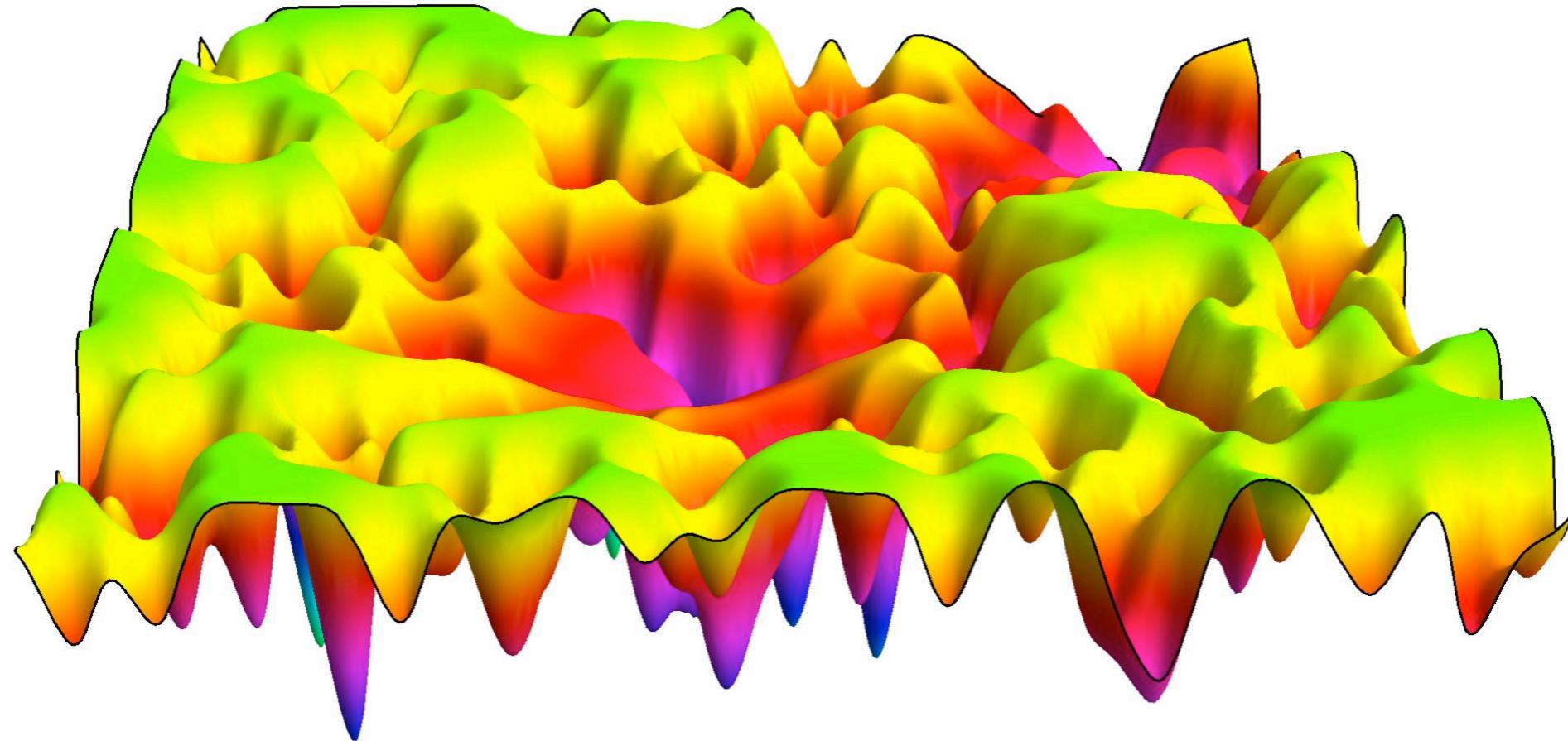


Data-driven discovery and enhanced sampling in slow collective variables



K. Shmilovich and A.L. Ferguson "Girsanov Reweighting Enhanced Sampling Technique (GREST): On-the-fly data-driven discovery of and enhanced sampling in slow collective variables" (submitted, 2023)

Brin Math Research Center: Rare Events, Analysis, Numerics, and Applications
27 February 2023
Andrew Ferguson, U. Chicago



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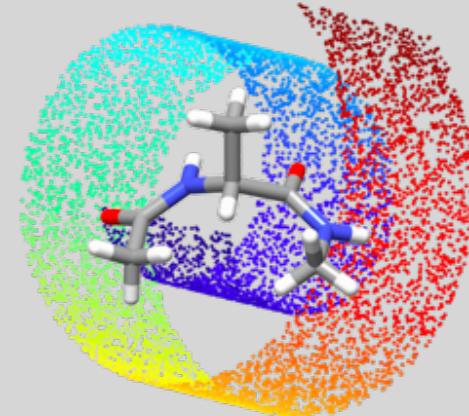
Acknowledgements



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Prof. Aaron Esser-Kahn (UofC)
Prof. Juan de Pablo (UofC)
Prof. Amish Patel (UPenn)

We are hiring for three postdocs

Ferguson Laboratory



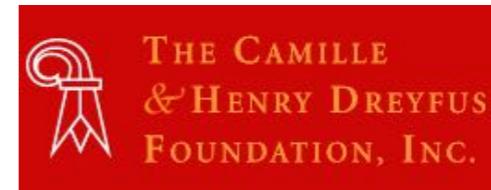
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Disclosure



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evozyne

<https://evozyne.com>

Data-driven protein engineering startup in Chicago
ALF is a co-founder, consultant, equity holder, and IP licensor
No funding or role of Evozyne in the reported work

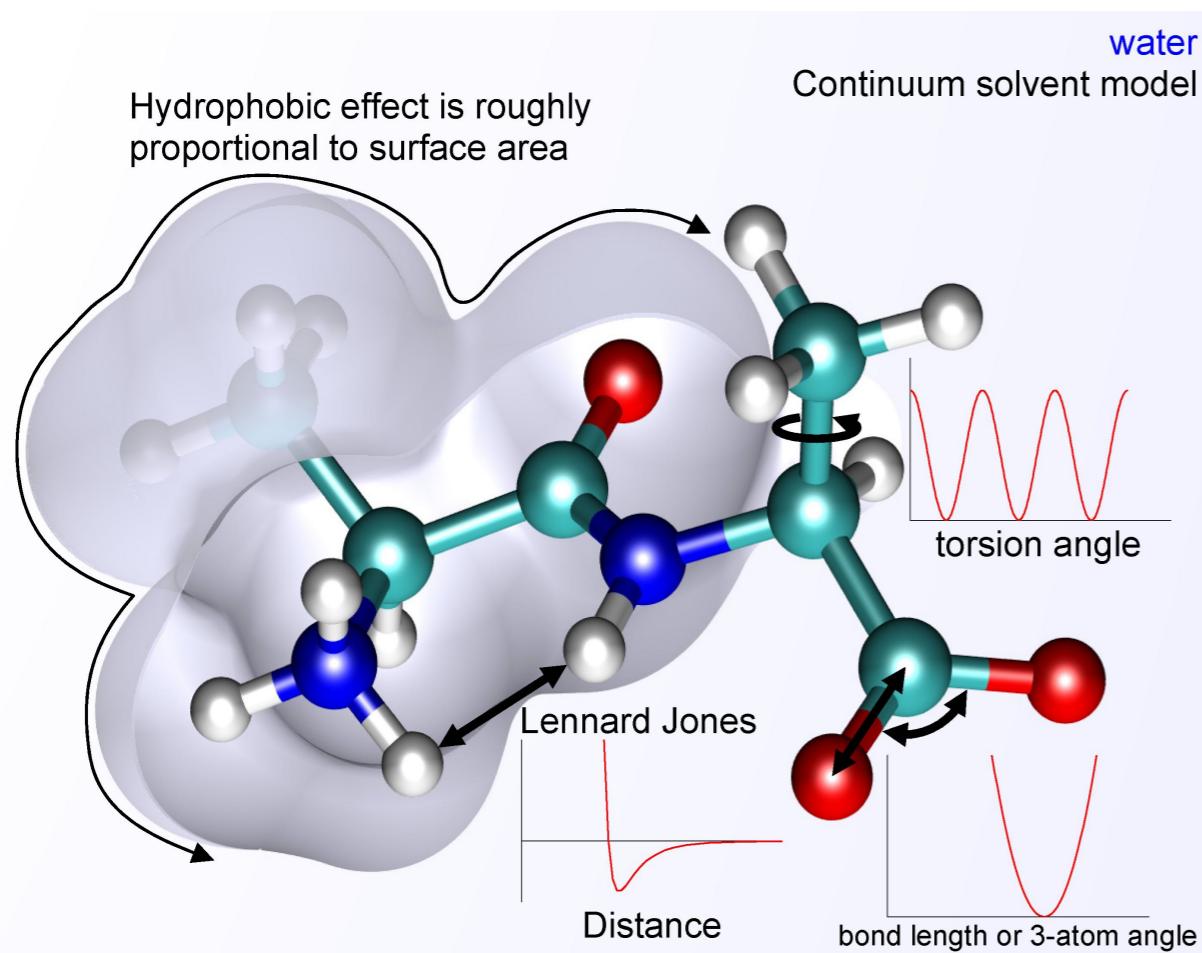
Limitations of molecular simulation

Two limitations in existing simulations are the approximations in the potential energy functions and the lengths of the simulations. The first introduces systematic errors and the second statistical errors.

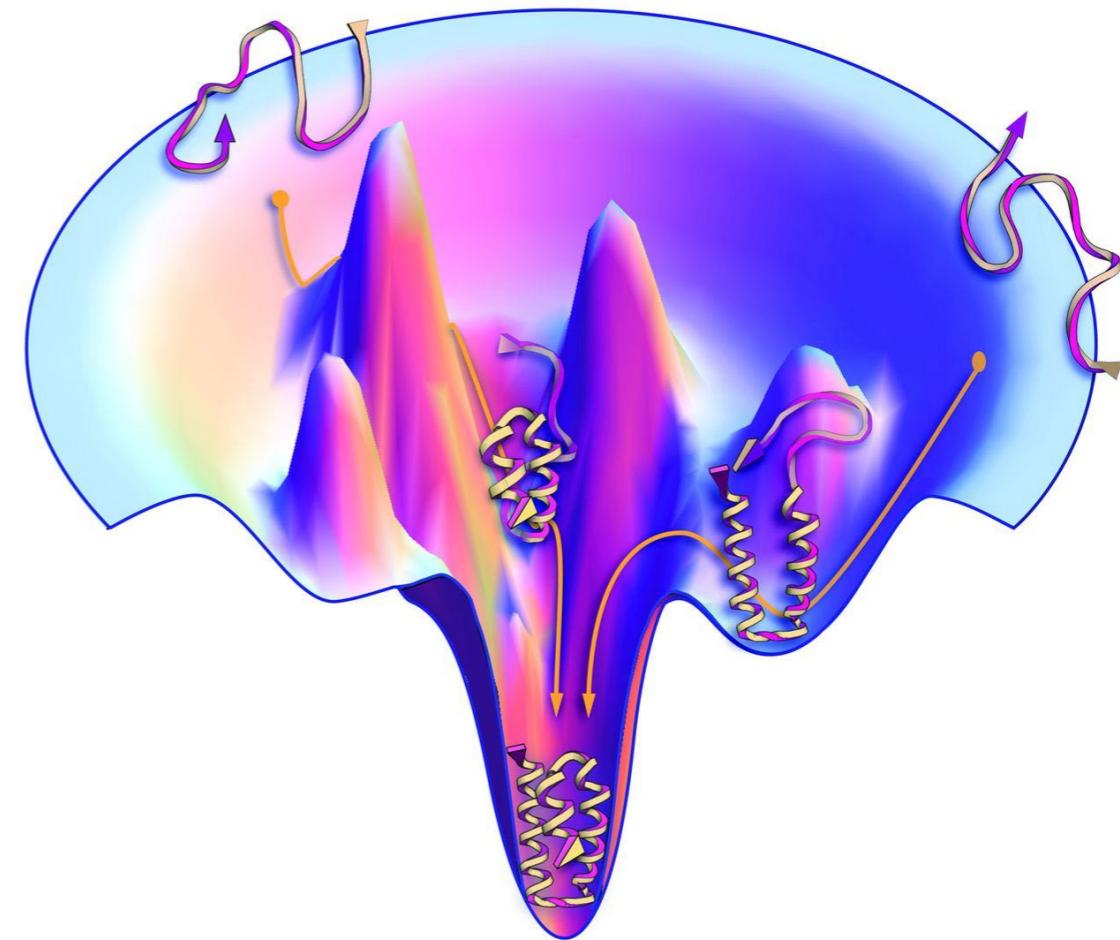
— M. Karplus & G. Petsko *Nature* (1990)



1. Accurate force fields (systematic errors)



2. Sampling configurational space (statistical errors)



Enhanced sampling

Tempering

Simulated annealing
Multicanonical algorithm
Replica exchange
Hamiltonian exchange
Parallel tempering
...

Collective variable bias

Umbrella sampling
Hyperdynamics
Metadynamics
Adaptive force biasing
Wang-Landau
...

Path sampling

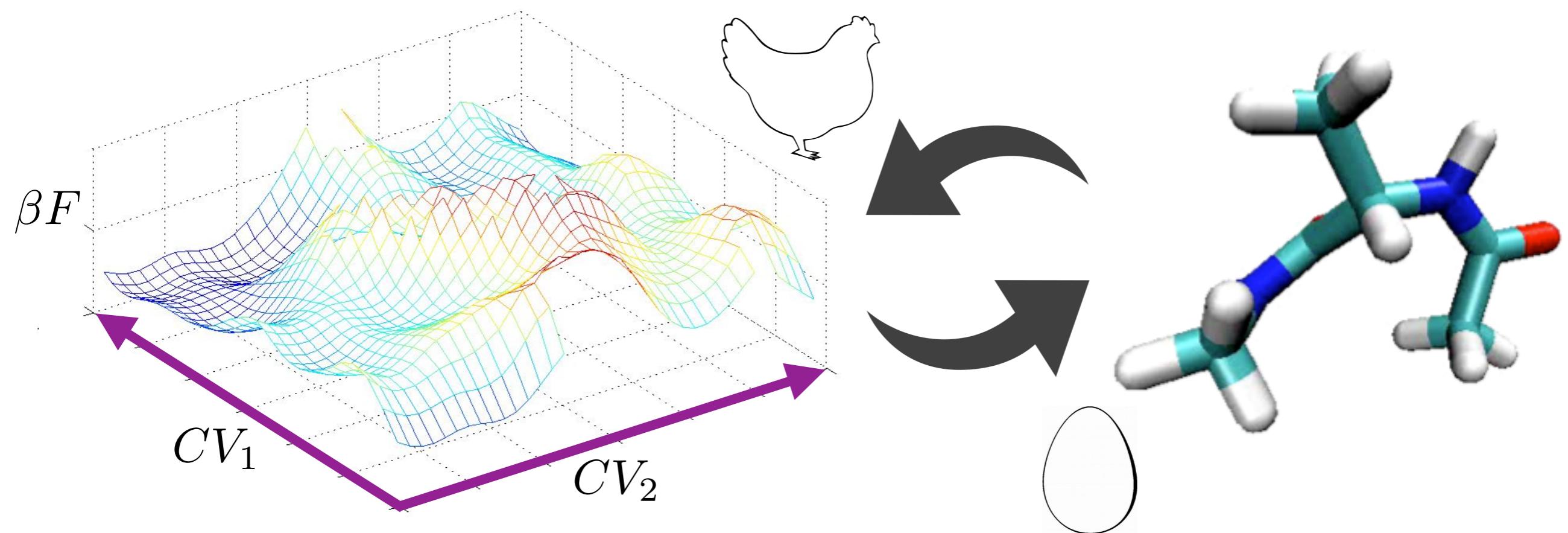
Transition path sampling
Transition interface sampling
Milestoning
Forward flux sampling
Weighted ensemble
...

