

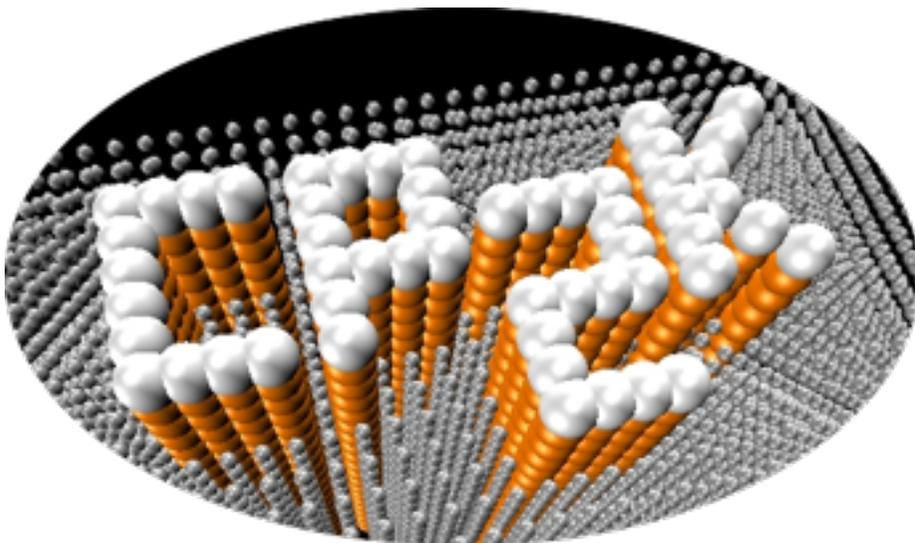
3rd CP2K tutorial: Enabling the Power of Imagination in MD Simulations

June 17-21 2013, Zürich

Sampling Free Energy Surfaces by MD

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<http://www.cp2k.org>

OUTLINE

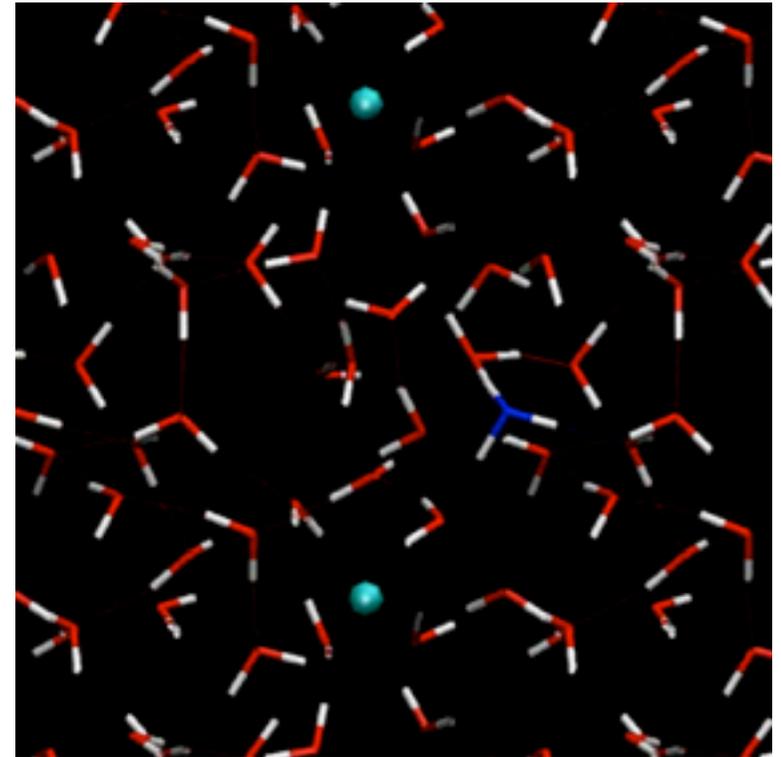
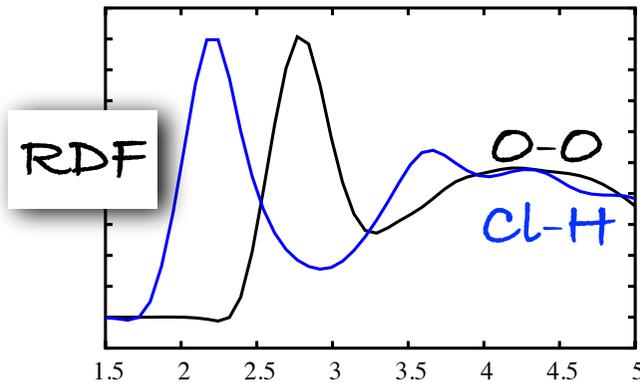
 Free energy in statistical mechanics

