

FEAT: Free energy Estimators with Adaptive Transport

José Miguel Hernández-Lobato
Department of Engineering
University of Cambridge

<http://jmhl.org>, jmh233@cam.ac.uk

Collaborators



Jiajun He



Yuanqi Du



Francisco
Vargas



Yuanqing
Wang



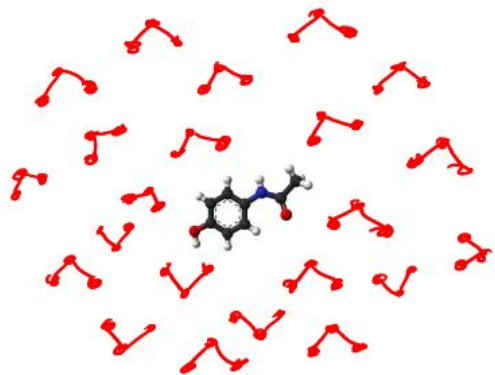
Carla P.
Gomes



Eric
Vanden-Eijnden

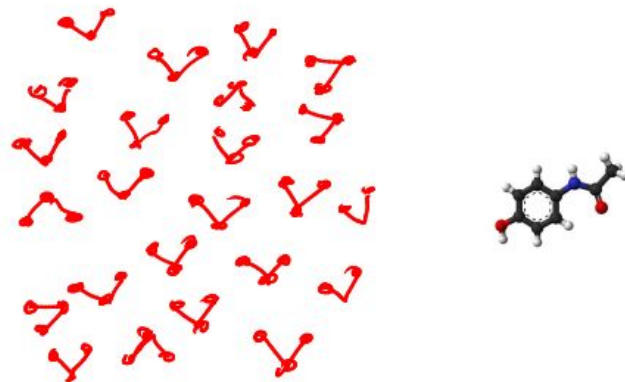
What state of a molecular system is more likely?

State A



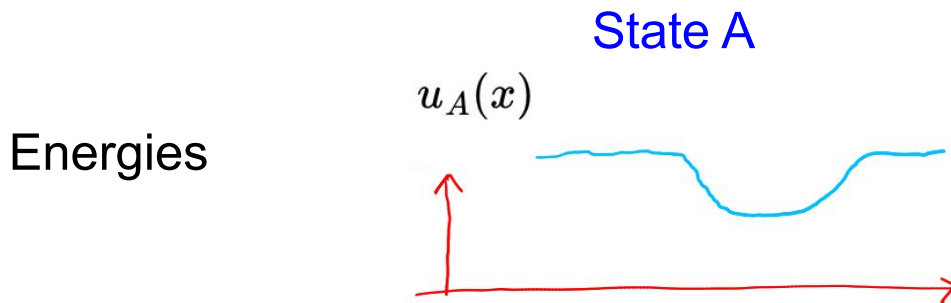
Drug-like molecule surrounded
by water molecules

State B

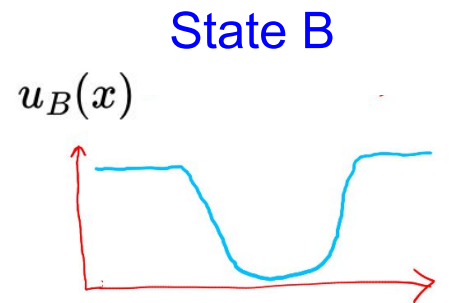


Drug-like molecule separated
from water molecules

From energies to probabilities

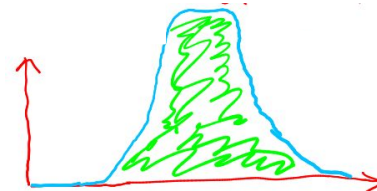
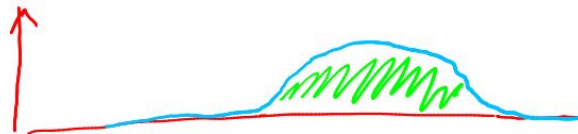