- **Tempering** modifies T or Hamiltonian to accelerate barrier crossing
→ substantial CPU time expended on conditions not of direct interest
- **CV biasing** efficiently directs sampling along relevant order parameters
→ presupposes *a priori* availability of “good” CVs
- **Path sampling** efficiently simulates reactive paths between states
→ requires *a priori* knowledge of metastable states and connectivity

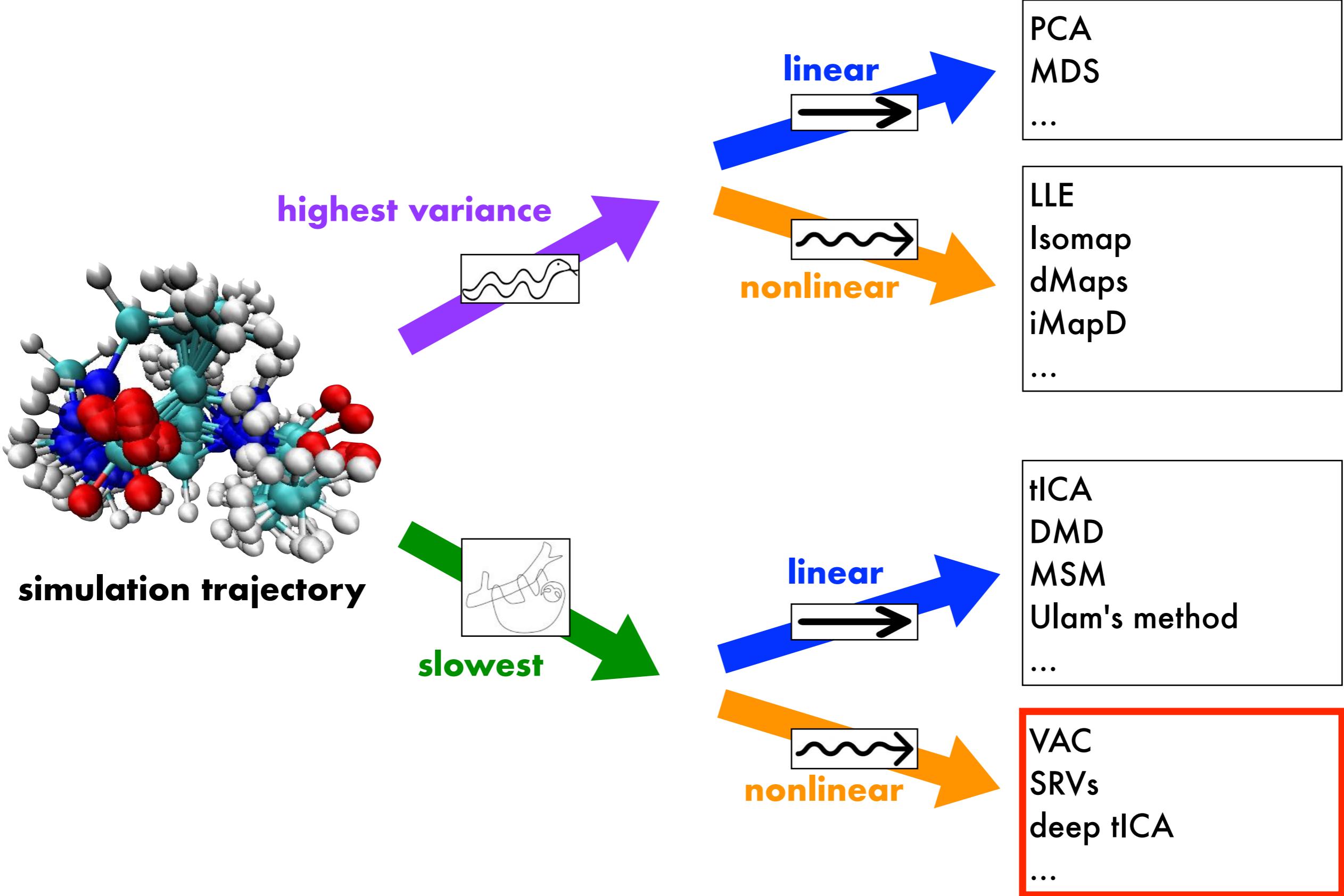
CV discovery + enhanced sampling

- Challenging to intuit CVs for all but the most trivial systems
- Data-driven CV discovery and enhanced sampling is typically **iterative**

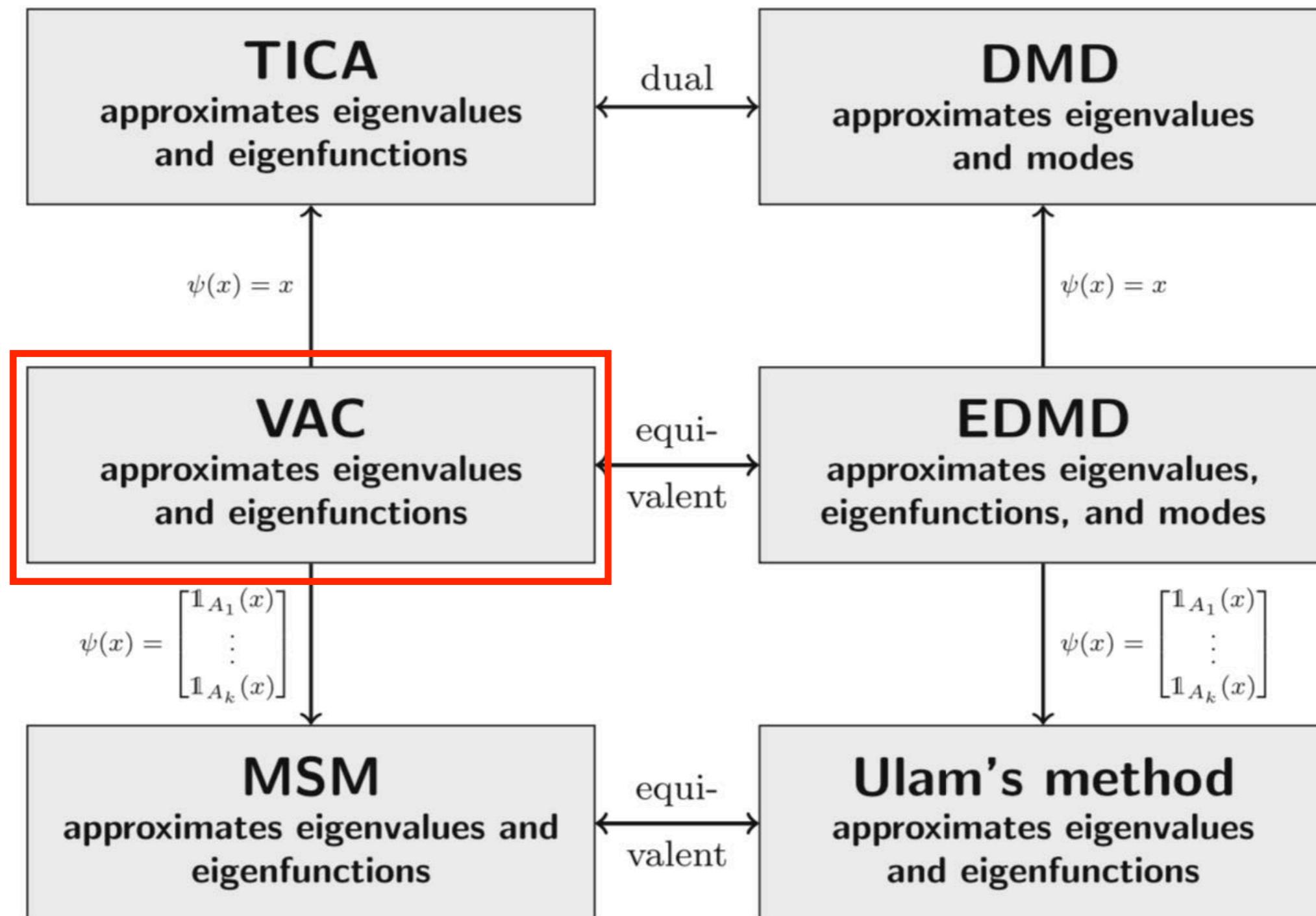
Good CVs required to drive sampling of configurational space (chicken)
Trajectories with good sampling needed to discover good CVs (egg)



CV discovery + enhanced sampling



Inference of slow CVs



Slow CVs: Transfer operator theory

- Dynamical evolution of a system is governed by the **transfer operator** that propagates probability densities through time in discrete steps of size τ

$$u_{t+\tau}(x) = \mathcal{T} \circ u_t(x)$$

transfer or Perron-Frobenius operator on densities
(self-adjoint to Koopman operator on observables)

probability distribution over state space x
at time t scaled by equilibrium distribution
 $u_t(x) = p_t(x)/\pi(x)$

- For equilibrium systems \mathcal{T} is **self-adjoint** \Rightarrow possesses an orthonormal basis of real eigenfunctions and eigenvalues

$$\mathcal{T} \circ \psi_i(x) = \lambda_i \psi_i(x) \quad \langle \psi_i | \psi_j \rangle_\pi = \delta_{ij} \quad 1 = \lambda_0 > \lambda_1 \geq \lambda_2 \geq \dots$$

inner product wrt equilibrium distribution

Transfer operator

- Any state function $\chi(x)$ can be projected into this orthonormal basis

$$\chi_t(x) = \sum_i \langle \psi_i | \chi_t \rangle_\pi \psi_i(x)$$

↑ ↑
expansion coefficients basis functions

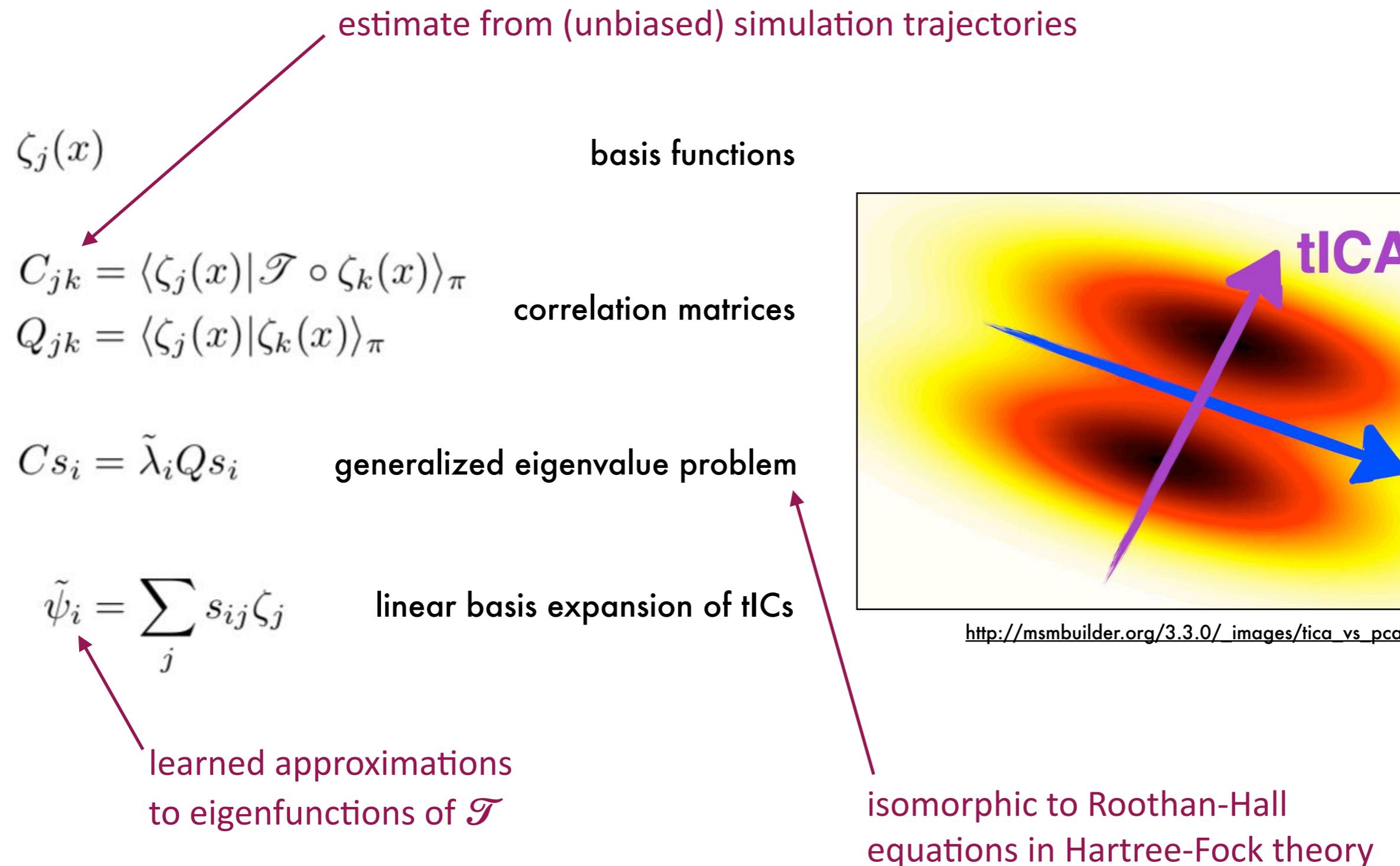
- The time evolution of $\chi(x)$ after k applications of \mathcal{T} becomes

$$\begin{aligned}\chi_{t+k\tau}(x) &= \mathcal{T}^k \circ \chi_t(x) = \sum_i \langle \psi_i | \chi_t \rangle_\pi \mathcal{T}^k \psi_i(x) \\ &= \sum_i \langle \psi_i | \chi_t \rangle_\pi \lambda_i^k \psi_i(x) \\ &= \sum_i \langle \psi_i | \chi_t \rangle_\pi \exp\left(-\frac{k\tau}{t_i}\right) \psi_i(x)\end{aligned}\quad \begin{array}{l} \text{implied timescale of } (\psi_i, \lambda_i) \\ \downarrow \\ t_i = -\frac{\tau}{\log \lambda_i} \end{array}$$

