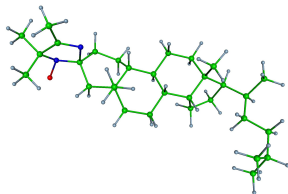
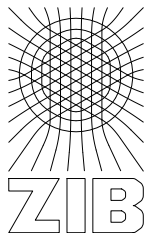


# Computation of equilibrium densities in metastable dynamical systems by domain decomposition

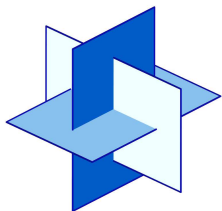
Susanna Kube



ICNAAM 2008



Zuse Institute Berlin



DFG Research Center

MATHEON

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## Cooperations:

FU Berlin:

Biocomputing: Christof Schütte

Applied Mathematics: Caroline Lasser

Computational Molecular Biology: Frank Noé

DFG Research Center **“Matheon”**

Modelling of molecules in **classical MD**:

$$H(q, p) = \frac{1}{2} p^\top M^{-1} p + V(q)$$

$$V = V_{\text{bond}} + V_{\text{angle}} + V_{\text{torsion}} + V_{\text{Coulomb}} + V_{\text{VdW}}$$

$q \in \mathbb{R}^{3s}$ : positions of all atoms,       $p \in \mathbb{R}^{3s}$ : momenta

Corresponding **equations of motion**:

$$\dot{q} = M^{-1} p, \quad \dot{p} = -\nabla V$$

**Invariant density** in the **canonical ensemble** (Boltzmann):

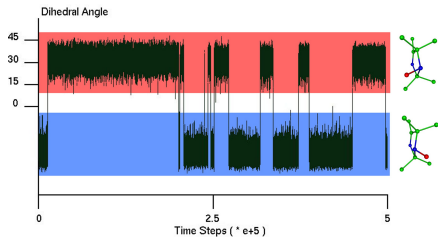
$$\mu(\mathbf{q}, \mathbf{p}) = \underbrace{\frac{1}{Z_p} \exp\left(-\frac{\beta}{2} \mathbf{p}^\top \mathbf{M}^{-1} \mathbf{p}\right)}_{=\eta(\mathbf{p})} \underbrace{\frac{1}{Z_q} \exp(-\beta V(\mathbf{q}))}_{=\pi(\mathbf{q})}, \quad \beta = 1/k_B \mathcal{T}$$

Ensemble averages:

$$\langle A \rangle \equiv \int_{\Omega} A(\mathbf{q}) \pi(\mathbf{q}) d\mathbf{q} \approx \frac{1}{M} \sum_{k=1}^M A(\mathbf{q}_k), \quad \mathbf{q}_k \sim \pi(\mathbf{q})$$

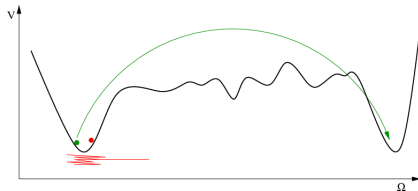
**HMC** (Hamiltonian dynamics with randomized momenta):

$$\mathbf{q}_{n+1} = \Pi_{\mathbf{q}} \Phi^{\mathcal{T}}(\mathbf{q}_n, \mathbf{p}_n), \quad \mathbf{p}_n \sim \eta(\mathbf{p})$$

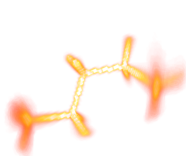


stationary  
density?

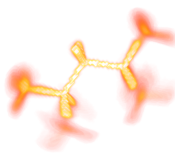
The trajectory gets trapped in valleys of the potential energy surface.



# Example: Pentane



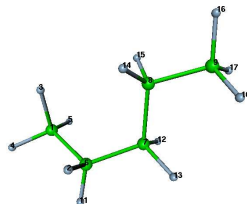
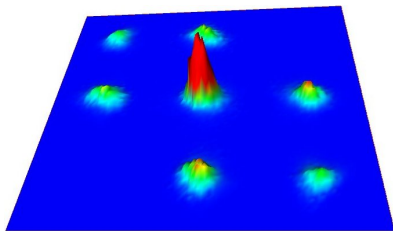
(g+/t)

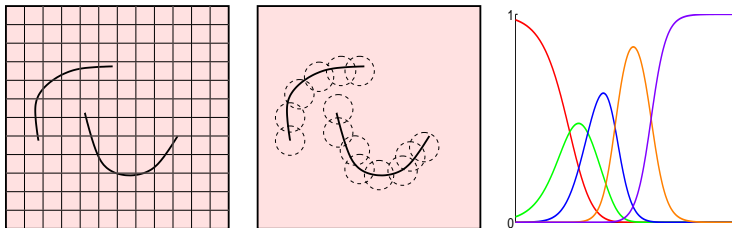


(transition state)