 Free energy difference by improving the sampling along the evolution order parameters

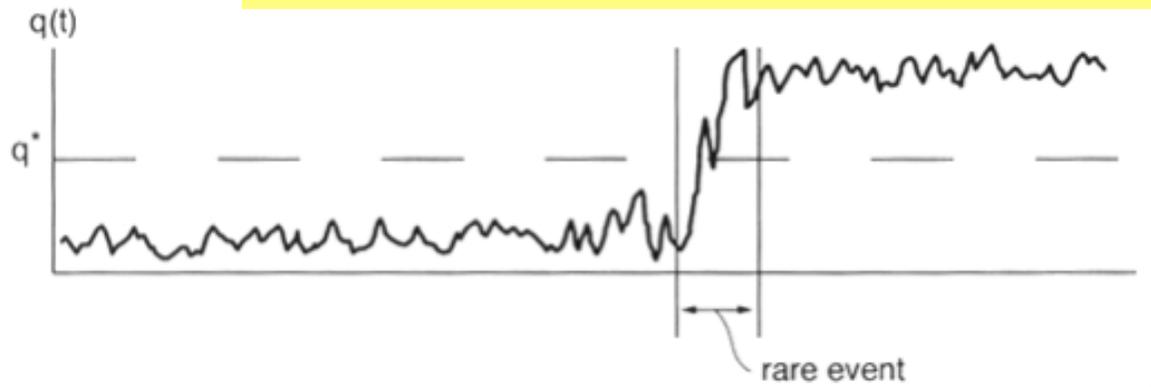
 Enhanced exploration of the configuration space and disclosure of mechanisms of transformation

Complex Processes by MD

- * Choose a suitable model of the system
- * Determine the thermodynamic conditions \Rightarrow Ensemble averages
- * Equilibrium sampling of physical quantities



Predictive power frustrated by sampling fast degrees of freedom with time-steps from < 0.1 fs (CPMD) up to 1 fs (MM)

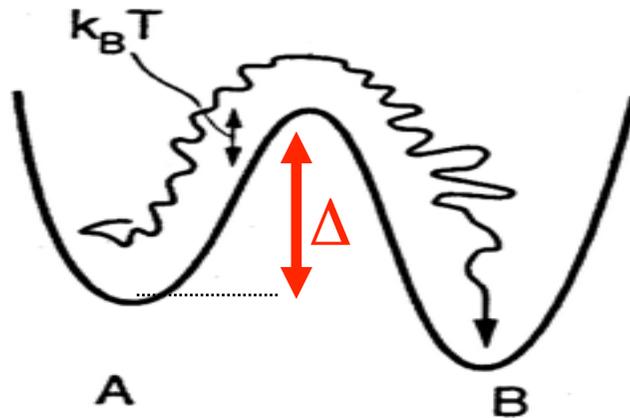


\sim few ns

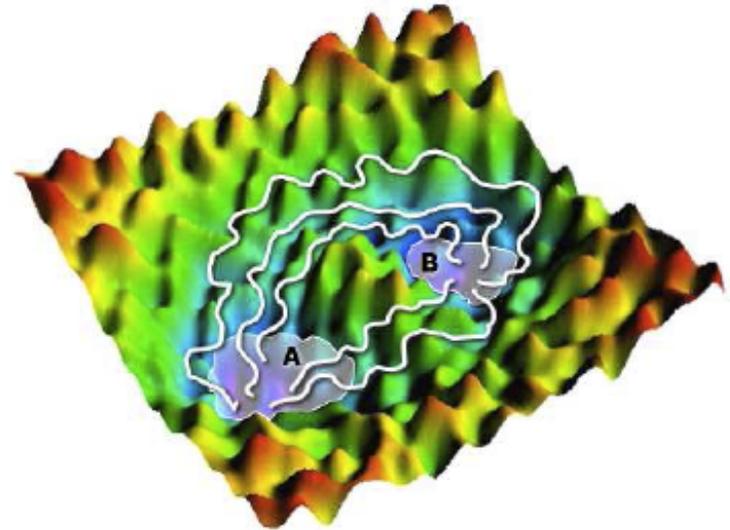
Rare Events

Phase Transitions, Conformational Rearrangements,
Chemical Reactions, Nucleation, Diffusion, Growth, etc.