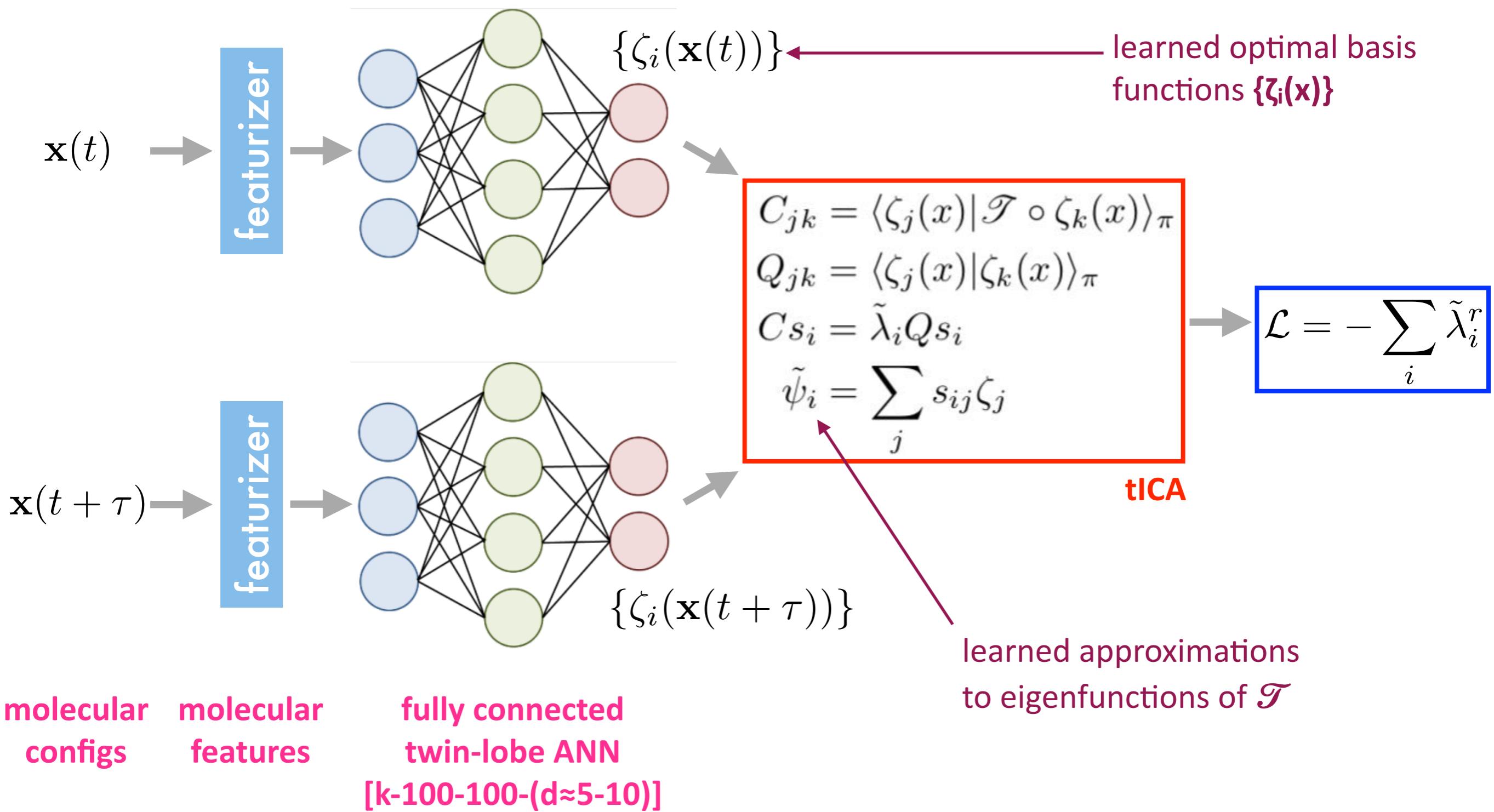
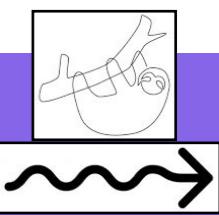
Leading eigenfunctions of \mathcal{T} are slowest dynamical modes



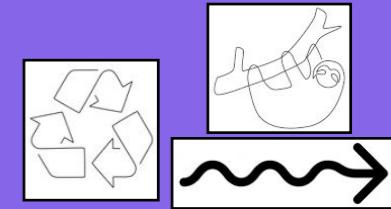
Time-lagged ICA (tICA)



State-free reversible VAMPnets (SRVs)

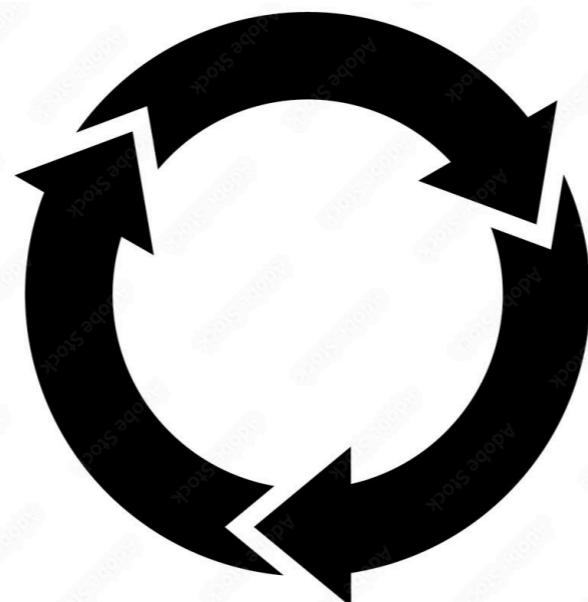


Iterative slow CV discovery and enhanced sampling?

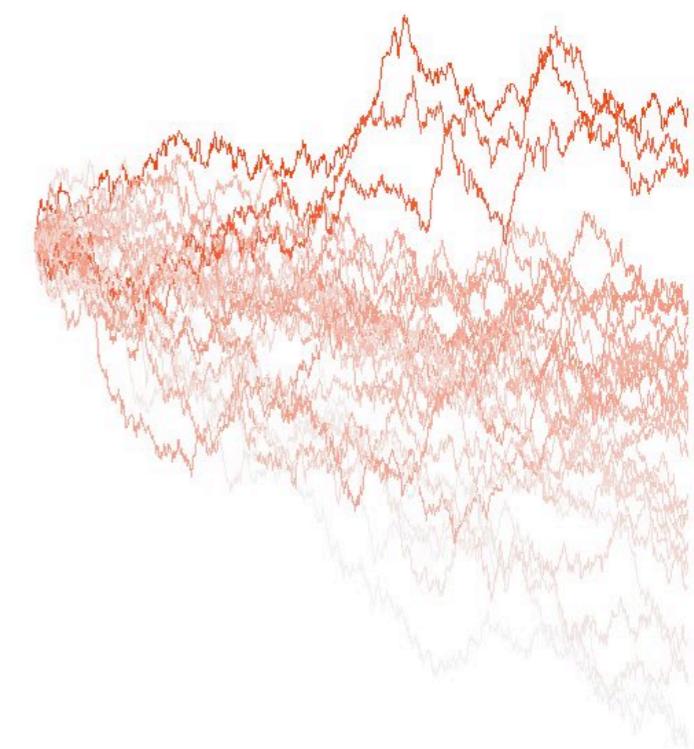


- Slow CVs are inherently dynamical variables estimated as path observables
- **Thermodynamic reweighting** of individual configurations is not enough, must also **dynamically reweight** the paths

enhanced sampling
 $\tilde{V}(x) = V(x) + U(\xi(x))$



thermodynamic and dynamical reweighting
 $W(\omega_{t \rightarrow t+\tau}) = g(x_t) \times M(\omega_{t \rightarrow t+\tau})$

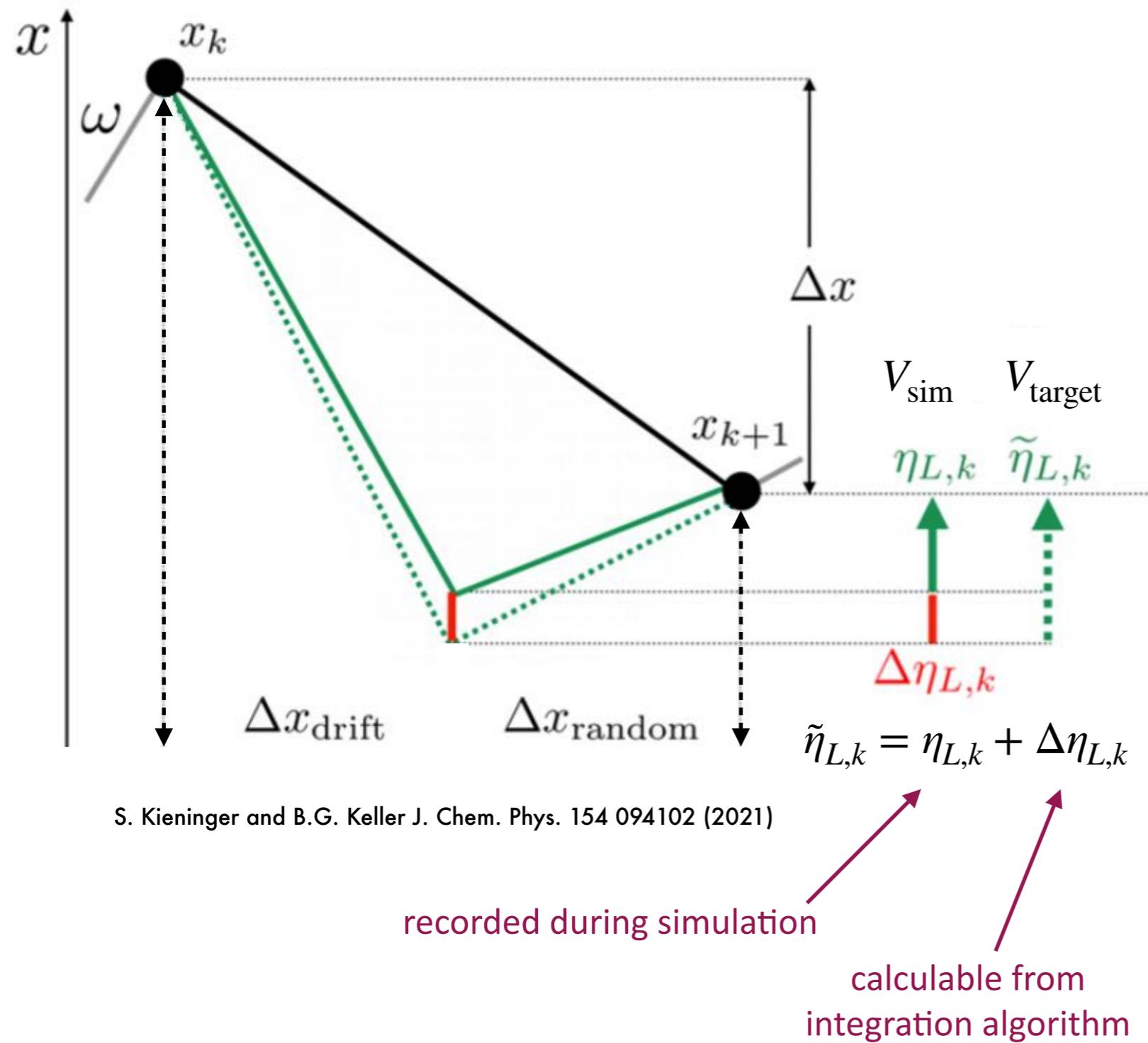


slow CV learning
 $\xi = \text{ML} (\omega_{t \rightarrow t+\tau}, W(\omega_{t \rightarrow t+\tau}))$

PYTORCH

Girsanov reweighting

- Elegant framework for dynamical reweighting of phase space paths under stochastic integrators (e.g., Langevin dynamics)



$$V_{\text{sim}} = V_{\text{target}} - U_{\text{bias}}$$

$$P_{\text{sim}}(\eta) = \frac{1}{(2\pi)^{n/2}} \exp\left(-\frac{1}{2} \sum_{k=0}^{n-1} \eta_k^2\right)$$

$$P_{\text{target}}(\tilde{\eta}) = \frac{1}{(2\pi)^{n/2}} \exp\left(-\frac{1}{2} \sum_{k=0}^{n-1} \tilde{\eta}_k^2\right)$$

$$M(\omega) = \frac{P_{\text{target}}(\tilde{\eta})}{P_{\text{sim}}(\eta)}$$

$$= \exp\left(-\frac{1}{2} \sum_{k=0}^{n-1} \eta_k \cdot \Delta\eta_k\right)$$

$$\times \exp\left(-\frac{1}{2} \sum_{k=0}^{n-1} \eta_k^2\right)$$

Girsanov reweighting

- Exact expressions known for
 - Brownian dynamics (overdamped Langevin) by Euler-Maruyama
 - (underdamped) Langevin dynamics by Izaguirre, Sweet, and Pande (ISP)
- Applied by Weber & Pande and Keller & co. to estimate MSMs from biased simulations collected under pre-defined CV

Brownian dynamics (E-M)

$$M_o(\omega_o, \eta_o; \Delta t | x_0) = \exp\left(-\sum_{k=0}^{n-1} \sqrt{\frac{\Delta t}{2k_B T \xi m}} \nabla U(x_k) \cdot \eta_{o,k}\right) \cdot \exp\left(-\frac{1}{2} \sum_{k=0}^{n-1} \frac{\Delta t}{2k_B T \xi m} (\nabla U(x_k))^2\right).$$

Langevin dynamics (ISP)

$$M_L(\omega_L, \eta_L; \Delta t | (x_0, v_0)) = \exp\left(-\frac{1 - \exp(-\xi \Delta t)}{\sqrt{1 - \exp(-2\xi \Delta t)}} \cdot \frac{\sum_{k=0}^{n-1} \nabla U(x_k) \eta_{L,k}}{\sqrt{k_B T \xi^2 m}}\right) \times \exp\left(-\frac{(1 - \exp(-\xi \Delta t))^2}{1 - \exp(-2\xi \Delta t)} \cdot \frac{\sum_{k=0}^{n-1} \nabla U^2(x_k)}{2k_B T \xi^2 m}\right).$$