$$p_A(x) = \exp\{-u_A(x) - f_A\}$$



$$p_B(x) = \exp\{-u_B(x) - f_B\}$$

Probabilities



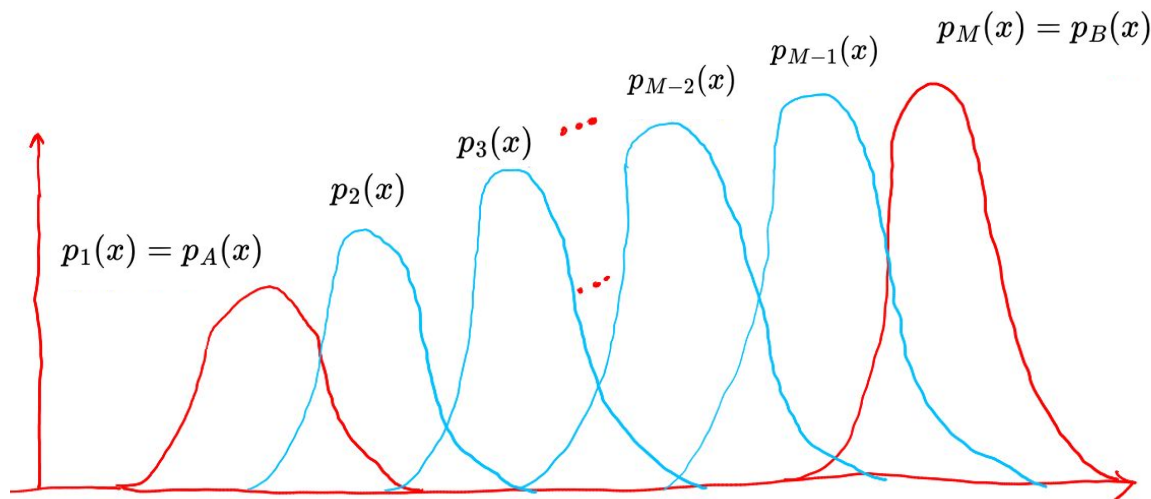
What is the relative probability mass associated with each state?

What is $\Delta f = f_A - f_B$? Δf is usually called the **free energy difference**.

Solution with MBAR

$$p(i_{n,m} = k | x_{n,m}) = \frac{\exp[h - u_k(x_{n,m}) - F_k]}{\sum_j \exp[h - u_j(x_{n,m}) - F_j]}$$

Use **interpolating distributions** with high overlap:

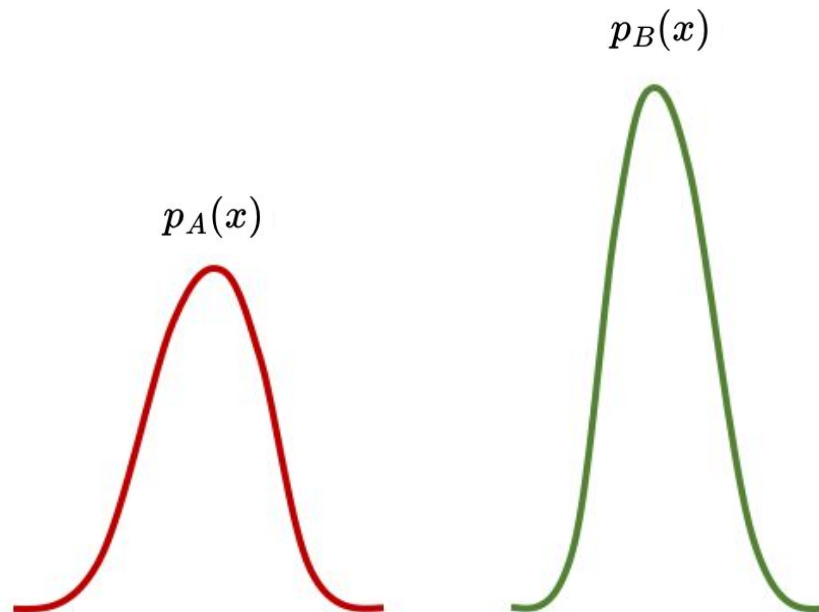


We assume we have N samples from each distribution:

