

Using stochastic approximation and self-adjusted mixture sampling for molecular design: Some collected notes

Patrick B. Grinaway,^{1,*} Julie M. Behr,^{1,†} Zhiqiang Tan,^{2,‡} and John D. Chodera^{1,§}

¹Computational and Systems Biology Program, Sloan Kettering Institute,
Memorial Sloan Kettering Cancer Center, New York, NY 10065

²Department of Statistics, Rutgers University, Piscataway, NJ 08854

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We collect some notes on using stochastic approximation with multiple self-adjusted mixture sampling simulations for molecular design problems.

Keywords: stochastic approximation; molecular simulation; computer aided ligand design; expanded ensemble; self-adjusted mixture sampling

MAXIMIZING LIGAND BINDING AFFINITY FOR A PROTEIN

THE ALGORITHM

Suppose we are trying to design a ligand that maximizes the binding affinity to a target protein. Let the index $j \in \{1, \dots, m\}$ denote the ligand identity.

We define two systems with associated probability densities:

Suppose we have a protein:ligand:solvent system with probability density

$$p_{1j}(x) = e^{\zeta_{1j}^*} q_{1j}(x), \quad j = 1, \dots, m \quad (1)$$

where we note the dimensionality of x may depend on the index j .

We also have a ligand:solvent system with probability density

$$p_{2j}(x) = e^{\zeta_{2j}^*} q_{2j}(x) \quad j = 1, \dots, m \quad (2)$$

Consider the expanded ensembles

$$(j, x)_1 \sim Q_1(\zeta_1, \zeta_2) \equiv \pi_{1j}(\zeta_1, \zeta_2) e^{-\zeta_{1j}} q_{1j}(x) \quad (3)$$

$$(j, x)_2 \sim Q_2(\zeta_1, \zeta_2) \equiv \pi_{2j}(\zeta_1, \zeta_2) e^{-\zeta_{2j}} q_{2j}(x) \quad (4)$$

To identify ligands with high binding affinity for a protein, we propose to design a chain with the limiting marginal distribution proportional to the binding affinity

$$p_{1j}, p_{2j} \propto K_j \equiv e^{\zeta_{2j}^* - \zeta_{1j}^*} \quad (5)$$

To do this, we allow the π to depend on $\zeta_1 \equiv \{\zeta_{11}, \dots, \zeta_{1m}\}$ and $\zeta_2 \equiv \{\zeta_{21}, \dots, \zeta_{2m}\}$ such that

$$\pi_{1j} = \pi_{2j} \equiv \pi_j(\zeta_1, \zeta_2) \quad (6)$$

Specifically, we propose

$$\pi_{1j} \pi_{2j} \equiv e^{\zeta_{2j} - \zeta_{1j}} = \frac{e^{\zeta_{2j} - \zeta_{1j}}}{\sum_{k=1}^m e^{\zeta_{2k} - \zeta_{1k}}} \quad (7)$$

The algorithm is as follows: At iteration n ,

- Sample from the expanded ensemble defined by Q_1 and Q_2 for the current $(\zeta_1^{(n-1)}, \zeta_2^{(n-1)})$.

$$(l, x)_1 \sim Q_1(\zeta_1^{(n-1)}, \zeta_2^{(n-1)}) \quad (8)$$

$$(l, x)_2 \sim Q_2(\zeta_1^{(n-1)}, \zeta_2^{(n-1)}) \quad (9)$$

- Update estimates of $(\zeta_1^{(n-1)}, \zeta_2^{(n-1)})$.

$$\zeta_{1j}^{(n)} = \zeta_{1j}^{(n-1)} + n^{-1} \frac{\delta_j(l_1)}{\pi_j(\zeta_1^{(n-1)}, \zeta_2^{(n-1)})} \quad (10)$$

$$\zeta_{2j}^{(n)} = \zeta_{2j}^{(n-1)} + n^{-1} \frac{\delta_j(l_2)}{\pi_j(\zeta_1^{(n-1)}, \zeta_2^{(n-1)})} \quad (11)$$

Set $\zeta_{11}^{(n)} = \zeta_{21}^{(n)} = 0$.