S. Kieninger and B.G. Keller J. Chem. Phys. 154 094102 (2021)

Girsanov reweighting within SRVs

$$C_{jk} = \langle \zeta_j(\mathbf{x}_t) | \zeta_k(\mathbf{x}_{t+\tau}) \rangle_{V_{\text{target}}}$$

$$\approx \frac{1}{L} \sum_{l=1}^L W(\omega_{t \rightarrow t+\tau}) \zeta_j(\mathbf{x}_t^{(l)}) \zeta_k(\mathbf{x}_{t+\tau}^{(l)})$$

$$Q_{jk} = \langle \zeta_j(\mathbf{x}_t) | \zeta_k(\mathbf{x}_t) \rangle_{V_{\text{target}}}$$

$$\approx \frac{1}{L} \sum_{l=1}^L W(\omega_{t \rightarrow t+\tau}) \zeta_j(\mathbf{x}_t^{(l)}) \zeta_k(\mathbf{x}_t^{(l)})$$

$$C_{S_i} = \tilde{\lambda}_i Q_{S_i}$$

$$\tilde{\psi}_i = \sum_j s_{ij} \zeta_j$$

learned optimal basis functions $\{\zeta_i(\mathbf{x})\}$

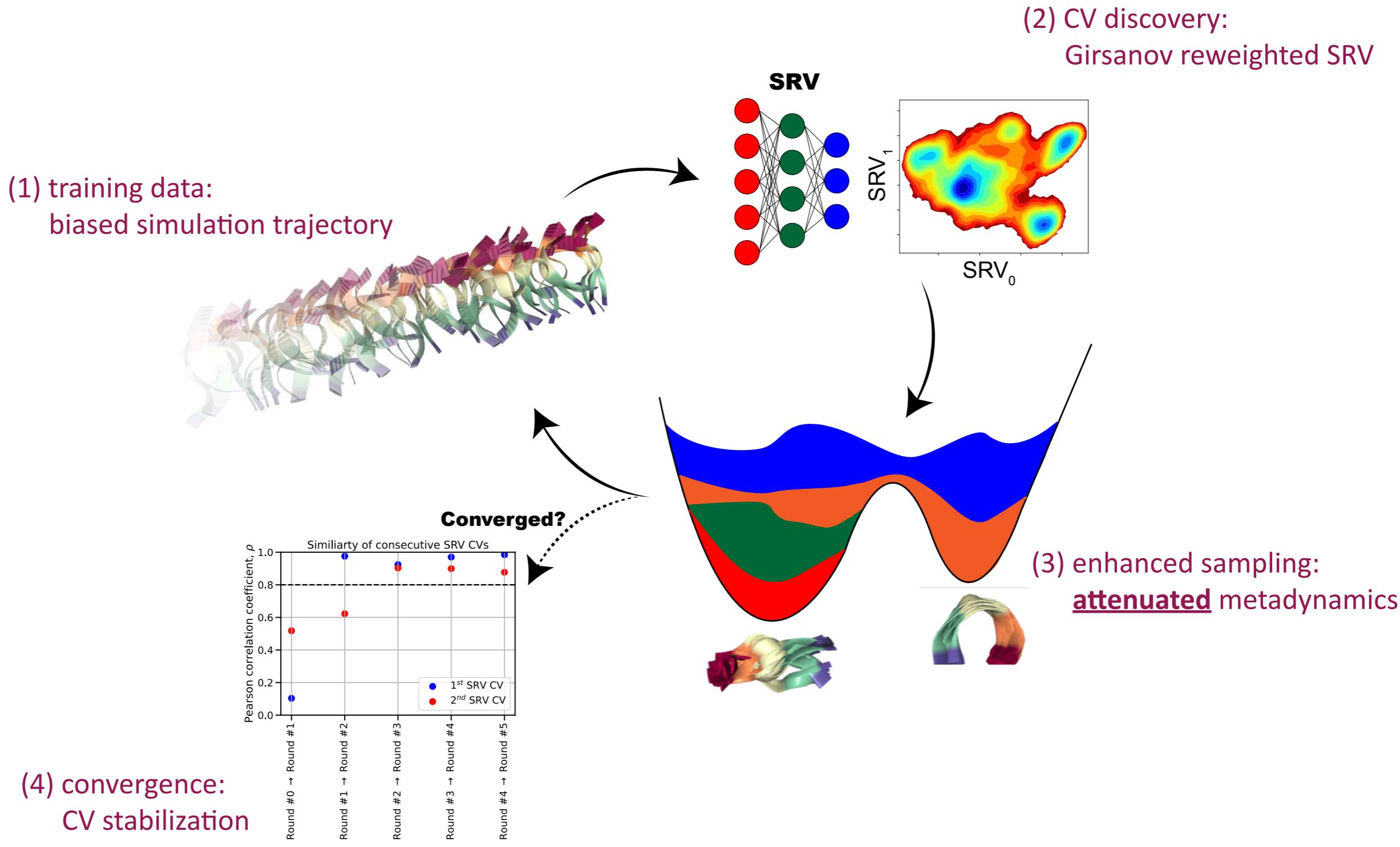
learned approximations to eigenfunctions of \mathcal{T}

$$W(\omega_{t \rightarrow t+\tau}) = g(\mathbf{x}_t) \times M(\omega_{t \rightarrow t+\tau})$$

$g(\mathbf{x}_t) = \exp(-\beta U_{\text{bias}}(\mathbf{x}_t))$
thermodynamic weighting of initial state

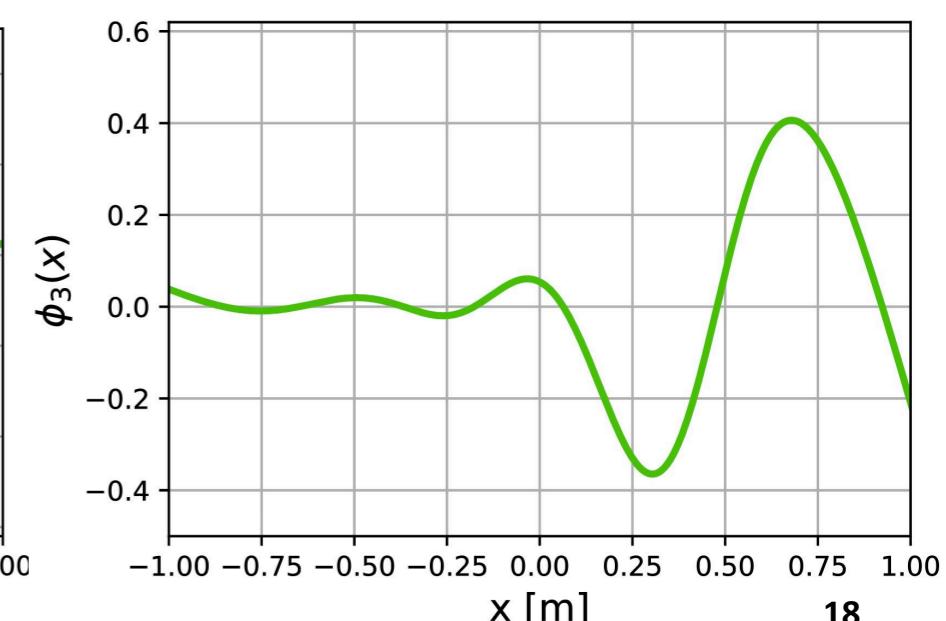
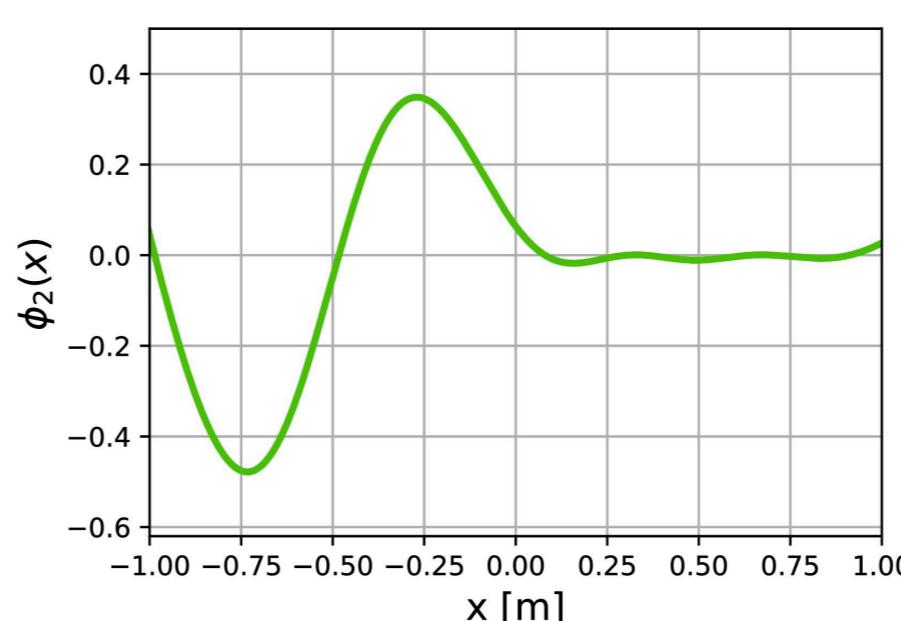
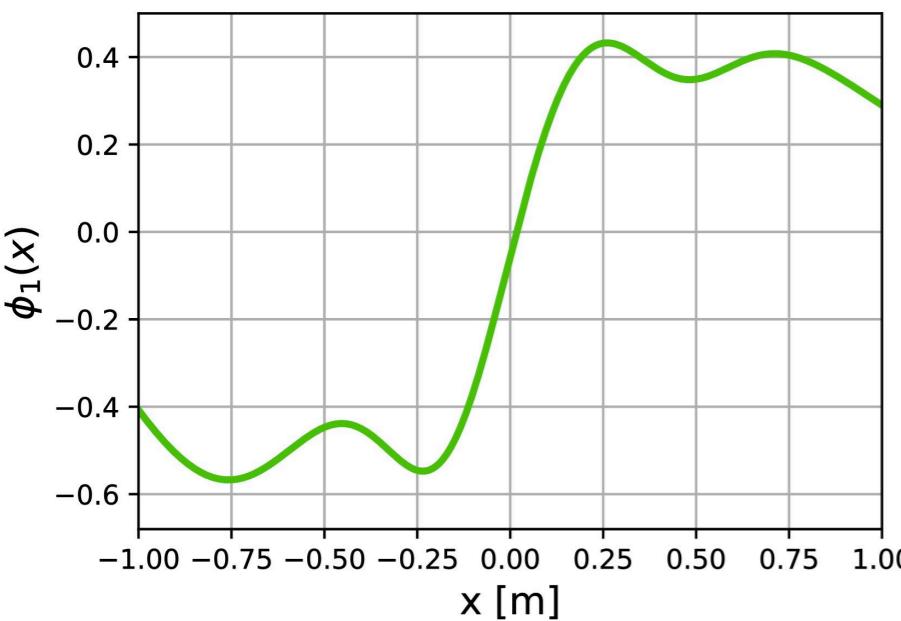
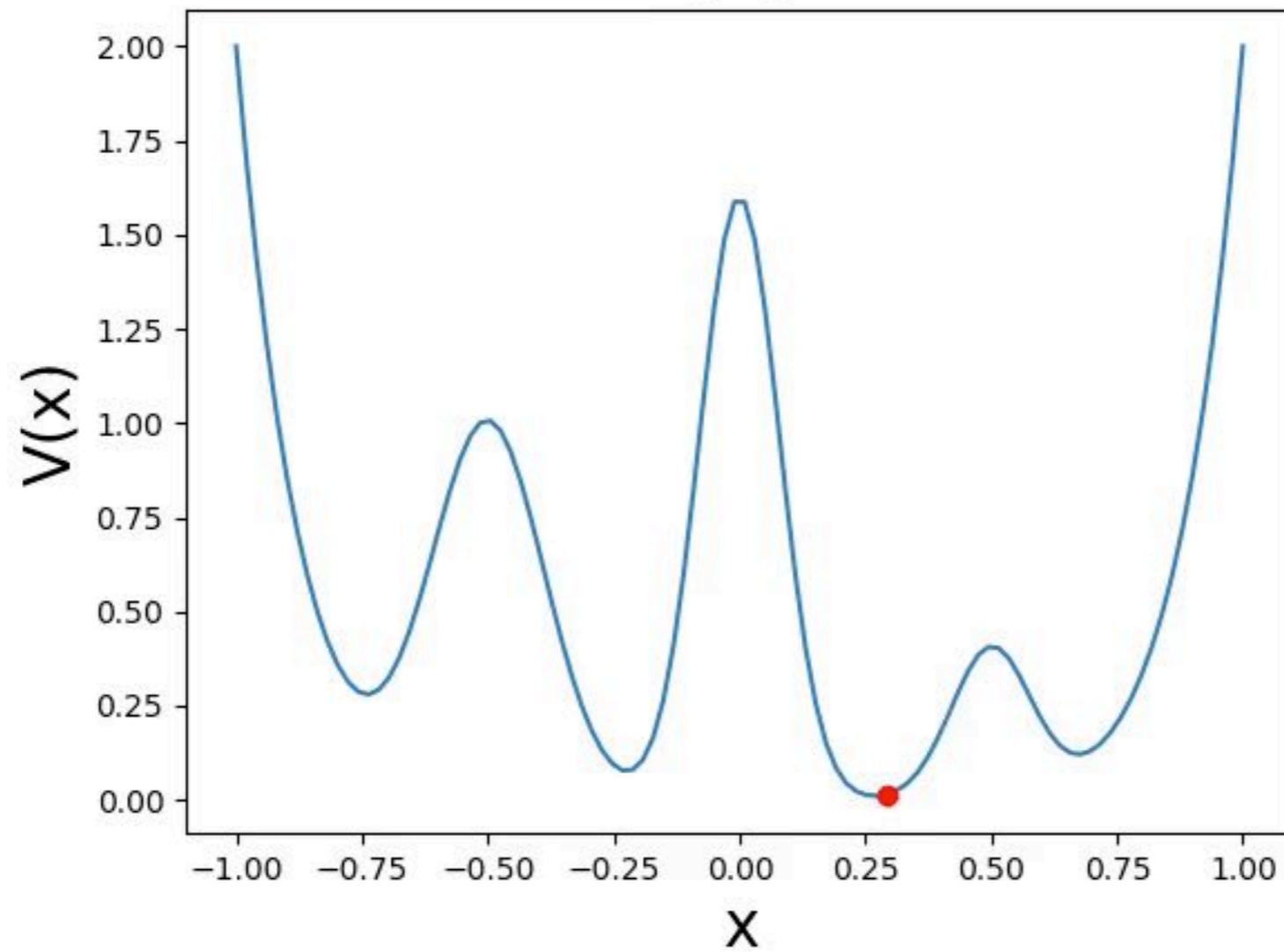
$M(\omega_{t \rightarrow t+\tau})$
Girsanov reweighting of phase space path