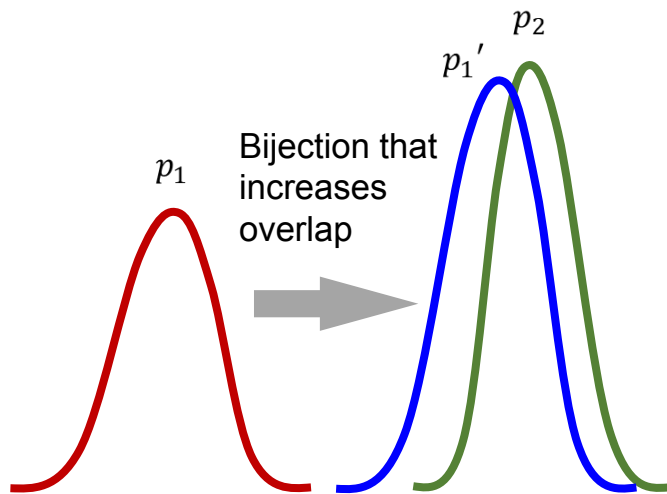
Data = $\{x_{n,m}\}$ where $x_{n,m} \sim p_m(x)$ $n = 1, \dots, N$ $m = 1, \dots, M$

MBAR is great but what if

- No samples from intermediate distributions $p_2(x), \dots, p_{M-1}(x)$ are available.
- We only have samples from $p_A(x)$ and $p_B(x)$.



Target FEP (use mapping to increase overlap)



$$x_0 \sim p_0(x) \quad \frac{dx_t}{dt} = u_t(x_t)$$

$$x_t = x_0 + \int_{s=0}^{s=t} u_s(x_s) ds$$

$$\log p_t(x_t) = \log p_0(x_0) - \int_{s=0}^{s=t} \nabla \cdot u_s(x_s) ds$$

Limitations:

- Need to learn **flexible and accurate bijection**, e.g. using **flow matching** [3].
- But then **difficult** to calculate **changes in density due to transformation**.

[1] C. Jarzynski. Targeted free energy perturbation. *Physical Review E*, 65(4):046122, 2002.

[2] P. Wirnsberger, et al. Targeted free energy estimation via learned mappings. *The Journal of Chemical Physics*, 153(14), 2020.

[3] L. Zhao and L. Wang. Bounding free energy difference with flow matching. *Chinese Physics Letters*, 40(12):120201, 2023.

Proposed approach

Learn forward and backward SDEs running **from $t = 0$ to $t = 1$** and **from $t = 1$ to $t = 0$** that interpolate between distributions p_A and p_B :

$$dX_t = -\sigma^2 \nabla U_t(X_t) dt + u_t dt + \sigma \sqrt{2} \overrightarrow{dB}_t, X_0 \sim p_A$$

$$dX_t = \sigma^2 \nabla U_t(X_t) dt + u_t dt + \sigma \sqrt{2} \overleftarrow{dB}_t, X_1 \sim p_B$$