(t/g+)





partition of unity:

$$\sum_{i=1}^N \phi_i(\mathbf{q}) = 1, \quad \phi_i(\mathbf{q}) \geq 0 \quad \forall \mathbf{q} \in \Omega$$

$$S(i,j) \equiv \frac{\int_{\Omega} \phi_i(\mathbf{q}) \phi_j(\mathbf{q}) \pi(\mathbf{q}) d\mathbf{q}}{\int_{\Omega} \phi_i(\mathbf{q}) \pi(\mathbf{q}) d\mathbf{q}}$$

partial densities:

$$\pi_i(\mathbf{q}) \equiv \frac{\phi_i(\mathbf{q})\pi(\mathbf{q})}{\int_{\Omega} \phi_i(\mathbf{q})\pi(\mathbf{q})}$$

$$\pi_i(\mathbf{q}) \propto \exp(-\beta V_i(\mathbf{q})), \quad V_i(\mathbf{q}) = V(\mathbf{q}) - \beta^{-1} \log(\phi_i(\mathbf{q}))$$

Monte Carlo integration:

$$S(i,j) = \int_{\Omega} \phi_j(\mathbf{q})\pi_i(\mathbf{q}) d\mathbf{q} \approx \frac{1}{n_i} \sum_{k=1}^{n_i} \phi_j(\mathbf{q}_k^{(i)}), \quad \mathbf{q}_k^{(i)} \sim \pi_i(\mathbf{q}).$$

$$\pi(q) = \sum_{i=1}^N w_i \pi_i(q)$$

$w_i$  is the **statistical weight** of basis function  $\phi_i(q)$ :

$$w_i \equiv \int_{\Omega} \phi_i(q) \pi(q) dq$$

The vector  $\mathbf{w} = [w_1, \dots, w_N]^T$  is the left **eigenvector** of  $S$  corresponding to the eigenvalue  $\lambda = 1$  (Perron-Frobenius)!

$$\mathbf{w}^T S = \mathbf{w}^T$$

$$\tilde{S} = \bar{S} + E, \quad \tilde{\mathbf{w}}^\top \tilde{S} = \tilde{\mathbf{w}}^\top$$

- ▶ Row-wise correlated random matrices:

$$\mathbb{E}[E(i,j)] = 0 \quad \text{and} \quad \mathbb{E}[E(i,j)E(k,l)] = \delta_{ik} C_i(j,l)$$

- ▶ 1st order Taylor series expansion:

$$\tilde{\mathbf{w}} \doteq \bar{\mathbf{w}} + \sum_{i=1}^N G_i^{\mathbf{w}} E(i,:)^\top, \quad G_i^{\mathbf{w}}(k,l) = \left. \frac{\partial w_k}{\partial s_{il}} \right|_{\bar{A}}, \quad \bar{A} = \bar{S}^\top - I$$

- ▶ probability distribution:  $E(i,:) \sim \text{Dir}(\alpha_i)$
- ▶ parameter estimation from sampling data (ML estimator)

Central limit theorem:

$$\tilde{\mathbf{w}} \sim \text{MVN}(\bar{\mathbf{w}}, \sum_{i=1}^N G_i^{\mathbf{w}} C_i (G_i^{\mathbf{w}})^{\top})$$

Equilibration of sampling effort and uncertainties

$$i = \arg \max_k \|G_k^{\mathbf{w}} C_k (G_k^{\mathbf{w}})^{\top}\|_2$$

Increase number of sampling points in the selected basis function, update the row, and repeat the analysis.

If a maximal allowed number of points is reached  $\rightarrow$  refinement

- ▶ current basis

$$\{\phi_1, \dots, \phi_N\} : \Omega \rightarrow [0, 1]$$

- ▶ function selected for refinement:  $\phi_k(\mathbf{q})$
- ▶ temporal set of basis functions (partition of unity, positivity)

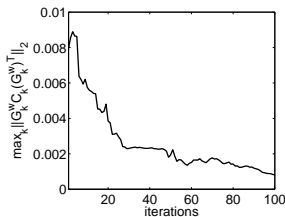
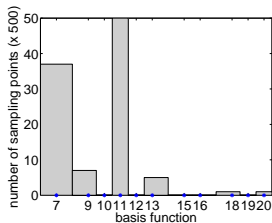
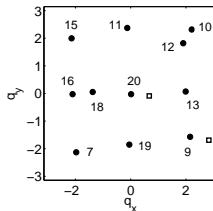
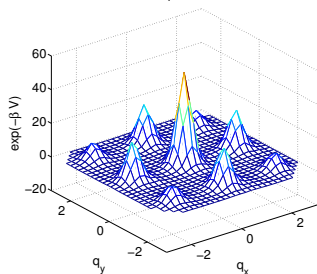
$$\{\tilde{\phi}_{k1}, \dots, \tilde{\phi}_{ks}\} : \Omega \rightarrow [0, 1]$$

- ▶ new basis functions

$$\phi_{ki}(\mathbf{q}) := \phi_k(\mathbf{q})\tilde{\phi}_{ki}(\mathbf{q}), \quad i = 1, \dots, s.$$

$$\{\phi_1, \dots, \phi_{k-1}, \phi_{k+1}, \dots, \phi_N, \phi_{k1}, \dots, \phi_{ks}\}$$

$\beta=0.4$



$$U = \int_{\Omega} V(q)\pi(q)dq = -5.03$$

$$U_0 = -2.63$$

$$U_{100} = -4.45$$

**Thank you for your attention!**

Further information

<http://www.zib.de/Numerik/DrugDesign/index.en.html>

Open positions:

- ▶ Matheon A4 (“Towards a mathematics of biomolecular flexibility”), PhD or Postdoc