GREST: Girsanov Reweighting Enhanced Sampling Technique

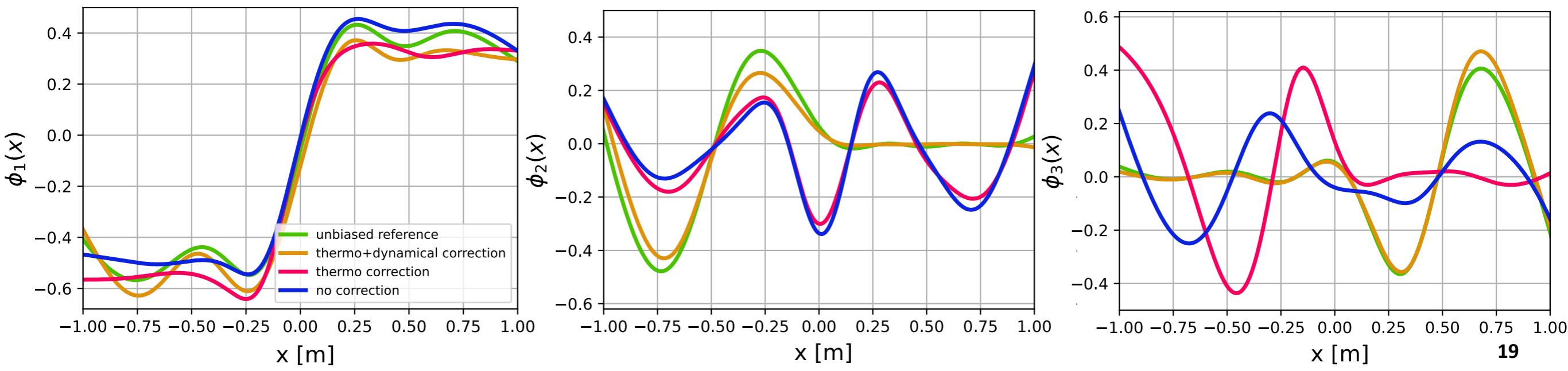
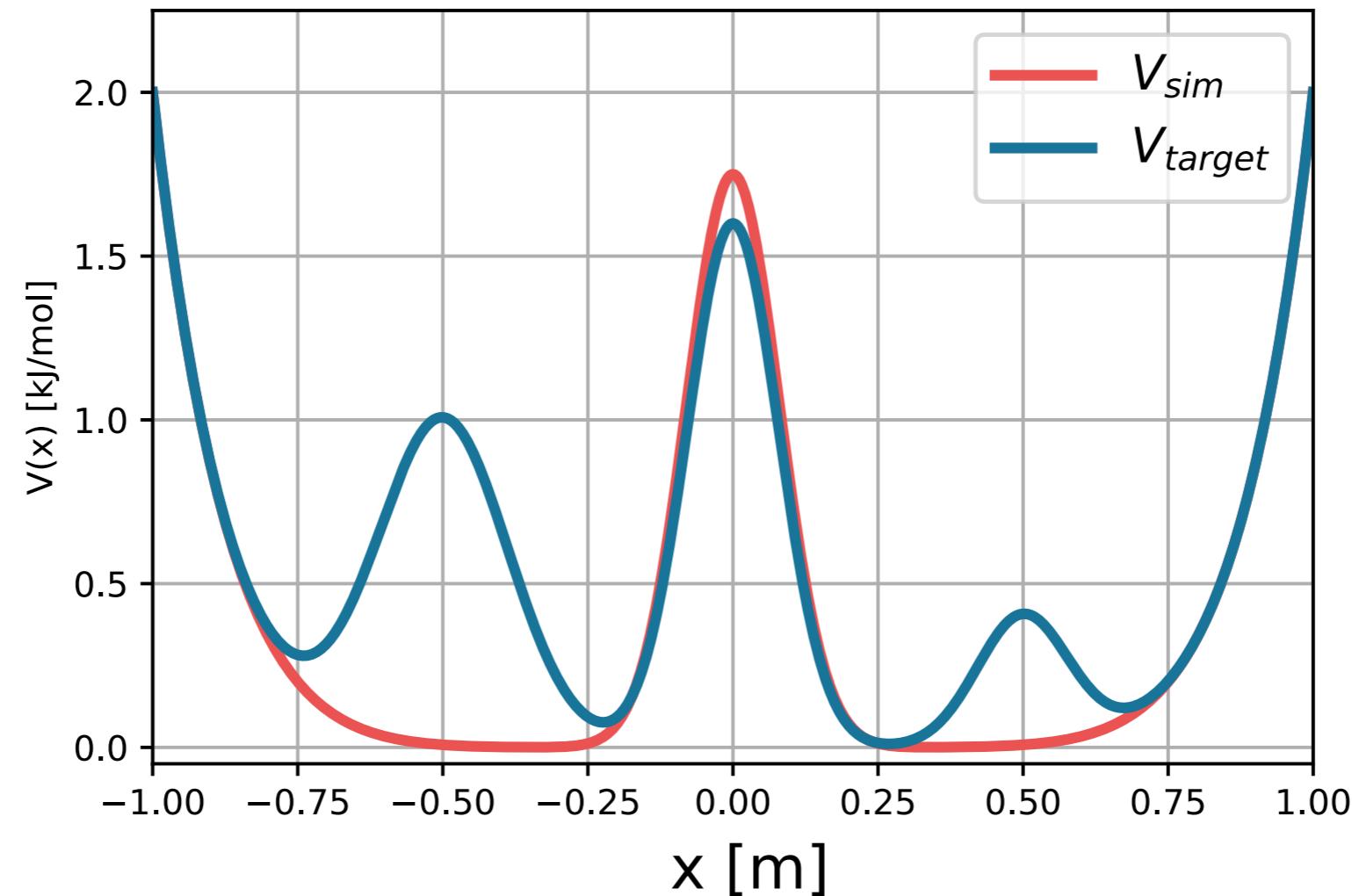


1D 4-well landscape

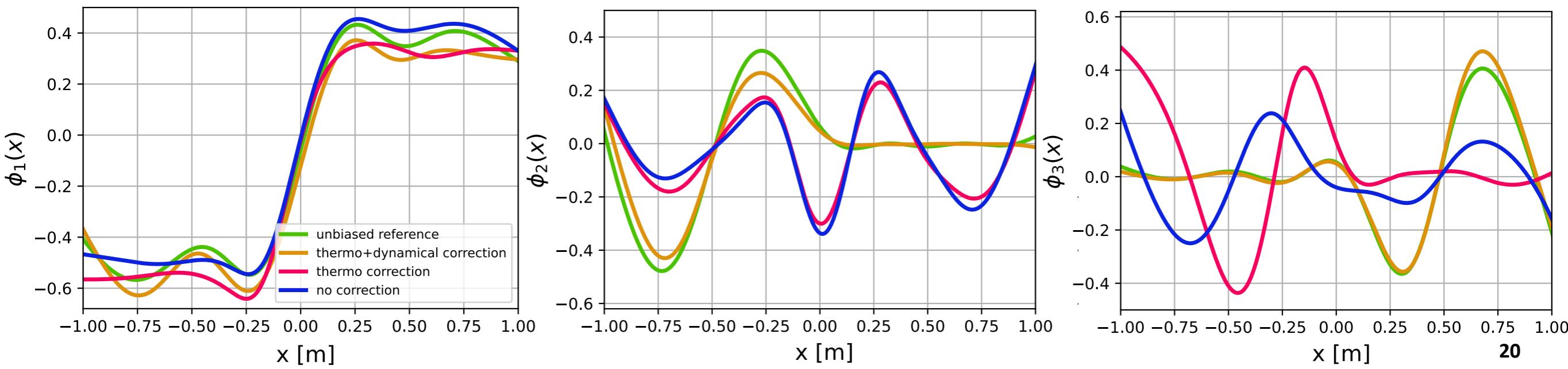
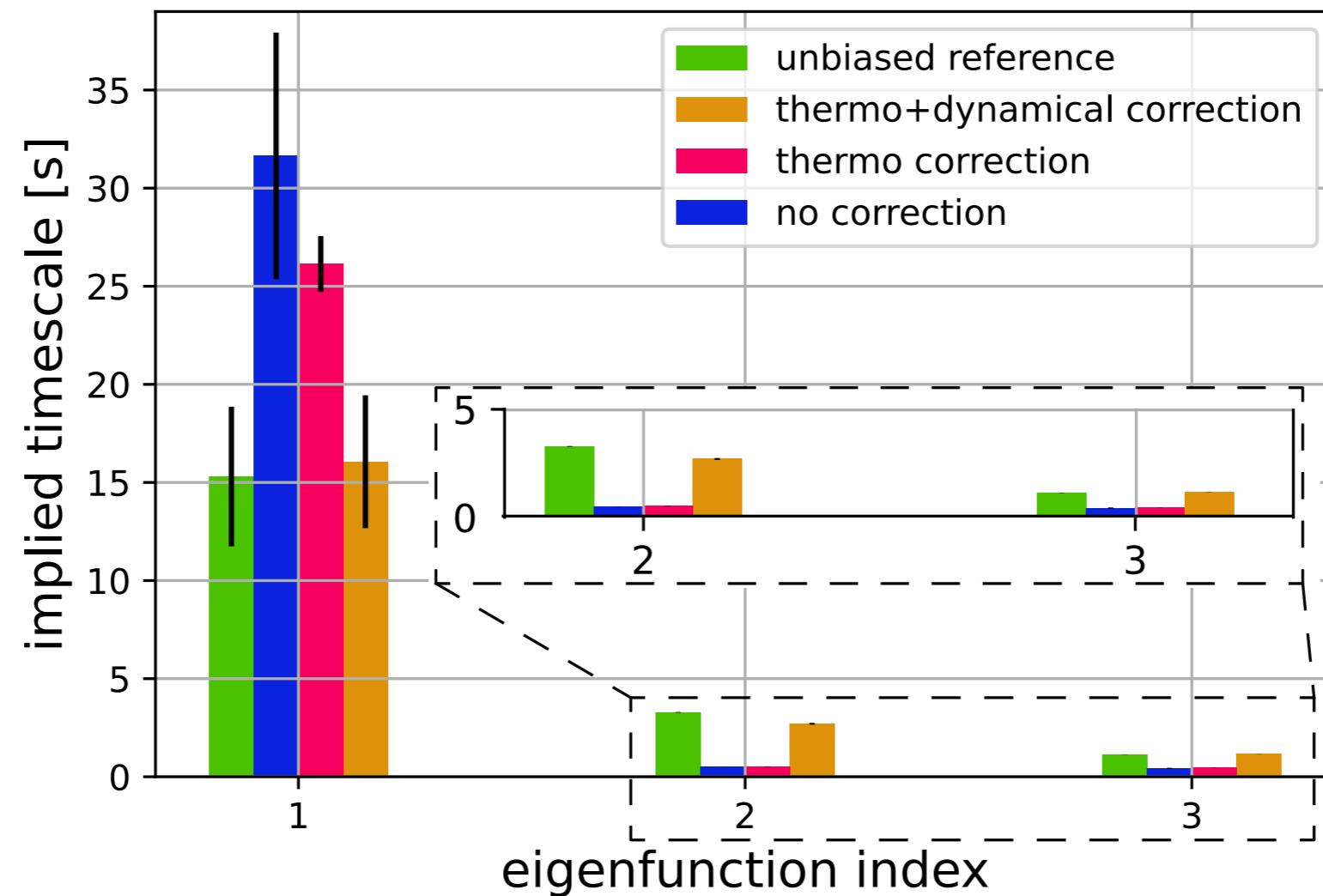
$t = 0$



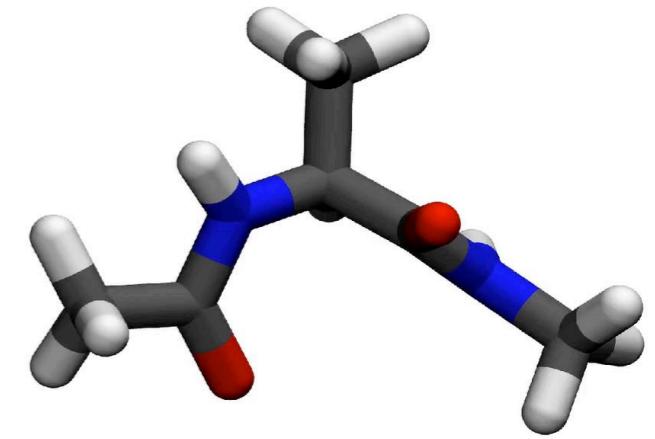
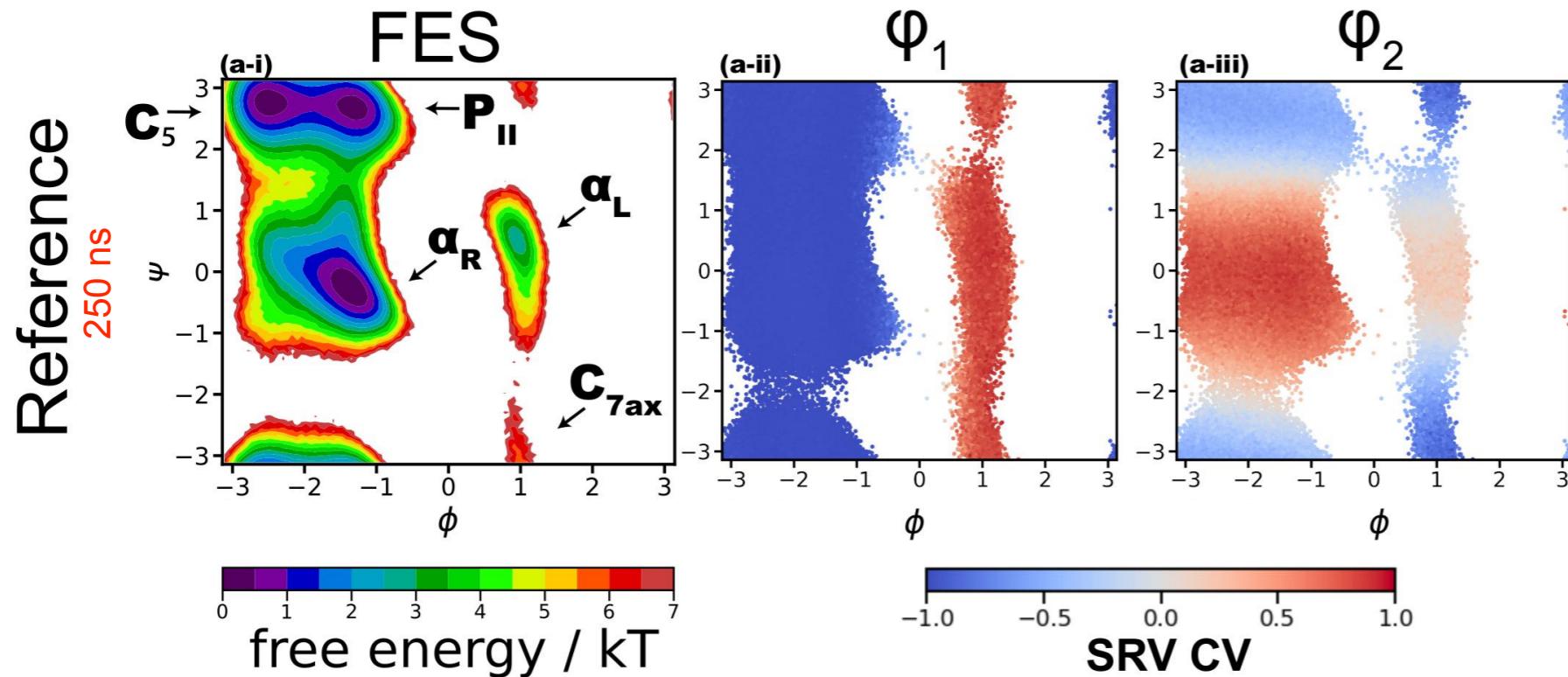
1D 4-well landscape