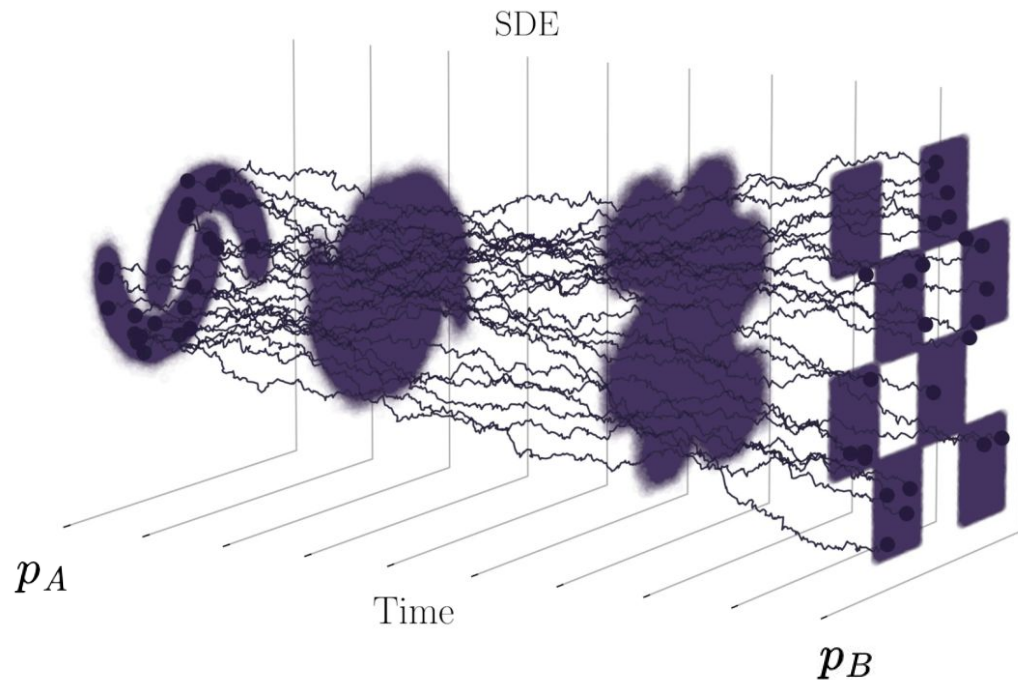
where

- $U_t(X_t)$ is the energy of the marginal distribution for X_t , ($U_0 = U_A$, $U_1 = U_B$).
- $u_t(X_t)$ is a control term ensuring the marginal of X_t has energy $U_t(X_t)$.

These SDEs are a **flexible mapping** between p_A and p_B .

Allow estimation of Δf without calculating expensive changes in density!

Example



Calculating Δf with forward and backward SDEs

Let $t_1 = 0, \dots, t_M = 1$ be times at which we discretize the SDEs trajectories:

$$\begin{aligned}\vec{X}_{t_{i+1}} &= \vec{X}_{t_i} - \sigma^2 \nabla U_{t_i}(\vec{X}_{t_i}) \Delta t_i + u_{t_i}(\vec{X}_{t_i}) \Delta t_i + \sqrt{2\Delta t_i} \sigma \eta_i, & \vec{X}_0 &\sim p_A \\ \overleftarrow{X}_{t_{i-1}} &= \overleftarrow{X}_{t_i} - \sigma^2 \nabla U_{t_i}(\overleftarrow{X}_{t_i}) \Delta t_{i-1} - u_{t_i}(\overleftarrow{X}_{t_i}) \Delta t_{i-1} + \sqrt{2\Delta t_{i-1}} \sigma \eta_i, & \overleftarrow{X}_1 &\sim p_B\end{aligned}$$

where $\Delta t_i = t_{i+1} - t_i$ and $\eta_i \sim \mathcal{N}(0, \mathbf{I})$. Then Δf can be obtained using **importance sampling**:

$$\Delta f = -\log \mathbb{E} \left[\underbrace{\frac{\exp(-U_b(\vec{X}_1)) \prod_{i=1}^M \mathcal{N}^-(\vec{X}_{t_{i-1}} | \vec{X}_{t_i})}{\exp(-U_a(\vec{X}_0)) \prod_{i=1}^M \mathcal{N}^+(\vec{X}_{t_i} | \vec{X}_{t_{i-1}})}}_{\exp -\widetilde{W}(\vec{X})} \right] = \log \mathbb{E} \left[\underbrace{\frac{\exp(-U_a(\overleftarrow{X}_0)) \prod_{i=1}^M \mathcal{N}^+(\overleftarrow{X}_{t_i} | \overleftarrow{X}_{t_{i-1}})}{\exp(-U_b(\overleftarrow{X}_1)) \prod_{i=1}^M \mathcal{N}^-(\overleftarrow{X}_{t_{i-1}} | \overleftarrow{X}_{t_i})}}_{\exp \widetilde{W}(\overleftarrow{X})} \right]$$

\widetilde{W} is **forward/backward log density ratio**. As $M \rightarrow \infty$, \widetilde{W} converges to the **generalized work**.