Activation Energies



Minimum energy pathways



Exploration of configurational space

- * Complex and high dimensional configurational space
- * Intrinsically multidimensional order parameter
- * Multitude of unknown intermediates and products
- * Entropic bottlenecks
- * Unforeseen processes, many irrelevant transition states
- * Diffusive trajectories

Hamiltonian MD

A system of N particles in a volume V is completely determined through its Hamiltonian

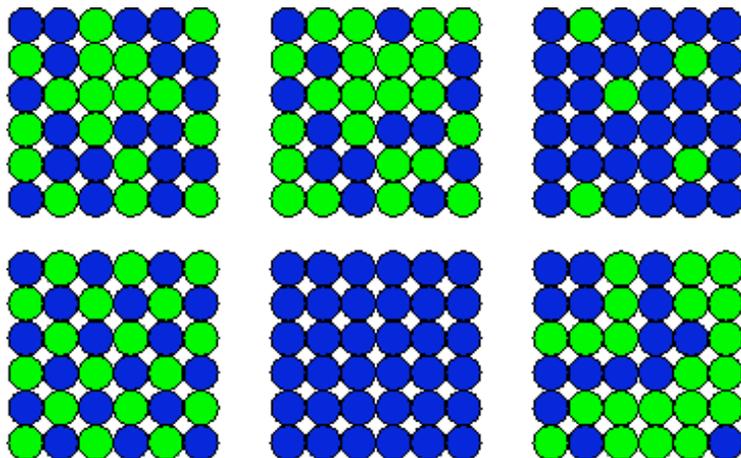
$$\mathcal{H}(\{\mathbf{R}_I\}, \{\mathbf{P}_I\}) = \sum_I \frac{\mathbf{P}_I^2}{2M_I} + U(\{\mathbf{R}_I\})$$

NVE-P total energy and linear momentum are constant of motion

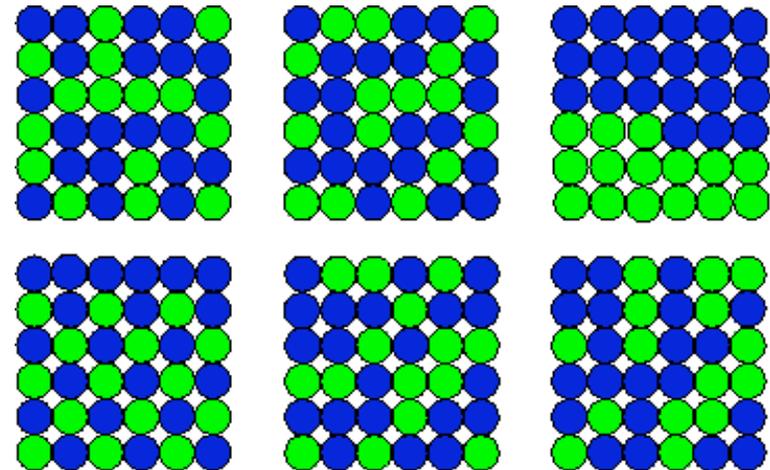
Choice of ensemble: portion of phase space, microstates
Difference in averages and fluctuations

Counters, blue on one side and green on the other, 6×6 checkerboard

Each pattern is a microstate



Subset belonging to macrostate "15 green"



Canonical Partition Function

The Laplace transform of the density of state

$$Q(N, V, T) = \int \exp(-\beta E) \Omega(N, V, E) dE$$

Probability of the macrostate at a given T

$$Q(N, V, T) = \frac{1}{N! h^{3N}} \int \exp[-\beta \mathcal{H}(\mathbf{r}^N, \mathbf{p}^N)] d\mathbf{r}^N d\mathbf{p}^N = \frac{1}{\Lambda(\beta)^{3N} N!} Z(N, V, T)$$

one dimensional integral over E replaced by configurational integral
analytic kinetic part is integrated out

$$Z(N, V, T) = \int e^{-\beta U(\mathbf{r})} d\mathbf{r}$$

**configurational partition
function**

Free Energy

Helmholtz free energy or thermodynamic potential

$$A = -\frac{1}{\beta} \ln Q(N, V, T)$$

Thermodynamics

Statistical Mechanics

$$\Delta A = -\frac{1}{\beta} \ln \left(\frac{Z_1}{Z_0} \right)$$

entropic and enthalpic contributions

$$Q_0 \propto \int_{\Gamma_0} e^{-\beta \mathcal{H}(\mathbf{r}, \mathbf{p})} d\mathbf{r} d\mathbf{p}$$

