Advancing Chemical Potential Calculations of Polymers Introducing the Effect of Conformational Change



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Background

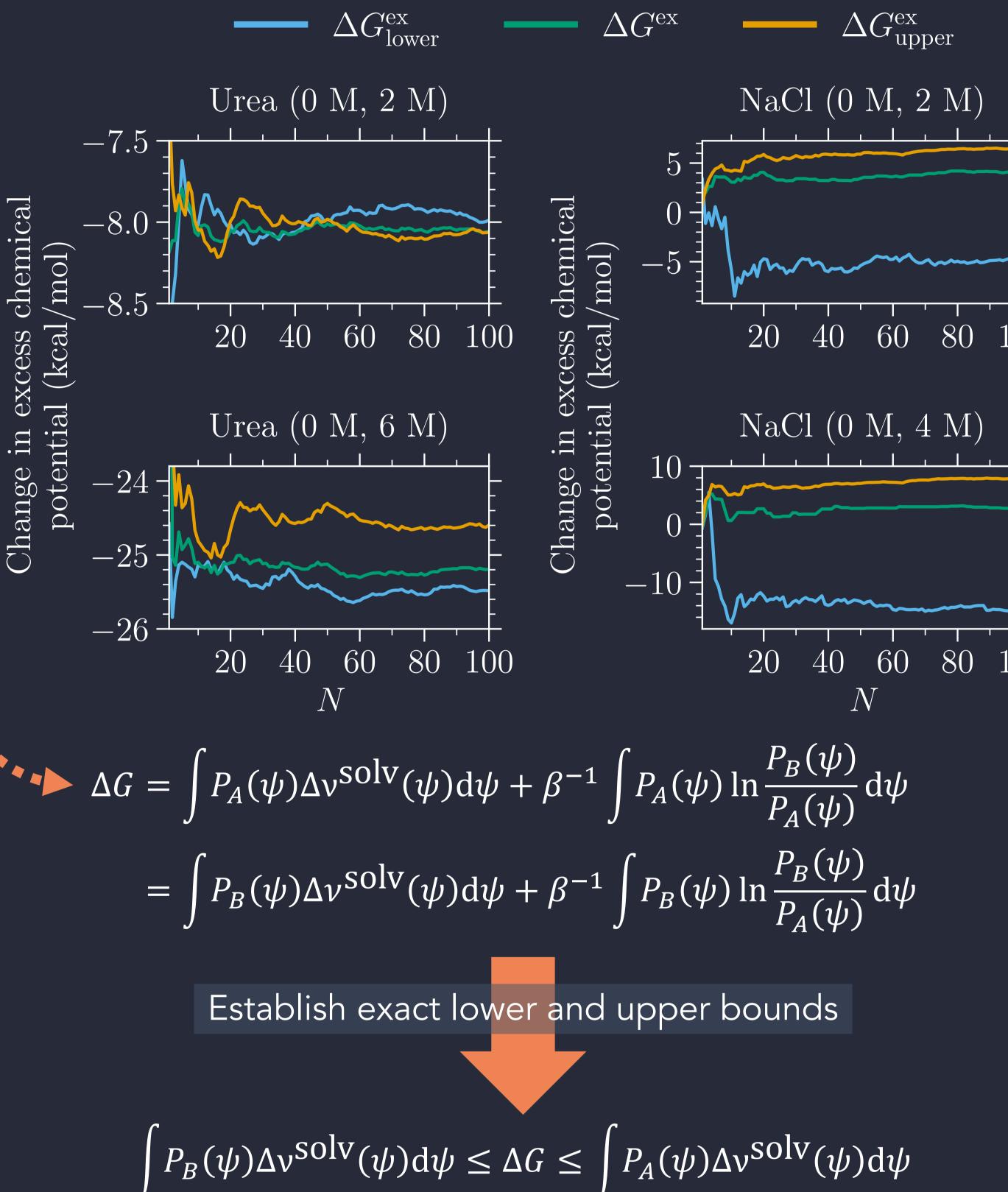
Soft matter systems, like polymers and globular proteins, can be structurally altered by thermal, mechanical, or chemical stress. For instance, proteins typically form compact, folded structures under native conditions but unfold under stress. To describe the thermodynamics of these transformations, obtaining knowledge about the excess chemical potential is essential. We previously developed a methodology to calculate changes in the excess chemical potential due to alterations in liquid composition for various molecular states. However, this method is a linear approximation and does not account for structural transitions induced by changes in solvent composition, which appears as higher-order terms and can be important when the solvent composition is changed much.

Objective

Here we introduce our latest computation scheme for the calculation of the change in free energy of flexible polymers being transferred between solvent compositions formulated though error minimization. The methodology has been applied to describe the solvation of polyethylene glycol (PEG) in aqueous mixtures of urea and NaCl.