NOTE: It may simplify things to break out the update of $\pi_j^{(n)}$ into an explicit recursion step instead of explicitly writing $\pi_j(\zeta_1, \zeta_2)$.

NOTE: It would be best if we can abandon using π_j and instead work in log space as $g_j \equiv -\log \pi_j$ instead, since any implementation involving π_j directly will run into numerical underflow/overflow issues otherwise.

See a simple example of this algorithm in action for a set of Gaussian distributions [at this link](#).

ANOTHER POSSIBILITY

Set $\pi_{1j}(\zeta_1, \zeta_2) \propto e^{\zeta_{2j} - \zeta_{1j}}$ but $\pi_{2j}(\zeta_1, \zeta_2) \propto 1$.
With this choice, we have

$$(j, x)_1 \sim Q_1(\zeta_1, \zeta_2) \equiv e^{-\zeta_{2j}} q_{1j}(x) \quad (12)$$

$$(j, x)_2 \sim Q_2(\zeta_1, \zeta_2) \equiv e^{-\zeta_{2j}} q_{2j}(x) \quad (13)$$

The corresponding algorithm is as follows: At iteration n ,

* patrick.grinaway@choderalab.org

† julie.behr@choderalab.org

‡ ztan@stat.rutgers.edu

§ Corresponding author; john.chodera@choderalab.org

- 51 • Sample $(l, x)_2$ from the expanded ensemble defined by
52 Q_2 for the current $\zeta_2^{(n-1)}$.

$$(l, x)_2 \sim Q_2(\zeta_1, \zeta_2) \equiv e^{-\zeta_{2j}} q_{2j}(x) \quad (14)$$

- 53 • Update estimate of $\zeta_2^{(n)}$.

$$\zeta_{2j}^{(n)} = \zeta_{2j}^{(n-1)} + n^{-1} \frac{\delta_j(l_2)}{m-1} \quad (15)$$

54 Set $\zeta_{21}^{(n)} = 0$.

- 55 • Sample $(l, x)_1$ from the expanded ensemble defined by
56 Q_1 for the current $\zeta_2^{(n)}$.

$$(l, x)_1 \sim Q_1(\zeta_1, \zeta_2) \equiv e^{-\zeta_{1j}} q_{1j}(x) \quad (16)$$

GENERALIZATION OF THE DESIGN PROBLEM

58 More generally, consider we have s different probability
59 densities

$$p_{ij}(x) = e^{\zeta_{ij}^*} q_{ij}(x), \quad i = 1, \dots, s, \quad j = 1, \dots, m \quad (17)$$

60 and we desire to design a chain where the marginal distribu-
61 tions of all s chains are

$$p_{ij} \propto \prod_{i'=1}^s e^{-\theta_{s'} \zeta_{i'j}^*} = \exp \left[- \sum_{i'=1}^s \theta_{s'} \zeta_{i'j}^* \right] \quad \forall i = 1, \dots, s \quad (18)$$

62 where the *design vector* $\Theta \equiv \{\theta_1, \dots, \theta_s\}$ specifies how dif-
63 ferent targets and antitargets are used in weighting the design
64 constraints.

81 We impose super-detailed balance:

$$\pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) \mathcal{P}[\mathcal{T}] \mathcal{A}[\mathcal{T}] = \pi(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) \mathcal{P}[\tilde{\mathcal{T}}] \mathcal{A}[\tilde{\mathcal{T}}] \quad (21)$$

82

$$\begin{aligned} & \pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}}) \phi(x_{\text{new}} | x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \Phi(x \rightarrow x' | \mathcal{M}_{\text{old}} \rightarrow \mathcal{M}_{\text{new}}) \mathcal{A}[\mathcal{T}] \\ & = \pi(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) P(\mathcal{M}_{\text{old}} | \mathcal{M}_{\text{new}}) \phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \Phi(x' \rightarrow x | \mathcal{M}_{\text{new}} \rightarrow \mathcal{M}_{\text{old}}) \mathcal{A}[\tilde{\mathcal{T}}] \end{aligned} \quad (22)$$