Macroscopic state 0 corresponds to a portion of the phase space : Γ_0

$$Q_0 \propto \int_{\Gamma} e^{-\beta \mathcal{H}_0(\mathbf{r}, \mathbf{p})} d\mathbf{r} d\mathbf{p}$$

Macroscopic state 0 corresponds to H_0

$$Q_0 \propto \int_{\Gamma} e^{-\beta_0 \mathcal{H}(\mathbf{r}, \mathbf{p})} d\mathbf{r} d\mathbf{p}$$

Macroscopic state 0 corresponds to a value of a macroscopic parameter, e.g T

Perturbation formalism

Reference (0) and target system (1)

$$\mathcal{H}_1(\mathbf{r}, \mathbf{p}) = \mathcal{H}_0(\mathbf{r}, \mathbf{p}) + \Delta\mathcal{H}(\mathbf{r}, \mathbf{p})$$

Probability of finding (0) in configuration (\mathbf{r}, \mathbf{p})

$$\mathcal{P}_0(\mathbf{r}, \mathbf{p}) = \frac{e^{-\beta\mathcal{H}_0(\mathbf{r}, \mathbf{p})}}{\int e^{\mathcal{H}(\mathbf{r}, \mathbf{p})} d\mathbf{r}d\mathbf{p}}$$

Free energy difference

$$\Delta A = -\frac{1}{\beta} \ln \frac{\int e^{-\beta\mathcal{H}_1} d\mathbf{r}^N d\mathbf{p}^N}{\int e^{-\beta\mathcal{H}_0} d\mathbf{r}^N d\mathbf{p}^N} = -\frac{1}{\beta} \ln \frac{\int e^{-\beta\mathcal{H}_0} e^{-\beta\Delta\mathcal{H}} d\mathbf{r}^N d\mathbf{p}^N}{\int e^{-\beta\mathcal{H}_0} d\mathbf{r}^N d\mathbf{p}^N}$$

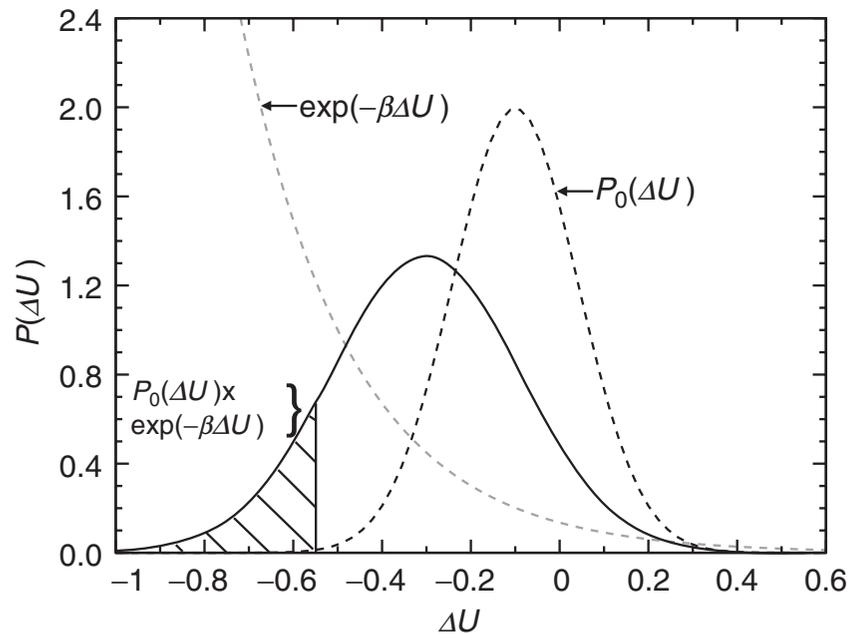
$$\Delta A = -\frac{1}{\beta} \ln \langle \exp [-\beta\Delta\mathcal{H}(\mathbf{r}^N, \mathbf{p}^N)] \rangle_0$$

