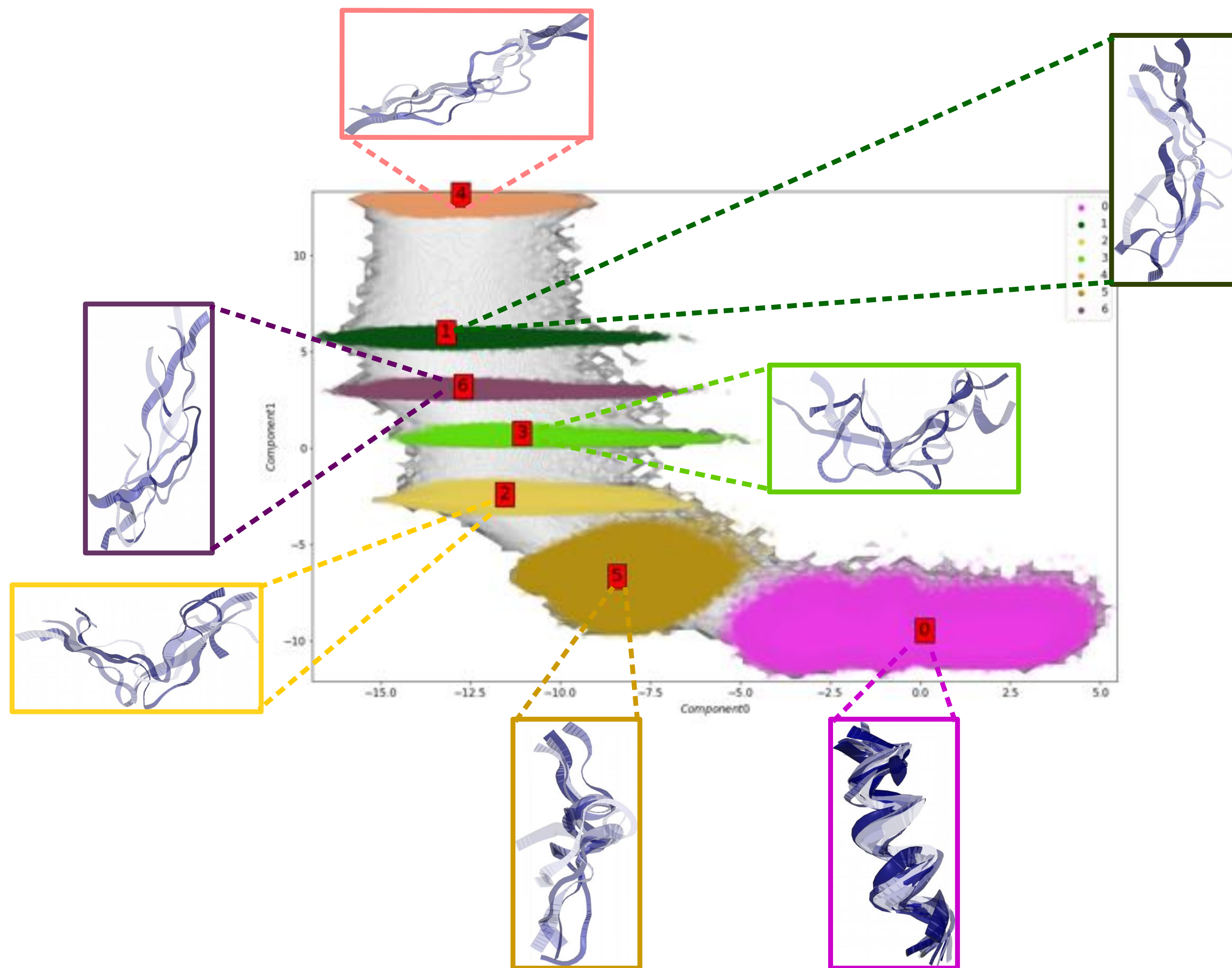
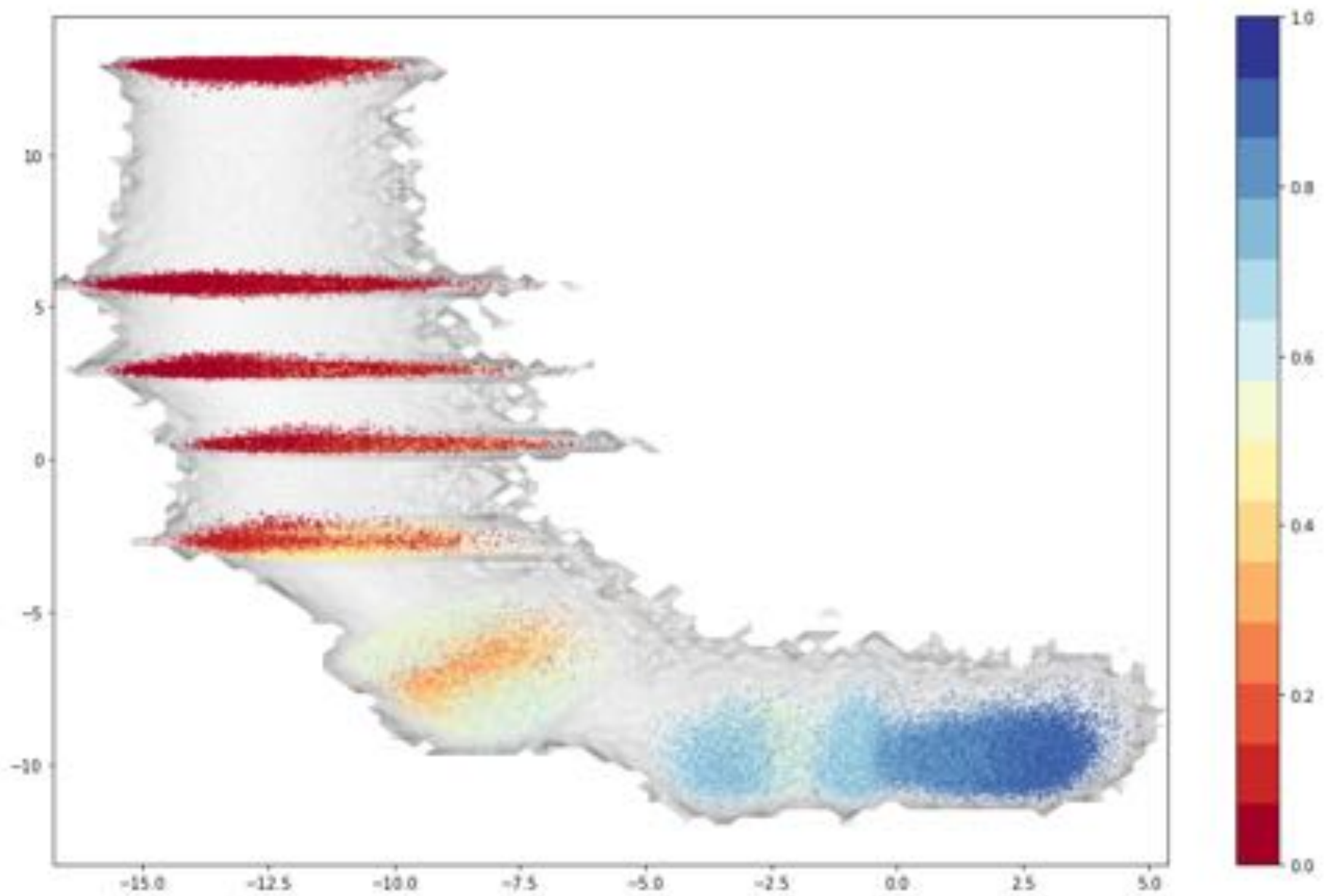
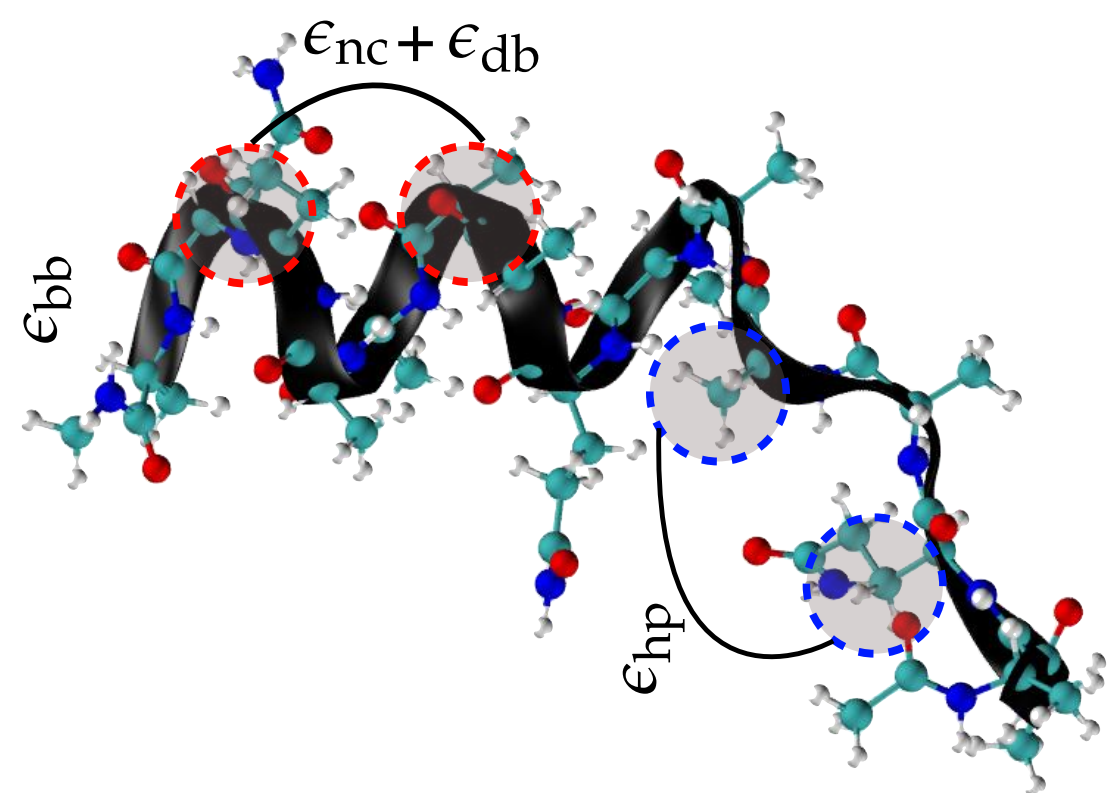




Interpretable embeddings for molecular kinetics using deep learning

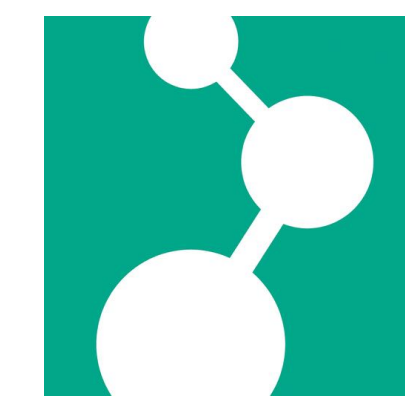


(AAQAA)₃





Variational autoencoders for dimensionality reduction and clustering of molecular dynamics data



Thank you for your attention!

Contact, papers, and current updates on my research @ RudzinskiResearch.com