83 Collecting terms to compute the acceptance criteria:

$$\frac{\mathcal{A}[\mathcal{T}]}{\mathcal{A}[\tilde{\mathcal{T}}]} = \frac{\pi(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) P(\mathcal{M}_{\text{old}} | \mathcal{M}_{\text{new}}) \phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \Phi(x' \rightarrow x | \mathcal{M}_{\text{new}} \rightarrow \mathcal{M}_{\text{old}})}{\pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}}) \phi(x_{\text{new}} | x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \Phi(x \rightarrow x' | \mathcal{M}_{\text{old}} \rightarrow \mathcal{M}_{\text{new}})} \quad (23)$$

$$= \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})} P(\mathcal{M}_{\text{old}} | \mathcal{M}_{\text{new}}) \phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})} P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}}) \phi(x_{\text{new}} | x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} e^{-\Delta S[x \rightarrow x' | \mathcal{M}_{\text{old}} \rightarrow \mathcal{M}_{\text{new}}]} \quad (24)$$

$$= \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})} P(\mathcal{M}_{\text{old}} | \mathcal{M}_{\text{new}}) \phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})} P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}}) \phi(x_{\text{new}} | x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} e^{-w[x \rightarrow x' | \lambda=0 \rightarrow 1]} \frac{e^{-u(x, \lambda=0)}}{e^{-u(x', \lambda=1)}} \quad (25)$$

84 **Two-stage scheme**

85 Starting from $(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}})$, the proposal scheme is:

65 We postulate that we can do this by defining $\pi_i(Z)$ for $Z \equiv$
66 $\{\zeta_1, \dots, \zeta_s\}$ as

$$p_{ij}(Z | \Theta) \propto \exp \left[- \sum_{i'=1}^s \theta_{s'} \zeta_{i'j} \right] \quad (19)$$

THE SAMPLING SCHEME

68 Suppose we are proposing a transition from a molecule
69 \mathcal{M}_{old} to \mathcal{M}_{new} , where the initial molecule has configuration
70 $x \equiv (x_{\text{core}}, x_{\text{old}})$ and the new molecule has configuration
71 $x' \equiv (x'_{\text{core}}, x'_{\text{new}})$:

$$\mathcal{T} : (x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) \rightarrow (x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) \quad (20)$$

72 [JDC: We still need to incorporate the stochastic nature of
73 the order of atom and torsion proposals into the ϕ terms.]

74

Hybrid scheme

75 Starting from $(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}})$, the proposal scheme is:

- 76 1. $\mathcal{M}_{\text{new}} \sim P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}})$
- 77 2. $x_{\text{new}} \sim \phi(x_{\text{new}} | x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})$
- 78 3. $(x'_{\text{new}}, x'_{\text{old}}, x'_{\text{core}}) \sim \Phi(x \rightarrow x' | \mathcal{M}_{\text{old}} \rightarrow \mathcal{M}_{\text{new}})$
- 79 4. Accept $(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}})$ with probability
80 $\min\{1, \mathcal{A}[\mathcal{T}]\}$.

- 86 1. $\mathcal{M}_{\text{new}} \sim P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}})$

- 87 2. $(x'_{\text{old}}, x'_{\text{core}}) \sim \Phi_{\text{delete}}(x \rightarrow x' | \mathcal{M}_{\text{old}})$

- 88 3. $x'_{\text{new}} \sim \phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})$ 90 5. Accept $(x''_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}})$ with probability
 89 4. $(x''_{\text{new}}, x''_{\text{core}}) \sim \Phi_{\text{insert}}(x' \rightarrow x''|\mathcal{M}_{\text{new}})$ 91 $\min\{1, \mathcal{A}[\mathcal{T}]\}.$