Integrating out the analytic kinetic part

$$\Delta A_{0,1} = -\frac{1}{\beta} \ln \langle e^{-\beta\Delta U} \rangle_0 \quad \langle \mathcal{F}(\mathbf{r}, \mathbf{p}) \rangle_1 = \frac{\langle \mathcal{F} e^{-\beta\Delta U} \rangle_0}{\langle e^{-\beta\Delta U} \rangle_0}$$

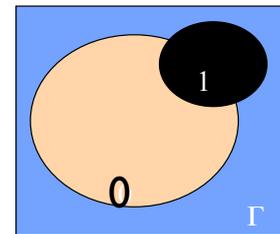
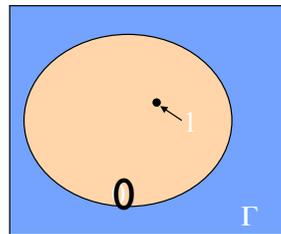
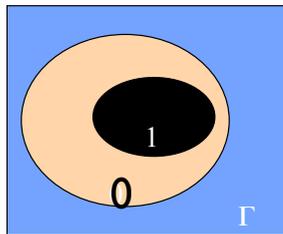
Limitations

$$\Delta A = -\frac{1}{\beta} \ln \int \exp[-\beta \Delta U] \mathcal{P}_0(\Delta U) d\Delta U$$



Shifted function
 Low- ΔU tail is poorly sampled
 low statistical accuracy
 but important contribution to ΔA

Accuracy \Rightarrow target and reference systems are similar \Rightarrow overlapping regions



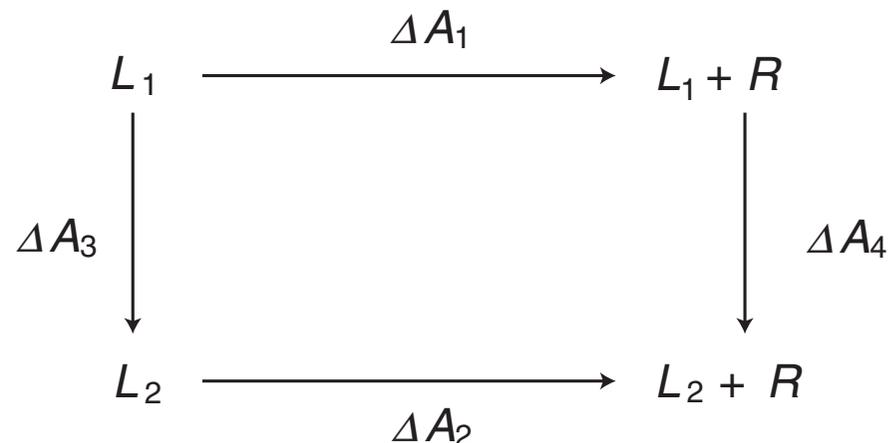
insufficient statistics or incomplete overlap \Rightarrow enhanced sampling

Alchemical Transformations

Protein ligand binding, host-guest chemistry, solvation properties, ...

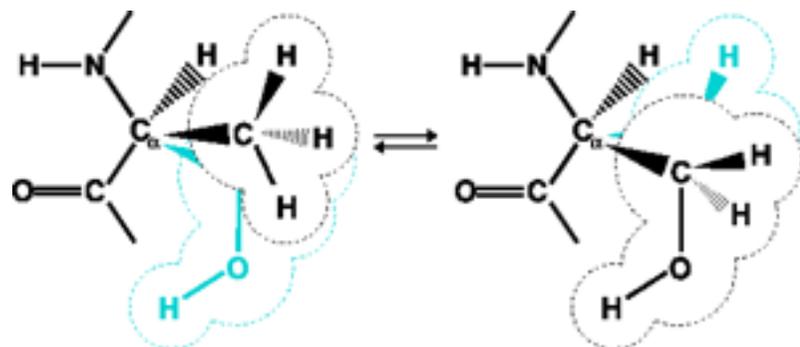
Transformation (0)→(1) as series of non-physical, intermediate states along a pathway characterized by the

“coupling parameter” λ



Free energy as continuous function of λ through $\mathcal{H}(\lambda)$

$$\mathcal{H}(\lambda) = \mathcal{H}_{\text{env}} + \lambda \mathcal{H}_0 + (1 - \lambda) \mathcal{H}_1$$