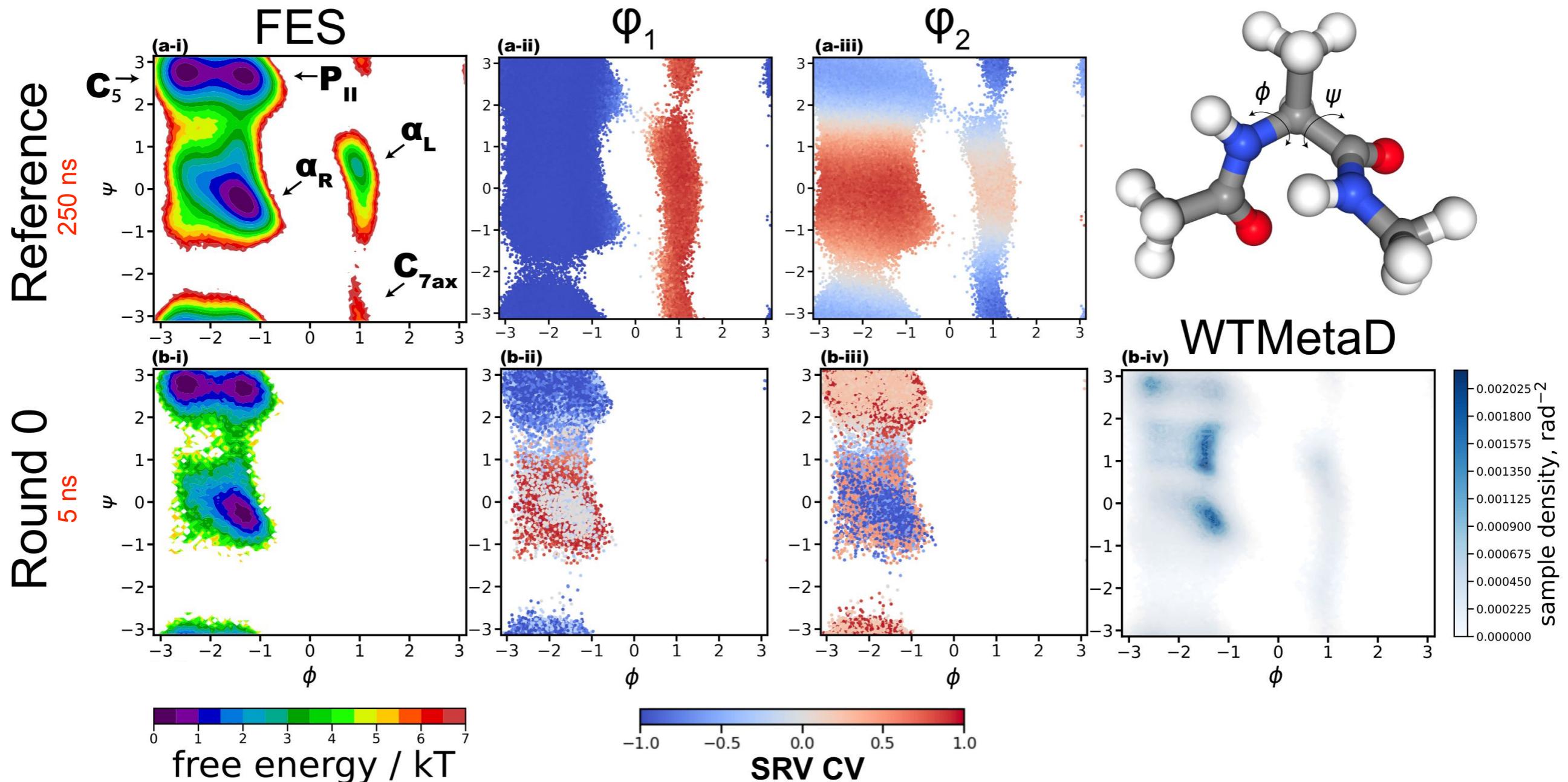
1D 4-well landscape



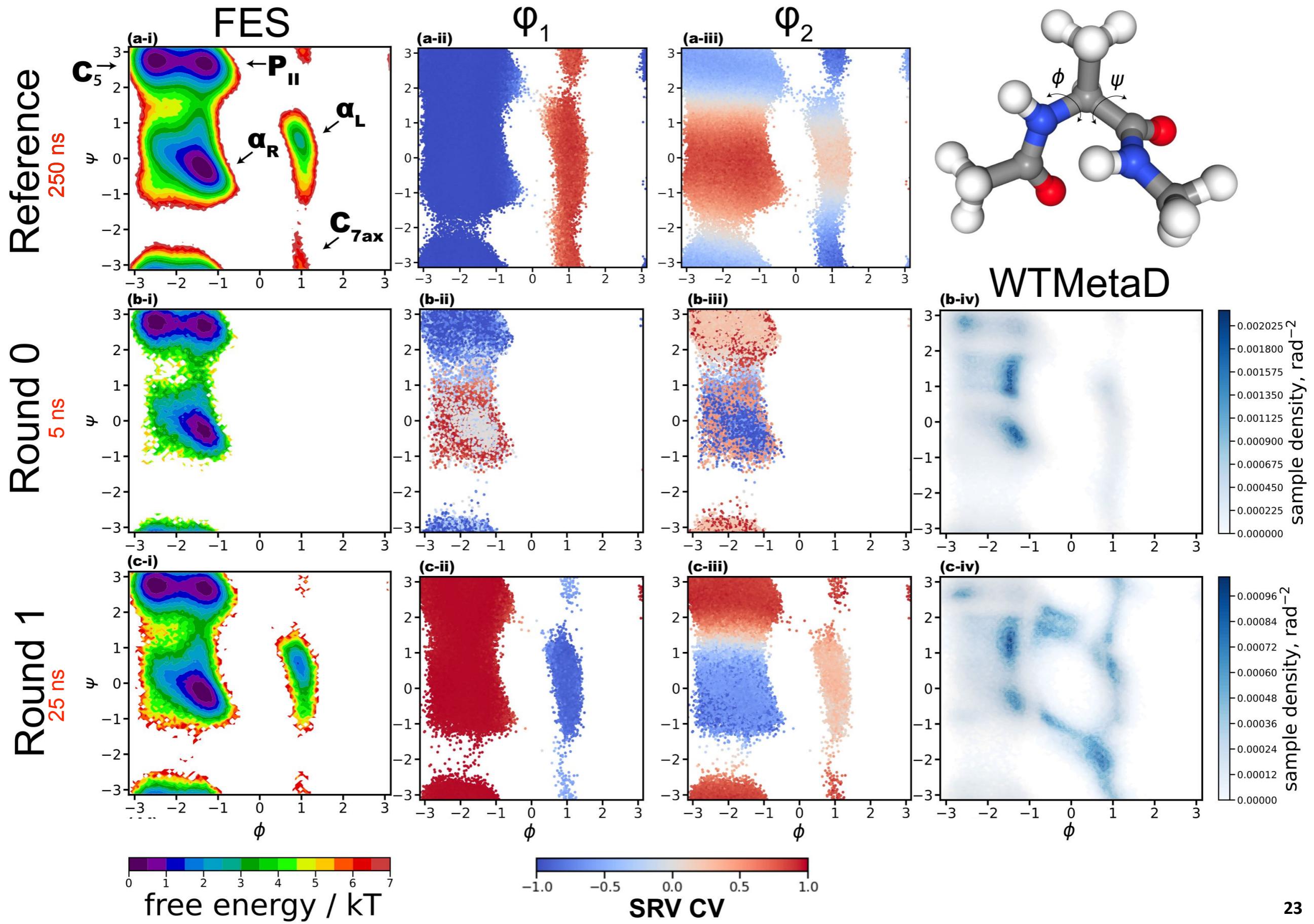
Alanine dipeptide



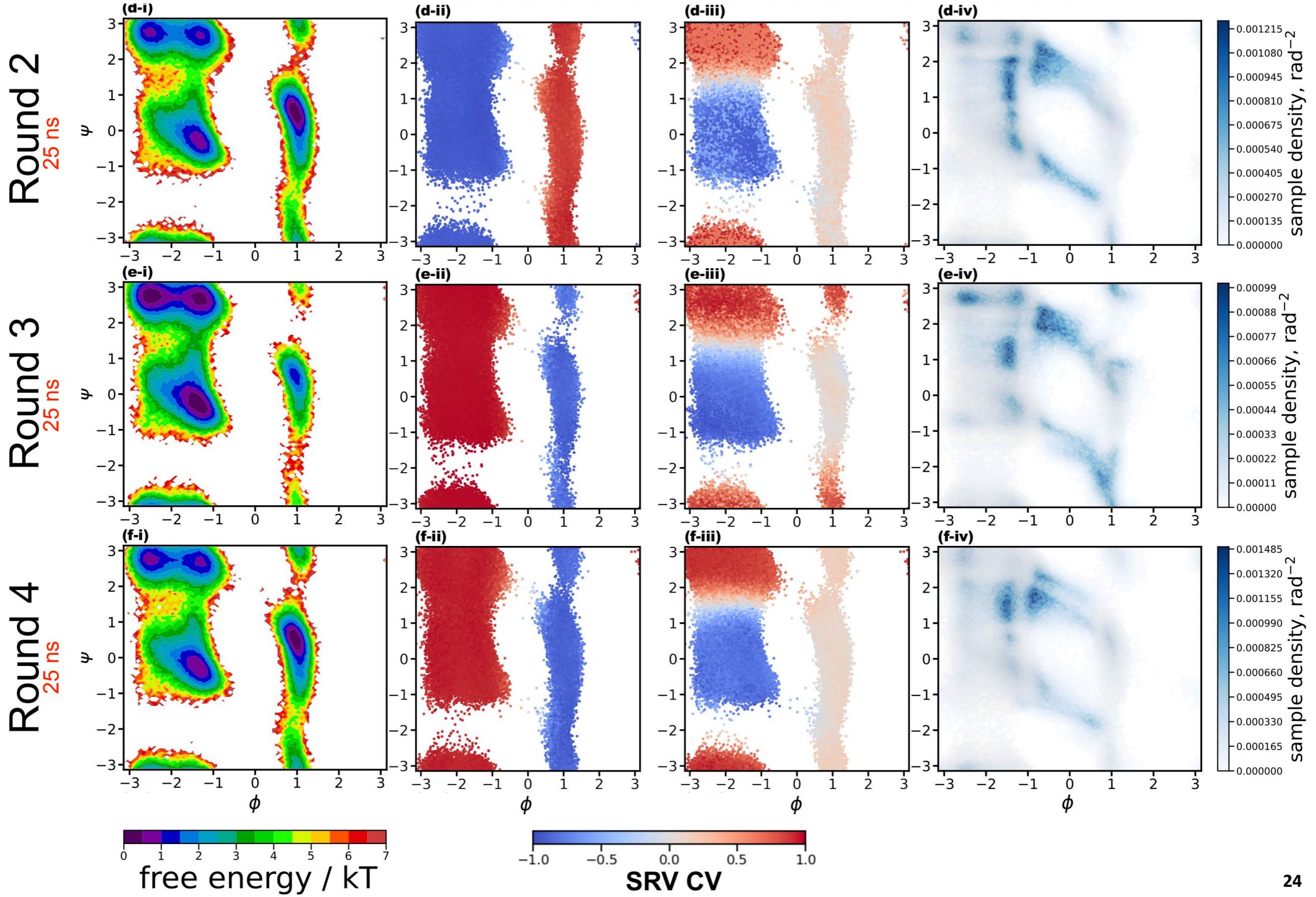
Alanine dipeptide



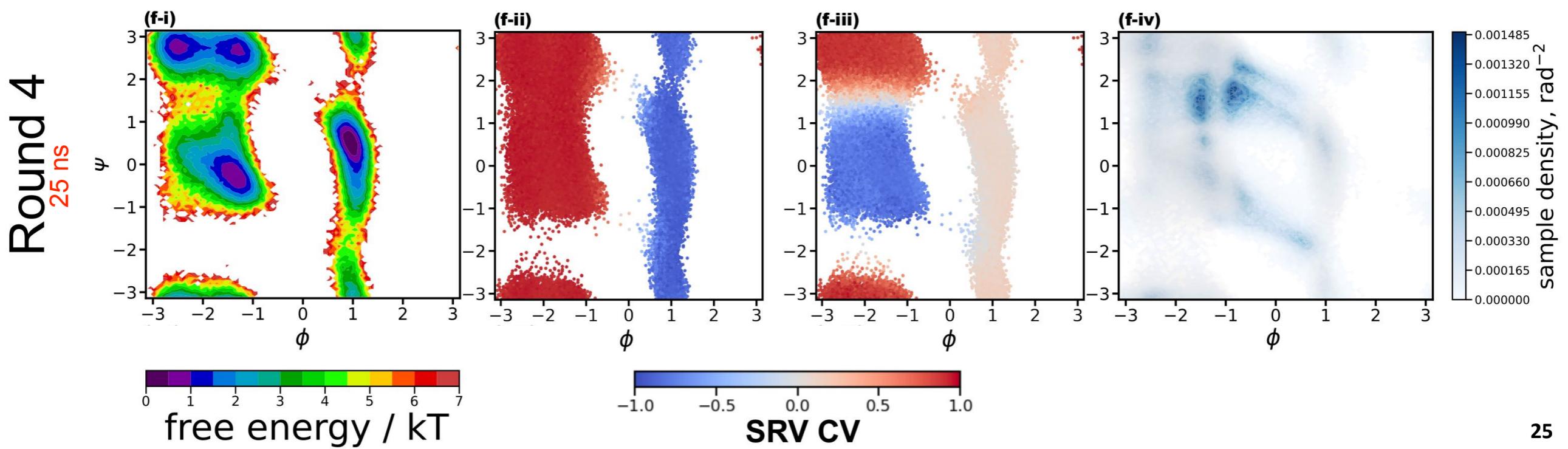
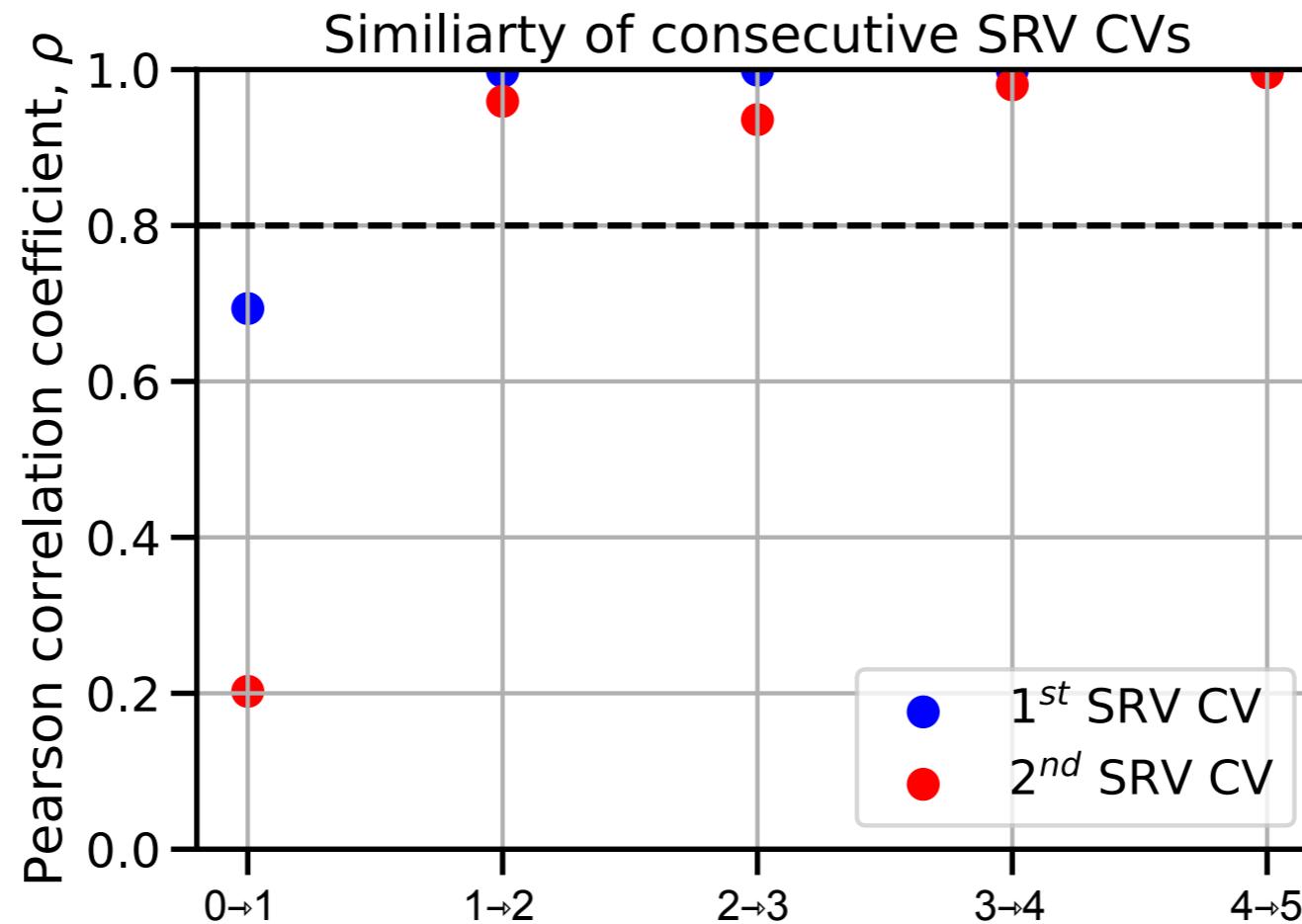
Alanine dipeptide