92 We impose super-detailed balance:

$$\pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) \mathcal{P}[\mathcal{T}] \mathcal{A}[\mathcal{T}] = \pi(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) \mathcal{P}[\tilde{\mathcal{T}}] \mathcal{A}[\tilde{\mathcal{T}}] \quad (26)$$

93

$$\begin{aligned} & \pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}}) \Phi_{\text{delete}}(x \rightarrow x'|\mathcal{M}_{\text{old}}) \phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \Phi_{\text{insert}}(x' \rightarrow x''|\mathcal{M}_{\text{new}}) \mathcal{A}[\mathcal{T}] \\ &= \pi(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}}) \Phi_{\text{delete}}(x'' \rightarrow x'|\mathcal{M}_{\text{new}}) \phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \Phi_{\text{insert}}(x' \rightarrow x|\mathcal{M}_{\text{old}}) \mathcal{A}[\tilde{\mathcal{T}}] \end{aligned} \quad (27)$$

94 Collecting terms to compute the acceptance criteria:

$$\begin{aligned} & \frac{\mathcal{A}[\mathcal{T}]}{\mathcal{A}[\tilde{\mathcal{T}}]} \\ &= \frac{\pi(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}}) \Phi_{\text{delete}}(x'' \rightarrow x'|\mathcal{M}_{\text{new}}) \phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \Phi_{\text{insert}}(x' \rightarrow x|\mathcal{M}_{\text{old}})}{\pi(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}}) \Phi_{\text{insert}}(x' \rightarrow x''|\mathcal{M}_{\text{new}}) \phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \Phi_{\text{delete}}(x \rightarrow x'|\mathcal{M}_{\text{old}})} \end{aligned} \quad (28)$$

$$= \frac{e^{-u(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})} P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}})}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})} P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}})} e^{-\Delta S_{\text{insert}}[x' \rightarrow x''|\mathcal{M}_{\text{new}}]} \frac{\phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{\phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} e^{-\Delta S_{\text{delete}}[x \rightarrow x'|\mathcal{M}_{\text{old}}]} \quad (29)$$

$$\begin{aligned} &= \frac{e^{-u(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})} P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}}) \phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})} P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}}) \phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} \\ &\times e^{-w_{\text{insert}}[x' \rightarrow x''|\mathcal{M}_{\text{new}}]} \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}; \mathcal{M}_{\text{new}}, \lambda=0)}}{e^{-u(x'_{\text{core}}, x'_{\text{new}}; \mathcal{M}_{\text{new}}, \lambda=1)}} e^{-w_{\text{delete}}[x \rightarrow x'|\mathcal{M}_{\text{old}}]} \frac{e^{-u(x_{\text{core}}, x_{\text{old}}; \mathcal{M}_{\text{old}}, \lambda=1)}}{e^{-u(x'_{\text{core}}, x'_{\text{old}}; \mathcal{M}_{\text{old}}, \lambda=0)}} \end{aligned} \quad (30)$$

$$\begin{aligned} &= \frac{e^{g(\mathcal{M}_{\text{new}})} P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}}) \phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{e^{g(\mathcal{M}_{\text{old}})} P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}}) \phi(x'_{\text{new}}|x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} e^{-w_{\text{insert}}[x' \rightarrow x''|\mathcal{M}_{\text{new}}]} e^{-w_{\text{delete}}[x \rightarrow x'|\mathcal{M}_{\text{old}}]} \\ &\times \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}; \mathcal{M}_{\text{new}}, \lambda=0)}}{e^{-u(x'_{\text{core}}, x'_{\text{new}}; \mathcal{M}_{\text{new}}, \lambda=1)}} \cdot \frac{e^{-u(x_{\text{core}}, x_{\text{old}}; \mathcal{M}_{\text{old}}, \lambda=1)}}{e^{-u(x_{\text{core}}, x_{\text{old}}; \mathcal{M}_{\text{old}}, \lambda=0)}} \cdot \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}; \mathcal{M}_{\text{new}}, \lambda=0)}}{e^{-u(x'_{\text{core}}, x'_{\text{old}}; \mathcal{M}_{\text{old}}, \lambda=0)}} \end{aligned} \quad (31)$$