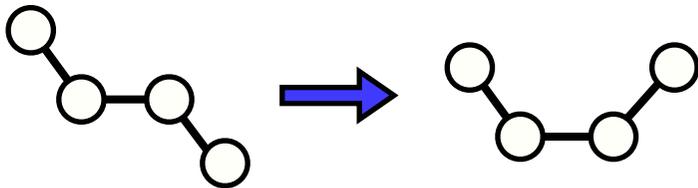
Point mutation of alanine into serine:
coexistence without seeing each other

Interaction with side chain tuned through λ

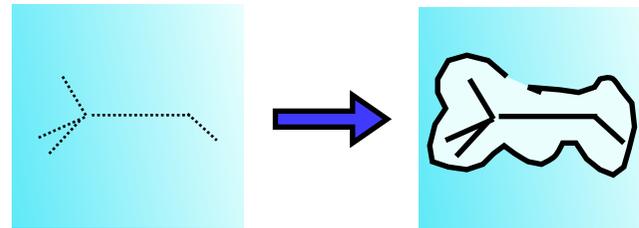
Order Parameters

Variables chosen to describe changes in the system

- **Reaction coordinate** : the order parameter corresponds to the pathway along which the transformation occur in nature
- **Collective variable** : fully represented as function of coordinates
- Indicating intermediate stages of the transformation: **mutation point**



torsion angle



annihilation non-bonded

- Different possible definitions of OP
- Effects on accuracy and efficiency of ΔA calculations
- Set up of system with desired values of OP
- Smoothness of the simulated path

Extended Ensemble

Select parameters, continuous functions of coordinates $\hat{\xi}_i(\mathbf{r}^N)$

Density of States

$$\Omega_{\xi}(N, V, E, \xi) = \int \delta[\mathcal{U}(\mathbf{r}^N) - E] \left(\prod_i \delta[\hat{\xi}_i(\mathbf{r}^N) - \xi_i] \right) d\mathbf{r}^N$$

$$\xi = \{\xi_i\}$$

Canonical Partition Function

$$Q_{\xi}(N, V, T, \xi) = \int e^{-\beta \mathcal{U}(\mathbf{r}^N)} \left(\prod_i \delta[\hat{\xi}_i(\mathbf{r}^N) - \xi_i] \right) d\mathbf{r}^N$$

Free Energy

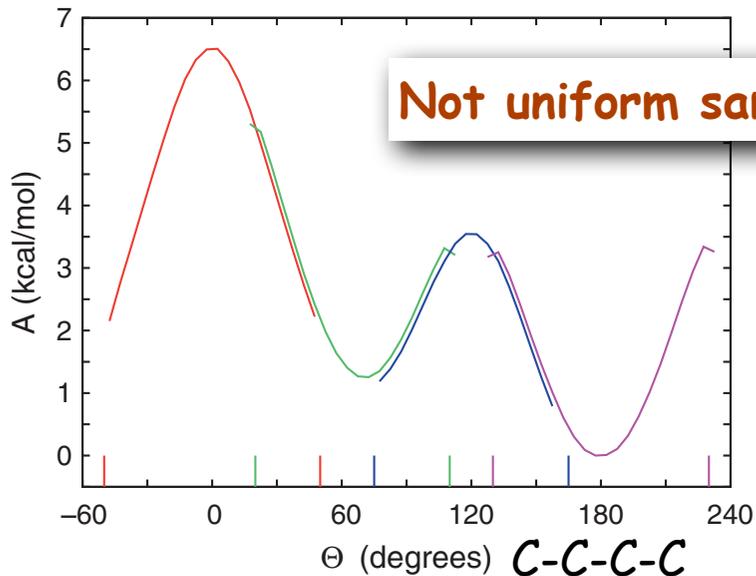
$$A_{\xi} = -\frac{1}{\beta} \ln Q_{\xi}$$

$\hat{\xi}_i(\mathbf{r}^N)$ must distinguish among metastable states

select specific configurations in the partition function

Stratification Scheme

Free energy butane isomerization



Probability distribution of the order parameter

$$\Delta A(\xi) = A(\xi_1) - A(\xi_0) = -\beta^{-1} \ln \frac{\mathcal{P}(\xi_1)}{\mathcal{P}(\xi_0)}$$