Improvement with BAR-style approach

Recall the importance sampling estimator:

$$\Delta f = -\log \mathbb{E} \left[\underbrace{\frac{\exp(-U_b(\vec{X}_1)) \prod_{i=1}^M \mathcal{N}^-(\vec{X}_{t_{i-1}} | \vec{X}_{t_i})}{\exp(-U_a(\vec{X}_0)) \prod_{i=1}^M \mathcal{N}^+(\vec{X}_{t_i} | \vec{X}_{t_{i-1}})}}_{\exp - \widetilde{W}(\vec{X})} \right] = \log \mathbb{E} \left[\underbrace{\frac{\exp(-U_a(\vec{X}_0)) \prod_{i=1}^M \mathcal{N}^+(\vec{X}_{t_i} | \vec{X}_{t_{i-1}})}{\exp(-U_b(\vec{X}_1)) \prod_{i=1}^M \mathcal{N}^-(\vec{X}_{t_{i-1}} | \vec{X}_{t_i})}}_{\exp \widetilde{W}(\vec{X})} \right]$$

Use instead a BAR-style **minimum variance method**:

$$\Delta f = \log \frac{\mathbf{E}[g(-\widetilde{W}(\vec{X}) + C)]}{\mathbf{E}[g(\widetilde{W}(\vec{X}) - C)]} \quad g(x) = \frac{1}{1 + \exp(x)}, C = \Delta f$$

1. Initialize C ;
2. Calculate Δf ; Set $C \leftarrow \Delta f$;
3. Repeat (2) until converge.

How to learn the SDEs?

We use the framework of **Stochastic Interpolants (SI)**, defining the **interpolant**

$$X_t = \alpha_t x_A + \beta_t x_B + \gamma_t \epsilon, \quad x_A \sim p_A, x_B \sim p_B, \epsilon \sim \mathcal{N}(0, \mathbf{I})$$

where $\alpha_0 = 1, \alpha_1 = 0$; $\beta_0 = 0, \beta_1 = 1$; and $\gamma_0 = \gamma_1 = 0$.

The distribution of the interpolant is the same as the one generated by the SDEs if

$$u_t(x) = \mathbb{E}[\partial_t X_t \mid X_t = x], \quad \nabla U_t(x) = \gamma_t^{-1} \mathbb{E}[\epsilon \mid X_t = x]$$

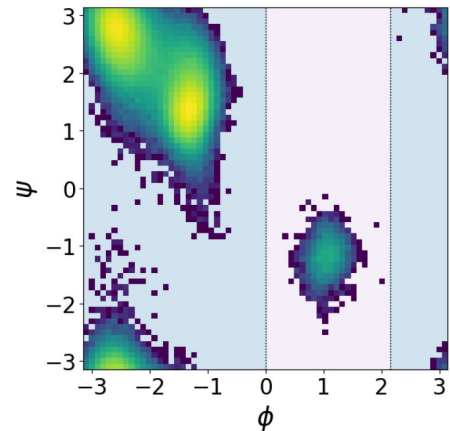
If we parametrize these functions as **neural networks** we can learn them by minimizing

$$\mathcal{L}_u(\psi) = \mathbb{E}_{t \sim \mathcal{U}(0,1)} \mathbb{E}_{x_A, x_B, \epsilon} \left[\lambda_t \left\| u_t^\psi(X_t) - \partial_t X_t \right\|^2 \right]$$

$$\mathcal{L}_U^{\text{DSM}}(\theta) = \mathbb{E}_{t \sim \mathcal{U}(0,1)} \mathbb{E}_{x_A, x_B, \epsilon} \left[\eta_t \left\| \nabla U_t^\theta(X_t) - \gamma_t^{-1} \epsilon \right\|^2 \right]$$

Experimental validation

- **GMM:**
16-mode **GMM** and a **40-mode GMM**.
- **LJ system:**
without **LJ potential** and system **with LJ potential**.
- **Alanine dipeptide – Solvation (ALDP-S):**
ALDP in **vacuum** and ALDP in **implicit solvent**.
- **Alanine dipeptide – Transition (ALDP-T):**
ALDP in **two meta-stable states**.