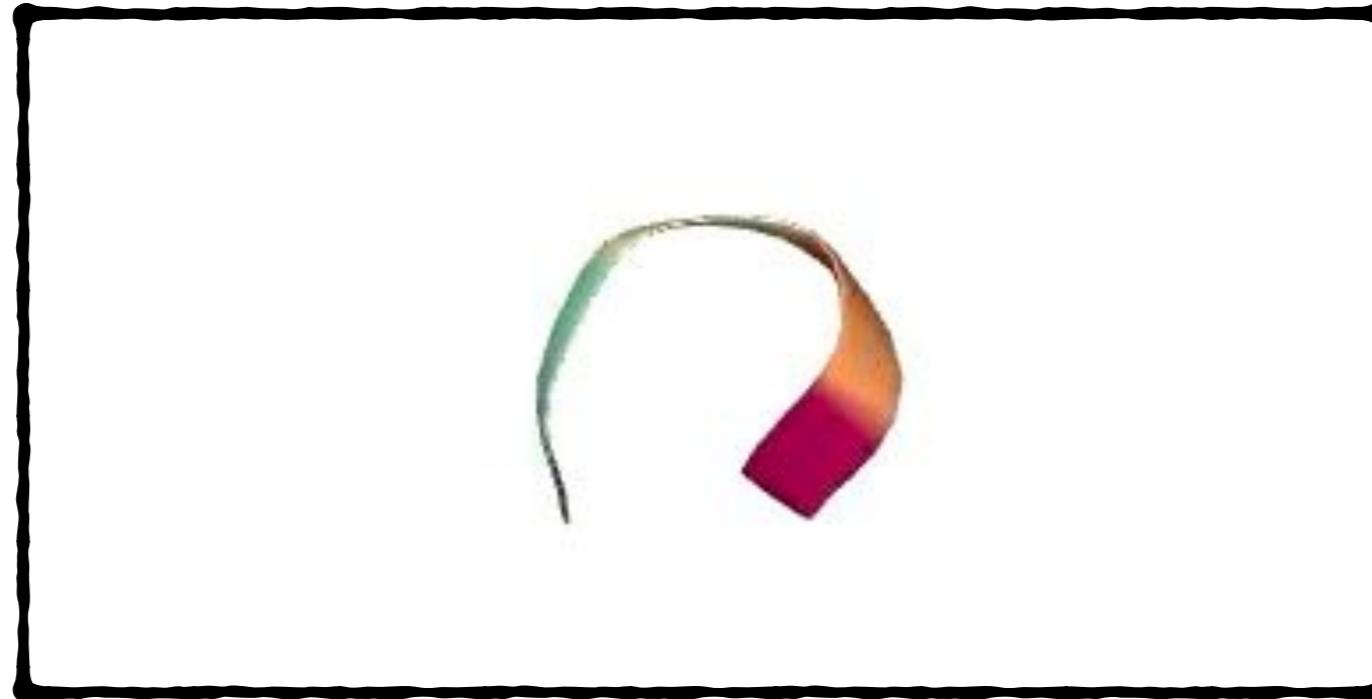
Alanine dipeptide



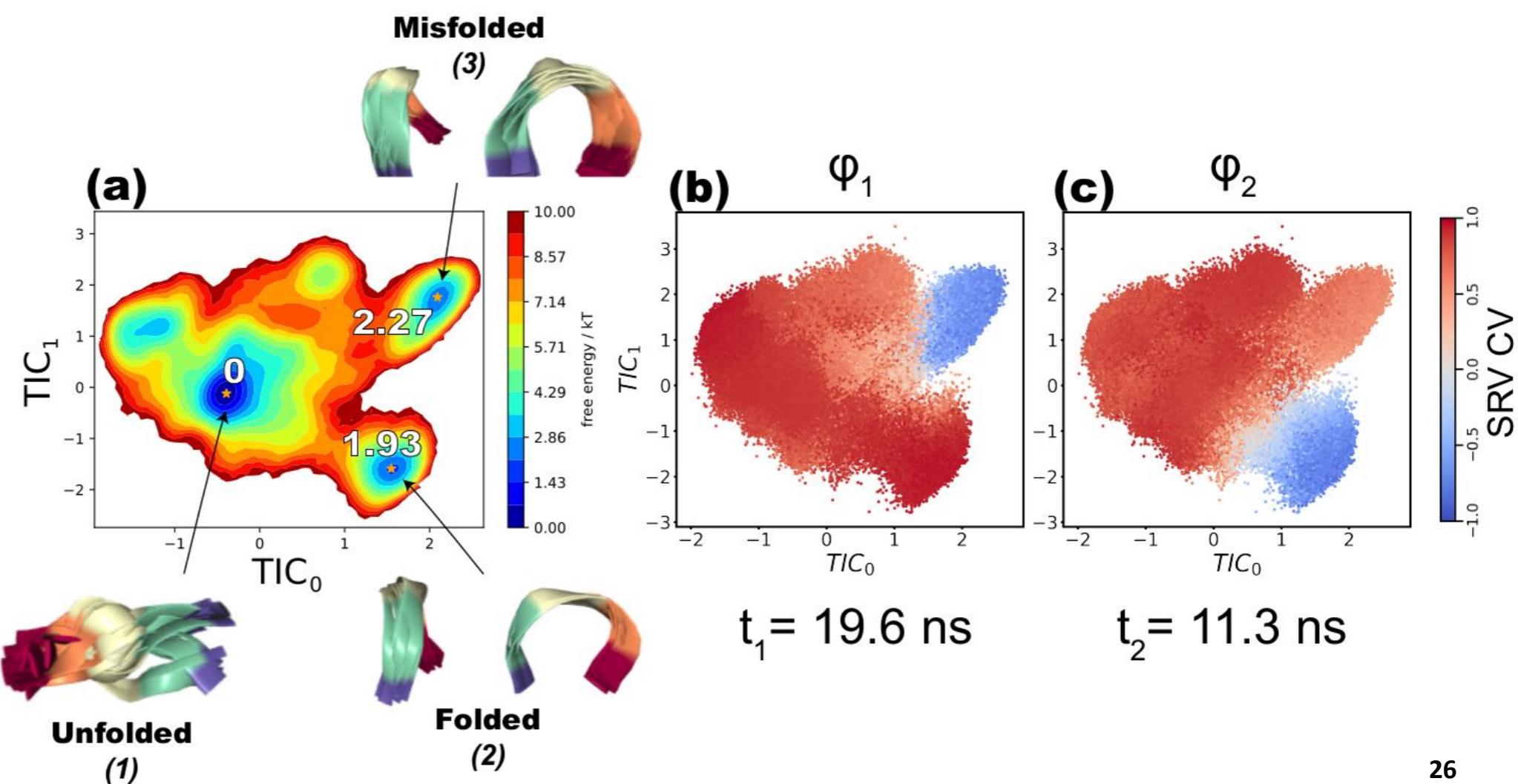
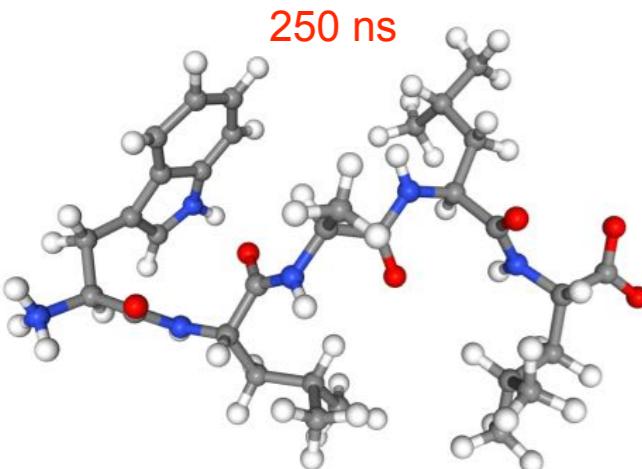
Alanine dipeptide