Formulation of New BAR-Like Method for Assessing Free **Energy Changes Across Solvent Conditions** "Solute-Only" BAR Usual BAR ψ : solute configuration X: system configuration $v_A^{\text{solv}}(\psi), v_B^{\text{solv}}(\psi)$: solvation free $U_A(\mathbf{X})$, $U_B(\mathbf{X})$: potential energies at states A, B energies in solvent conditions A, B $\Delta v^{\text{solv}}(\psi) = v_B^{\text{solv}}(\psi) - v_A^{\text{solv}}(\psi)$ $\Delta U(\mathbf{X}) = U_B(\mathbf{X}) - U_A(\mathbf{X})$ $P_A(\psi)$, $P_B(\psi)$: distribution $P_A(X)$, $P_B(X)$: distribution functions of ψ functions of X ΔG : free energy change from A to B identity connecting P_A and P_B $P_B(\psi) = P_A(\psi)e^{eta(\Delta G - \Delta v} \mathsf{solv}_{(\psi)})$ $P_B(\mathbf{X}) = P_A(\mathbf{X})e^{\beta(\Delta G - \Delta U(\mathbf{X}))}$ Fermi function N_A , N_B : number of samples in A, B $\mathcal{F}(x) = \frac{1}{1 + \exp(\beta x)}$ $D = \Delta G + \beta^{-1} \ln \frac{N_B}{N_A}$

Convergence of the "Solute-only BAR" Method and Establishing an Upper and Lower Limit of ΔG





 $\sum \mathcal{F}(\Delta U(\mathbf{X}) - D)$

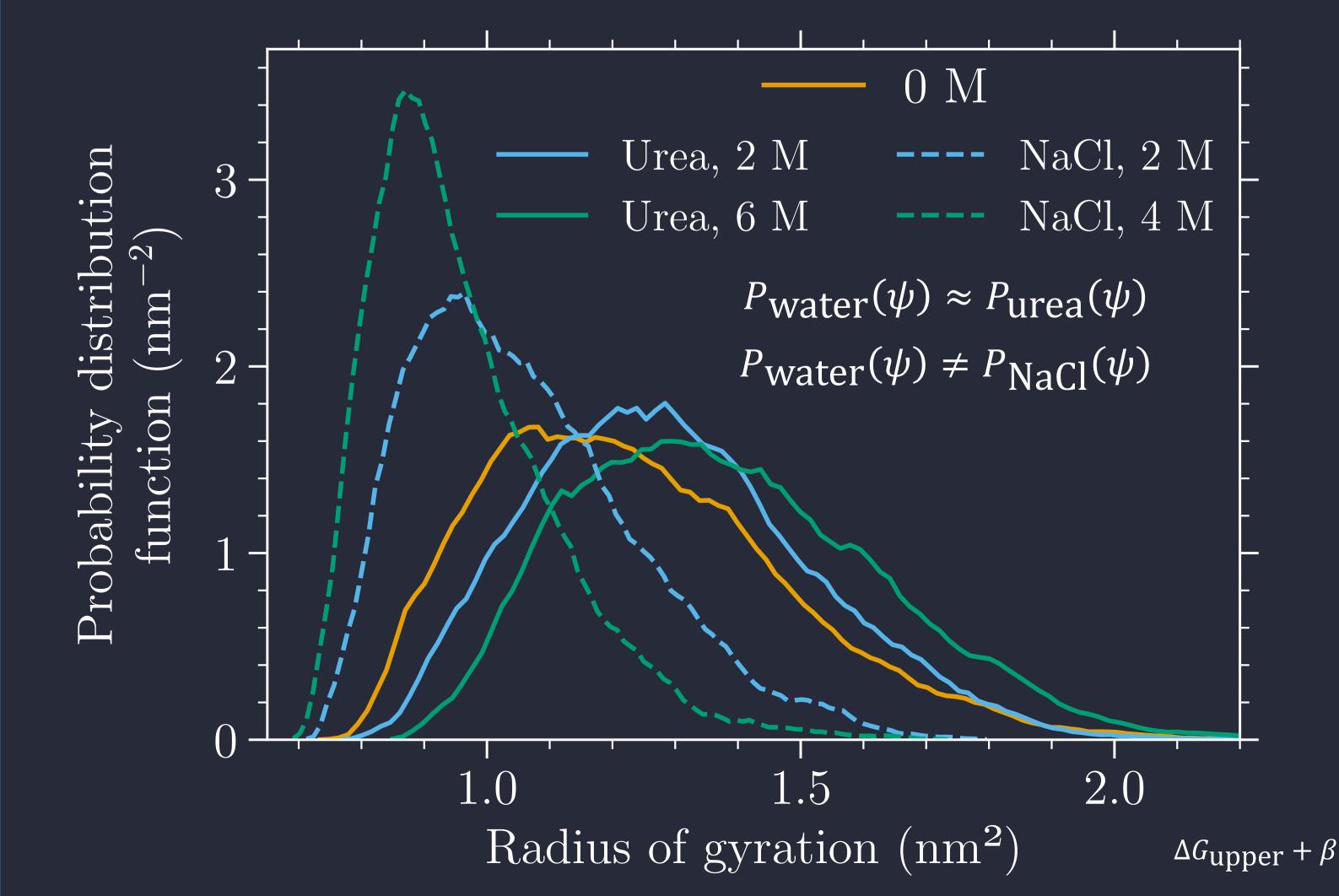
 $= \sum \mathcal{F}(-\Delta U(\mathbf{X}) + D)$

PEG Conformational Ensemble is Invariant with Urea, but Structure is Collapsed with the Addition of NaCl

 $\sum_{\psi \in A} \mathcal{F}\left(\Delta \nu^{\mathsf{solv}}(\psi) - D\right)$

 $= \sum_{i=D} \mathcal{F}\left(-\Delta v^{\mathsf{solv}}(\psi) + D\right)$

Error minimization for ΔG



Relative Entropy is the Statistical Thermodynamic Origin of the Inequality of the Lower and Upper