95 SAMPLING SCHEME IMPLEMENTATION

96 Hybrid scheme

97 Implementation

98 We break the **hybrid** acceptance criteria (Eq. 25) into the
 99 following components:

- 100 • The **stationary probability** from the initial and
 101 final chemical states, which is computed in the
 102 ExpandedEnsembleSampler:

$$\begin{aligned} \log P_{\text{stationary}} &= \log \frac{e^{-u(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})}}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})}} \quad (32) \\ &= \log P_{\text{final}} - \log P_{\text{initial}} \quad (33) \end{aligned}$$

103 which we further decompose into the **initial** and **final**
 104 **log probabilities** of chemical states:

$$\log P_{\text{final}} = -u(x'_{\text{core}}, x'_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}}) \quad (34)$$

$$\log P_{\text{initial}} = -u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}}) \quad (35) \quad 111$$

- 105 • The **chemical proposal probabilities**, which are com-
 106 puted by the ProposalEngine:

$$\log P_{\text{chemical}} = \log \frac{P(\mathcal{M}_{\text{old}}|\mathcal{M}_{\text{new}})}{P(\mathcal{M}_{\text{new}}|\mathcal{M}_{\text{old}})} \quad (36)$$

- 107 • The **geometry proposal probabilities**, computed by the
 108 GeometryEngine:

$$\begin{aligned} \log P_{\text{geometry}} &= \log \frac{\phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{\phi(x_{\text{new}}|x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} \quad (37) \\ &= \log P_{\text{reverse}} - \log P_{\text{forward}} \quad (38) \end{aligned}$$

which we further decompose into **reverse** and **forward**
 geometry proposal probabilities:

$$\log P_{\text{reverse}} = \log \phi(x'_{\text{old}}|x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \quad (39)$$

$$\log P_{\text{forward}} = \log \phi(x_{\text{new}}|x_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \quad (40)$$

and finally the **NCMC** component, computed by the

NCMCEngine:

$$\log P_{\text{NCMC}} = \log e^{-\Delta S[x \rightarrow x' | \mathcal{M}_{\text{old}} \rightarrow \mathcal{M}_{\text{new}}]} \quad (41)$$

$$= \log \left[e^{-w[x \rightarrow x' | \lambda=0 \rightarrow 1]} \frac{e^{-u(x, \lambda=0)}}{e^{-u(x', \lambda=1)}} \right] \quad (42)$$

$$= \log P_{\text{work}} + \log P_{\text{energy}} \quad (43)$$

which we further decompose into **work** and **energy** change contributions

$$\log P_{\text{work}} = -w[x \rightarrow x' | \lambda = 0 \rightarrow 1] \quad (44)$$

$$\log P_{\text{energy}} = u(x', \lambda = 1) - u(x, \lambda = 0) \quad (45)$$

JDC: Check that the energy component here is really the reduced potential for both GHMC and VV, rather than the total reduced energy.

With this definition of terms, the overall acceptance probability is therefore given as

$$\begin{aligned} \log P_{\text{accept}} &= \log P_{\text{final}} - \log P_{\text{initial}} + \log P_{\text{chemical}} \\ &\quad + \log P_{\text{reverse}} - \log P_{\text{forward}} \\ &\quad + \log P_{\text{work}} + \log P_{\text{energy}} \end{aligned} \quad (46)$$

Testing

We use several kinds of tests to ensure that the quantities described above are computed correctly.