Results

Method	GMM		LJ		
	$d = 40$	$d = 100$	$d = 55 \times 3$	$d = 79 \times 3$	$d = 128 \times 3$
Reference	0	0	234.77 \pm 0.09	357.43 \pm 3.43	595.98 \pm 0.58
Target FEP w. FM	0.09 \pm 0.26	-17.96 \pm 1.49	232.06 \pm 0.03	*	*
Neural TI	-181.63 \pm 6.65	-402.93 \pm 283.75	328.55 \pm 336.39	468.76 \pm 391.16	N/A
Ours	0.04 \pm 0.04	-5.34 \pm 1.52	232.47 \pm 0.15	356.74 \pm 0.79	595.04 \pm 6.52

Method	ALDP-S	ALDP-T
	$d = 22 \times 3$	$d = 22 \times 3$
Reference	29.43 \pm 0.01	-4.25 \pm 0.05
Target FEP w. FM	29.47 \pm 0.22	-4.78 \pm 0.32
Neural TI	24.93 \pm 3.13	-4.11 \pm 2.56
Ours	29.38 \pm 0.04	-4.56 \pm 0.08

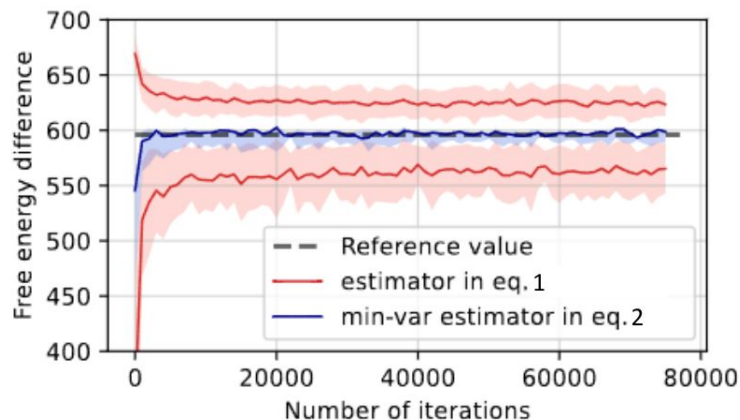
Results

$$\text{Eq 1: } \Delta f = -\log \mathbf{E}_{\tilde{\mathbf{p}}}[\exp(-W)] = \log \mathbf{E}_{\tilde{\mathbf{p}}}[\exp(W)]$$

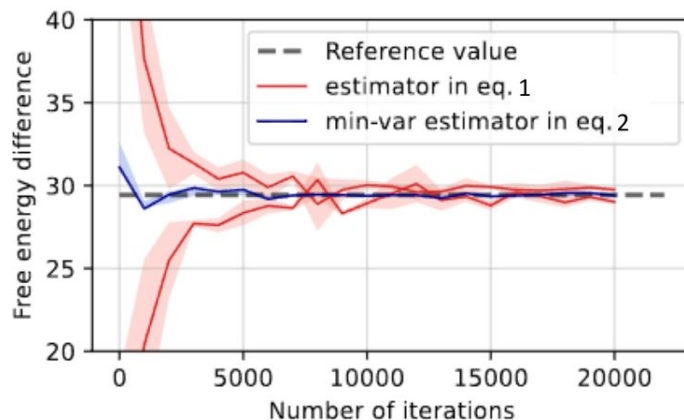
Importance sampling estimator

$$\text{Eq 2: } \Delta f = -\log \frac{\mathbf{E}_{\tilde{\mathbf{p}}}[g(W-C)]}{\mathbf{E}_{\tilde{\mathbf{p}}}[g(-W+C)]} + C$$

Minimum variance estimator



(b) LJ-128.



(c) ALDP-S.

Take home messages

FEAT: Free energy Estimators with Adaptive Transport is a method that

- calculates free energy differences using only samples from states A and B,
- does not need samples from intermediate distributions,
- uses SI framework to learn SDEs that flexibly map between distributions,
- avoids calculating expensive changes in density due to sample transport,
- uses a BAR-like minimum variance estimator,
- outperforms baselines like Neural TI and Neural target FEP,

Limitations: requires extra cost for training neural networks which may not be accurate enough in large systems.

Thanks!