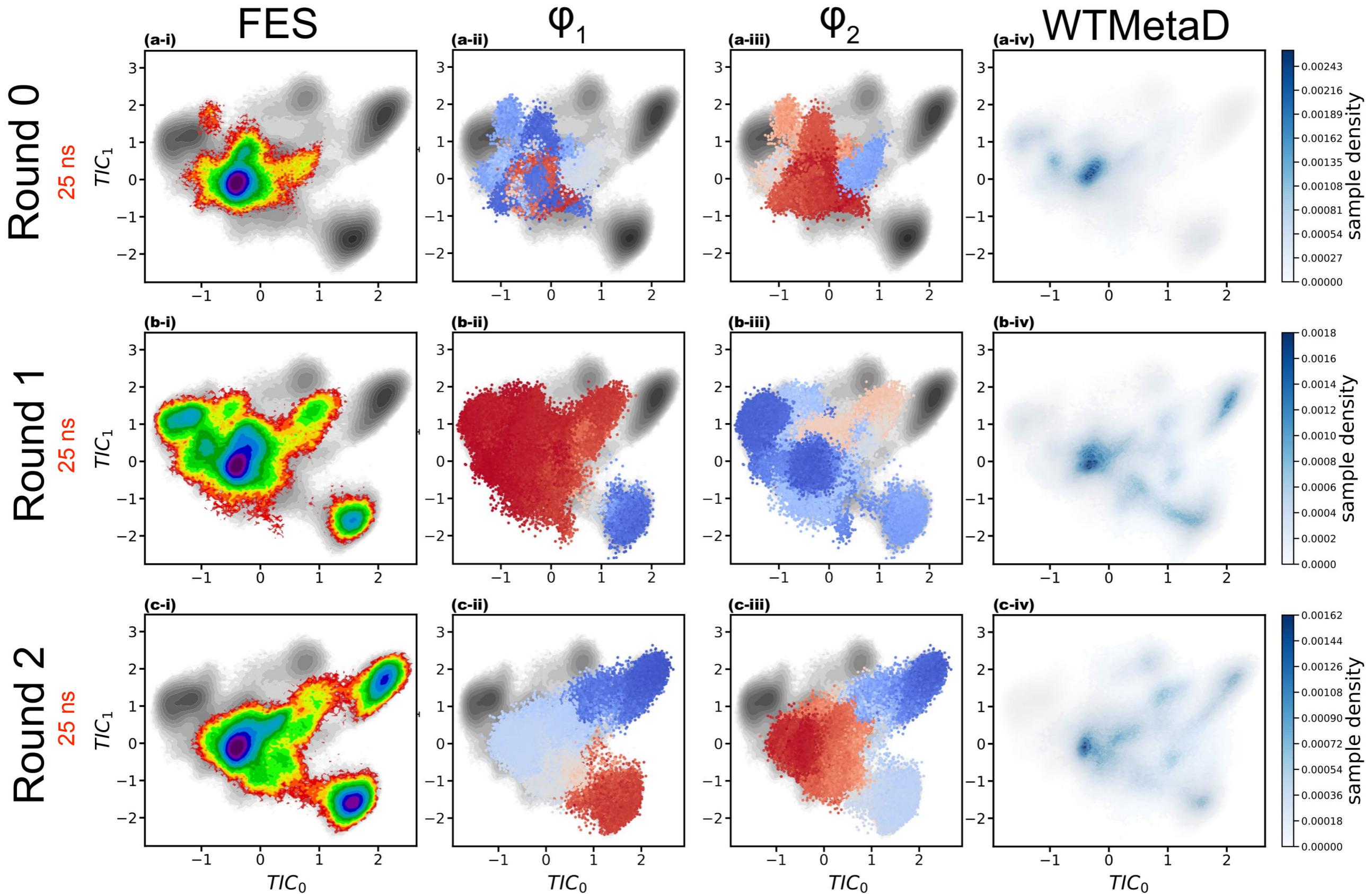
WLALL pentapeptide



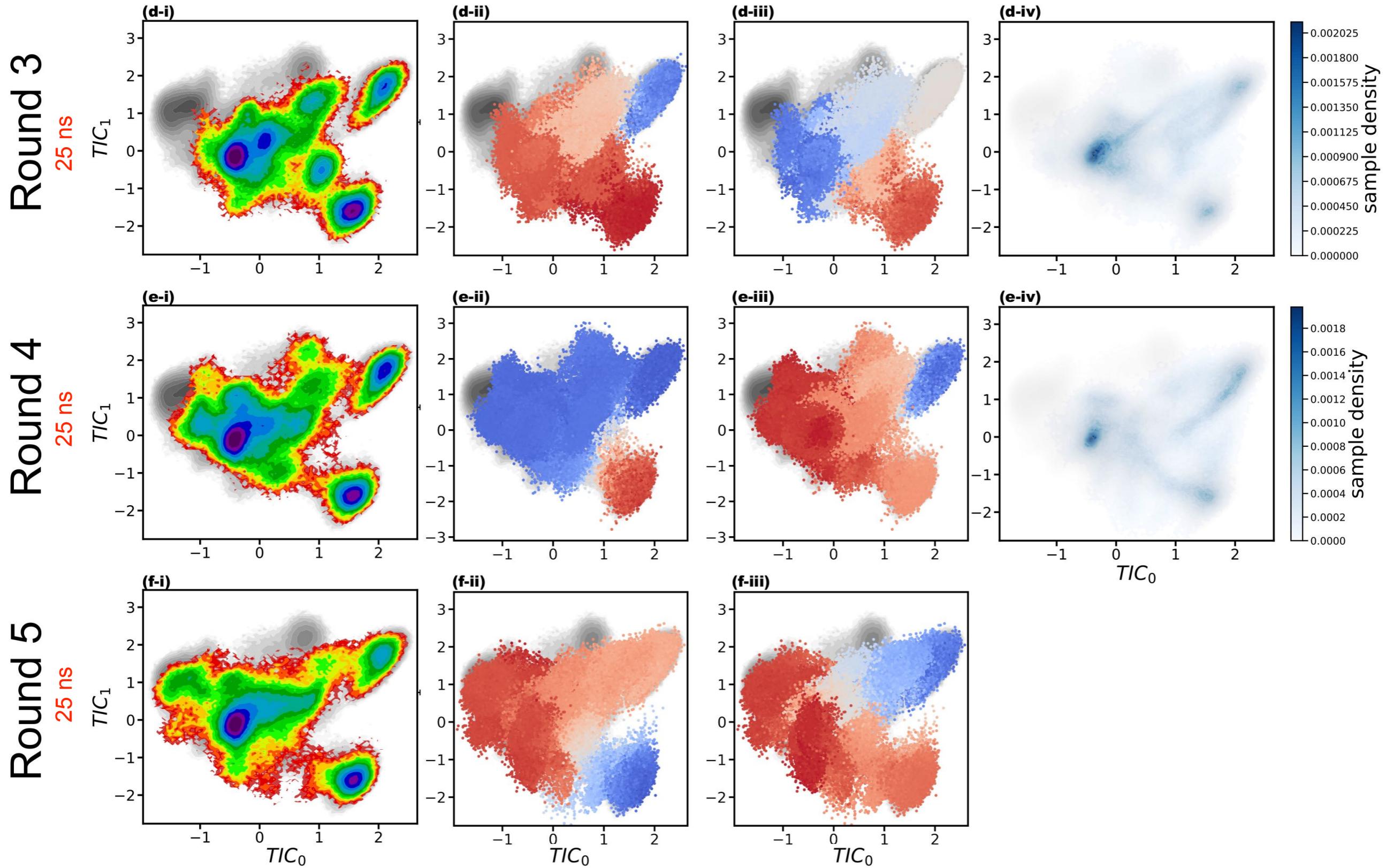
**Trp-Leu-Ala-Leu-Leu
(WLALL)**



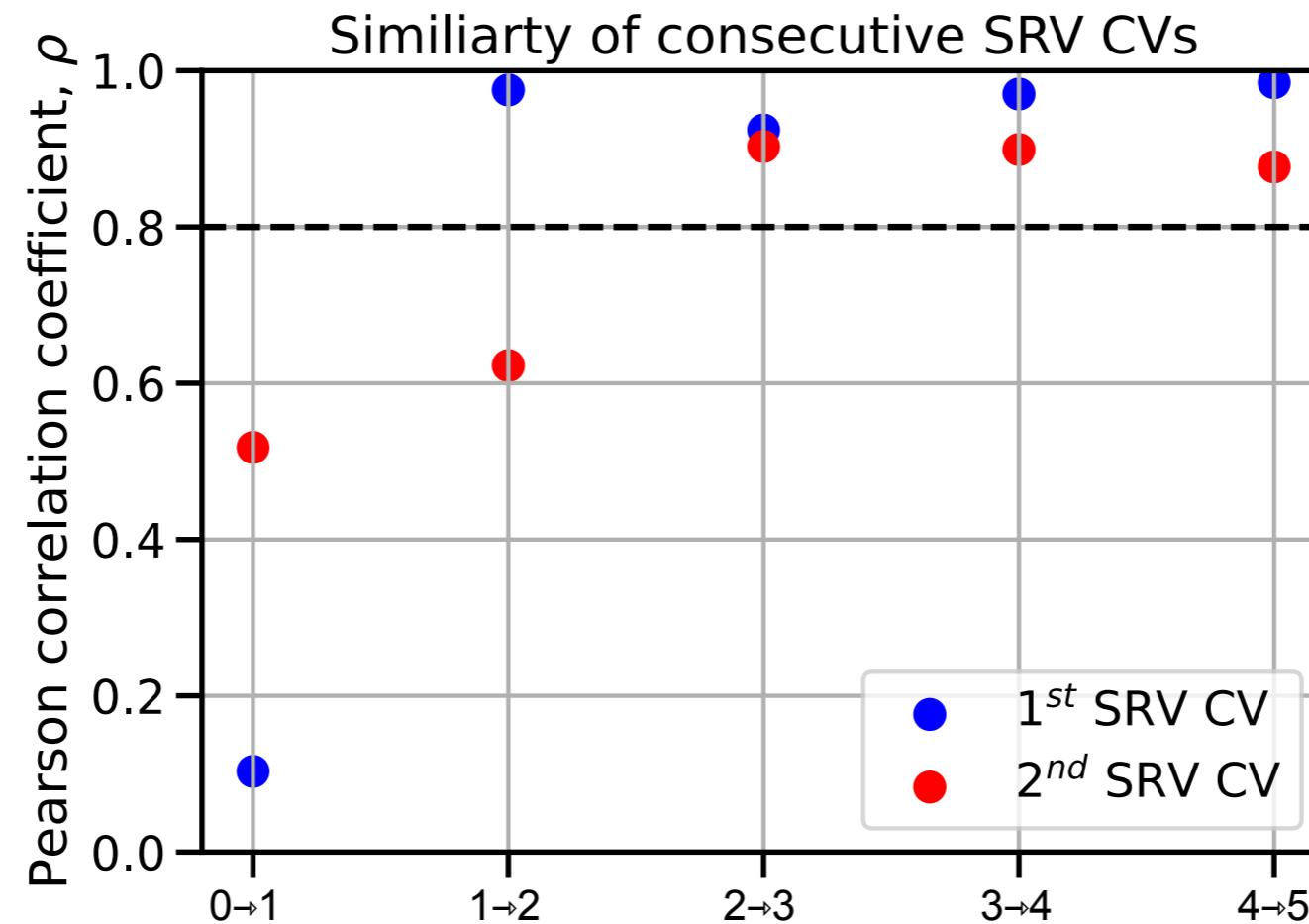
WLALL pentapeptide



WLALL pentapeptide

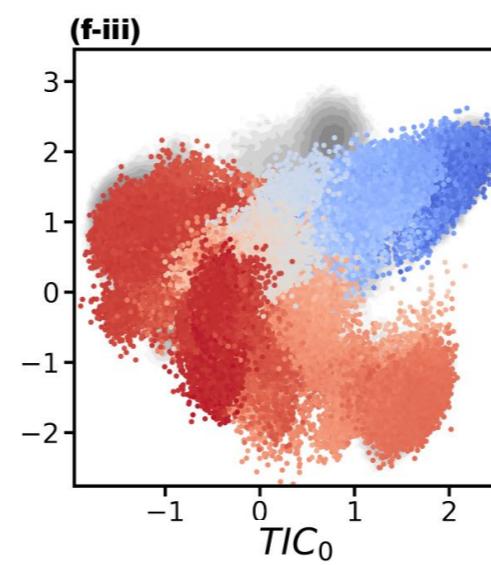
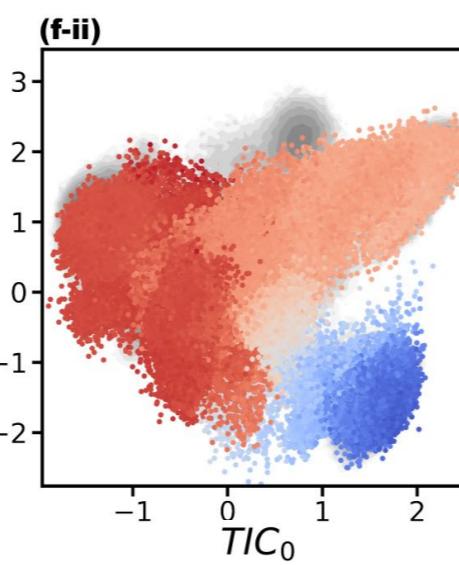
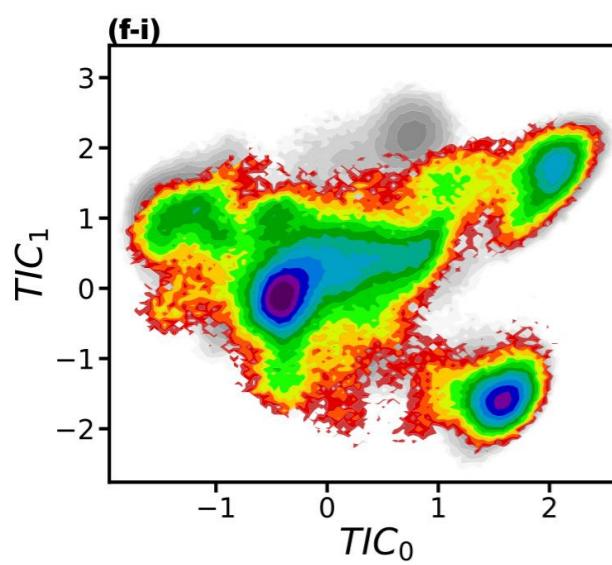


WLALL pentapeptide

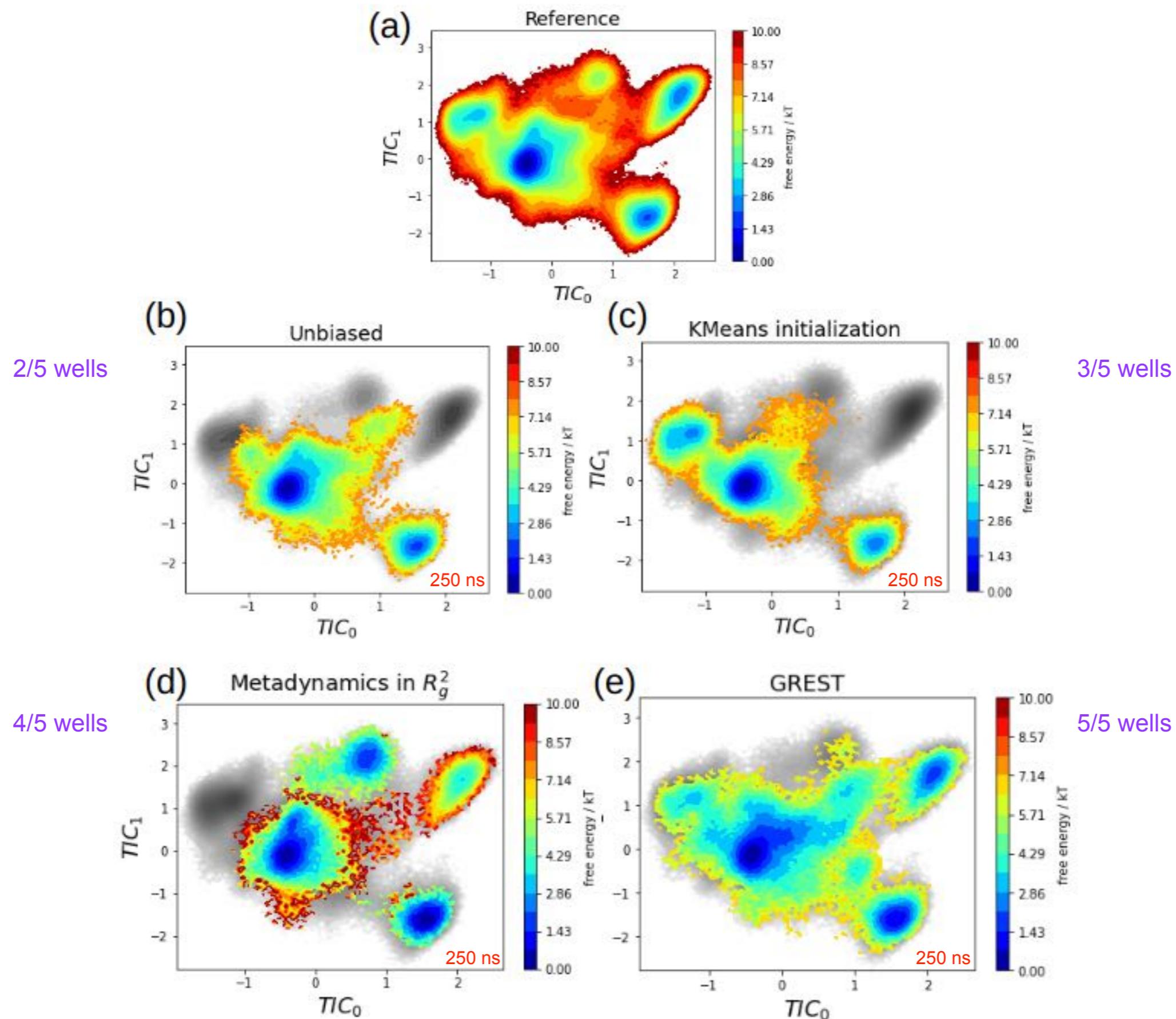


Round 5

25 ns



WLALL pentapeptide



Python package

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Kirill Shmilovich Merge branch 'main' of https://github.com/andrewlferguson/s...	63aeebc 2 weeks ago	82 commits
.github	Adding CI.yaml support for on-demand testing and for all branches	last year
.idea	Updated requirements.txt using PyCharm; removed code coverage ...	last year
devtools	Initial commit after CMS Cookiecutter creation, version 1.6	last year
docs	Adding pages to docs/api.rst	last year
example_notebooks/langevin_four...	Updating typo in comment block to snrv fit call in example notebooks	last year
snrv	Merge branch 'main' of https://github.com/andrewlferguson/snrv	2 weeks ago
.codecov.yml	Initial commit after CMS Cookiecutter creation, version 1.6	last year
.gitignore	Updated .gitignore to ignore emacs temporary files	last year
.lgtm.yml	Initial commit after CMS Cookiecutter creation, version 1.6	last year
CODE_OF_CONDUCT.md	Initial commit after CMS Cookiecutter creation, version 1.6	last year
LICENSE	Initial commit after CMS Cookiecutter creation, version 1.6	last year
MANIFEST.in	Initial commit after CMS Cookiecutter creation, version 1.6	last year
README.md	update README	last month
readthedocs.yml	Initial commit after CMS Cookiecutter creation, version 1.6	last year
requirements.txt	f	9 months ago
requirements.yml	add matplotlib dependency	9 months ago

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Releases v0.1.0 Latest on Jan 19, 2022

Packages No packages published Publish your first package

Contributors andrewlferguson Andrew L Ferguson KirillShmilovich Kirill Shmilovich mrjoness Mike Jones

Conclusions & Outlook

Girsanov reweighting enables rigorous iterative learning of and enhanced sampling in **slow molecular CVs**

Slow CVs reveal **important dynamical motions** and **optimal for enhanced sampling & long-time kinetic models**

Incorporation of **solvent coordinates** respecting permutational invariance

Application to **larger protein systems**

$$W(\omega_{t \rightarrow t+\tau}) = g(x_t) \times M(\omega_{t \rightarrow t+\tau})$$

$$\tilde{\lambda}_k = \frac{\langle u_k | \mathcal{T}(\tau) \circ u_k \rangle_\pi}{\langle u_k | u_k \rangle_\pi}$$