check_alchemical_null_elimination: This test ensures that the NCMC work is computed correctly by performing a *null transformation* in which the overall free energy change should be zero. We can show that the expectation of the exponentiated work should be given by the exponentiated free energy difference (due to Jarzynski [1]):

$$\begin{aligned} &E_{0 \rightarrow 1} [e^{-w[x \rightarrow x' | \lambda=0 \rightarrow 1]}] \\ &= \sum_{x_0 \cdots x_N} P_{0 \rightarrow 1}[x \rightarrow x'] e^{-w[x \rightarrow x' | \lambda=0 \rightarrow 1]} \\ &= \sum_{x_0 \cdots x_N} \pi_0(x_0) \left[\prod_{n=1}^N K_n(x_{n-1}, x_n) \right] e^{-\sum_{n=1}^N (u_n(x_n) - u_{n-1}(x_n))} \\ &= \sum_{x_0 \cdots x_N} \pi_0(x_0) \left[\prod_{n=1}^N K_n(x_{n-1}, x_n) \right] \prod_{n=1}^N \frac{q_n(x_n)}{q_{n-1}(x_n)} \\ &= \sum_{x_0 \cdots x_N} \pi_0(x_0) \left[\prod_{n=1}^N K_n(x_{n-1}, x_n) \right] \prod_{n=1}^N \frac{Z_n \pi_n(x_n)}{Z_{n-1} \pi_{n-1}(x_n)} \\ &= \frac{Z_N}{Z_0} \sum_{x_0 \cdots x_N} \pi_0(x_1) K_0(x_0, x_1) \left[\prod_{n=2}^N K_n(x_{n-1}, x_n) \right] \prod_{n=1}^N \frac{\pi_n(x_n)}{\pi_{n-1}(x_n)} \\ &= \frac{Z_N}{Z_0} \sum_{x_1 \cdots x_N} \pi_0(x_1) \frac{\pi_1(x_1)}{\pi_0(x_1)} \left[\prod_{n=2}^N K_n(x_{n-1}, x_n) \right] \prod_{n=1}^N \frac{\pi_n(x_n)}{\pi_{n-1}(x_n)} \\ &= \frac{Z_N}{Z_0} = e^{-(f_N - f_0)} = e^{-\Delta f_{0 \rightarrow 1}} \end{aligned} \quad (47)$$

We can test the $\log P_{\text{work}}$ component using the one-sided EXP estimator

$$\Delta f_{0 \rightarrow 1} = -\log E_{0 \rightarrow 1} [\exp \log P_{\text{work}}] \quad (48)$$

Note that we are only testing the **work** contribution here. The differential path action ($\log \Delta S = \log P_{\text{work}} + \log P_{\text{energy}}$) obeys different statistics. **JDC: Is there a similar test we could apply to $\log \Delta S$?**

The EXP estimator can produce heavily biased estimates, making the uncertainty estimates unreliable [2], so instead use the bidirectional BAR estimator to estimate switches in both directions when possible [3]. This also ensures that the NCMC method obeys the correct symmetry relations when run forward and backward. In particular, the protocol must be *symmetric* unless additional corrections for selecting the same protocol and its time-reverse are included [4].

check_harmonic_oscillator_ncmc: The same principles as above, applied to a harmonic oscillator. This scheme tests only the NCMCIntegrator, rather than the whole NCMCEngine.

Two-stage scheme

Implementation

We break the **two-stage** acceptance criteria (Eq 31) into the following components:

- The **stationary probability** from the initial and final chemical states, which is computed in the ExpandedEnsembleSampler:

$$\begin{aligned} \log P_{\text{stationary}} &= \log \frac{e^{-u(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}})}}{e^{-u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}})}} \\ &= \log P_{\text{final}} - \log P_{\text{initial}} \end{aligned} \quad (49)$$

which we further decompose into the **initial** and **final** log probabilities of chemical states:

$$\log P_{\text{final}} = -u(x''_{\text{core}}, x''_{\text{new}}, \mathcal{M}_{\text{new}}) + g(\mathcal{M}_{\text{new}}) \quad (51)$$

$$\log P_{\text{initial}} = -u(x_{\text{core}}, x_{\text{old}}, \mathcal{M}_{\text{old}}) + g(\mathcal{M}_{\text{old}}) \quad (52)$$

- The **chemical proposal probabilities**, which are computed by the ProposalEngine:

$$\log P_{\text{chemical}} = \log \frac{P(\mathcal{M}_{\text{old}} | \mathcal{M}_{\text{new}})}{P(\mathcal{M}_{\text{new}} | \mathcal{M}_{\text{old}})} \quad (53)$$

- The **geometry proposal probabilities**, computed by the GeometryEngine:

$$\log P_{\text{geometry}} = \log \frac{\phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}})}{\phi(x'_{\text{new}} | x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}})} \quad (54)$$

$$= \log P_{\text{reverse}} - \log P_{\text{forward}} \quad (55)$$

160 which we further decompose into **reverse** and **forward** 167
161 geometry proposal probabilities:

$$\log P_{\text{reverse}} = \log \phi(x'_{\text{old}} | x'_{\text{core}}, \mathcal{M}_{\text{new}}, \mathcal{M}_{\text{old}}) \quad (56)$$

$$\log P_{\text{forward}} = \log \phi(x'_{\text{new}} | x'_{\text{core}}, \mathcal{M}_{\text{old}}, \mathcal{M}_{\text{new}}) \quad (57)$$

162 and finally the **NCMC** components, computed by the 168
163 NCMCEngine, where there are now two stages (**delete** 169
164 and **insert**):

$$\log P_{\text{delete}} = \log e^{-\Delta S_{\text{delete}}[x \rightarrow x' | \mathcal{M}_{\text{old}}]} \quad (58)$$

$$= \log \left[e^{-w_{\text{delete}}[x \rightarrow x' | \mathcal{M}_{\text{old}}]} \frac{e^{-u(x, \mathcal{M}_{\text{old}}, \lambda=1)}}{e^{-u(x', \mathcal{M}_{\text{old}}, \lambda=0)}} \right]$$

$$= \log P_{\text{delete work}} + \log P_{\text{delete energy}} \quad (59)$$

$$\log P_{\text{insert}} = \log e^{-\Delta S_{\text{insert}}[x' \rightarrow x'' | \mathcal{M}_{\text{new}}]} \quad (60)$$

$$= \log \left[e^{-w_{\text{insert}}[x' \rightarrow x'' | \mathcal{M}_{\text{new}}]} \frac{e^{-u(x', \mathcal{M}_{\text{new}}, \lambda=0)}}{e^{-u(x'', \mathcal{M}_{\text{new}}, \lambda=1)}} \right]$$

$$= \log P_{\text{insert work}} + \log P_{\text{insert energy}} \quad (61)$$

165 which we further decompose into **work** and **energy**
166 change contributions for the separate **delete** and **insert**

NCMC stages:

$$\log P_{\text{delete work}} = -w_{\text{delete}}[x \rightarrow x' | \mathcal{M}_{\text{old}}] \quad (62)$$

$$\log P_{\text{delete energy}} = u(x', \mathcal{M}_{\text{old}}, \lambda = 0) - u(x, \mathcal{M}_{\text{old}}, \lambda = 1)$$

$$\log P_{\text{insert work}} = -w_{\text{insert}}[x' \rightarrow x'' | \mathcal{M}_{\text{new}}] \quad (63)$$

$$\log P_{\text{insert energy}} = u(x'', \mathcal{M}_{\text{new}}, \lambda = 1) - u(x', \mathcal{M}_{\text{new}}, \lambda = 0)$$

JDC: Check that the energy component here is really the reduced potential for both GHMC and VV, rather than the total reduced energy. ski

171 With this definition of terms, the overall acceptance proba-
172 bility is therefore given as

$$\begin{aligned} \log P_{\text{accept}} &= \log P_{\text{final}} - \log P_{\text{initial}} + \log P_{\text{chemical}} \\ &+ \log P_{\text{delete work}} + \log P_{\text{delete energy}} \\ &+ \log P_{\text{reverse}} - \log P_{\text{forward}} \\ &+ \log P_{\text{insert work}} + \log P_{\text{insert energy}} \end{aligned} \quad (64)$$

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