Histogram of M bins $\delta\xi = (\xi_1 - \xi_0)/M$

$$\mathcal{P}(\xi_0 + (i - 0.5)\delta\xi) = \frac{f_i}{\sum_j f_j}$$

Restrain the system within a window by harmonic potential

Overlapping windows

Efficient sampling

$$\tau = L\tau_w \propto \frac{(\xi_1 - \xi_0)^2}{LD_\xi}$$

Reconstruct the full probability by matching

Importance Sampling

Non-Boltzmann sampling to enhance the probability of important regions

$$\mathcal{P}^{(w)}(\xi, T) = \frac{\int w(\hat{\xi}(\mathbf{r}^N)) \exp[-\beta\mathcal{U}(\mathbf{r}^N)] \delta[\xi - \hat{\xi}(\mathbf{r}^N)] d\mathbf{r}^N}{\int w(\hat{\xi}(\mathbf{r}^N)) \exp[-\beta\mathcal{U}(\mathbf{r}^N)] d\mathbf{r}^N} \quad \text{positive bias } w(\xi)$$

Free energy differences

$$\Delta A^{(w)}(\xi) = -\beta^{-1} \ln \frac{w(\xi_0) \mathcal{P}^{(w)}(\xi, T)}{w(\xi) \mathcal{P}^{(w)}(\xi_0, T)} = -\beta^{-1} \left[\ln \frac{\mathcal{P}^{(w)}(\xi, T)}{\mathcal{P}^{(w_0)}(\xi, T)} + \ln w(\xi_0) - \ln w(\xi) \right]$$

Biasing potential

$$w(\xi) = \exp[-\beta V(\xi)] \quad \mathcal{U}^{(b)}(\mathbf{r}^N) = \mathcal{U}(\mathbf{r}^N) + V(\xi(\mathbf{r}^N))$$

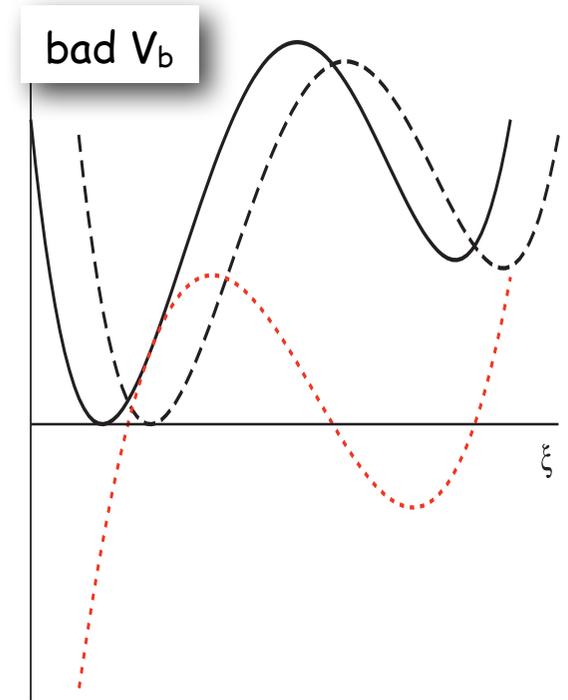
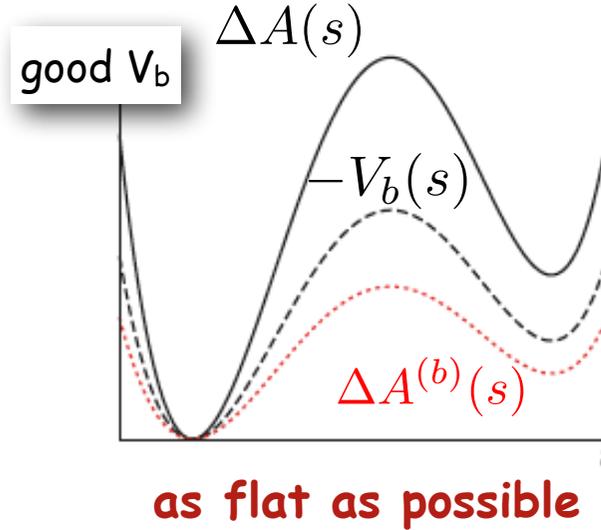
Umbrella Potential
connecting different regions of the phase space

umbrella sampling

Modify the underlying potential to obtain a uniform sampling : $V_b(s) = -\Delta A(s)$

$$U^{(p)}(x) = U(x) + V_b(s(x))$$

First guess
&
iterative improvement



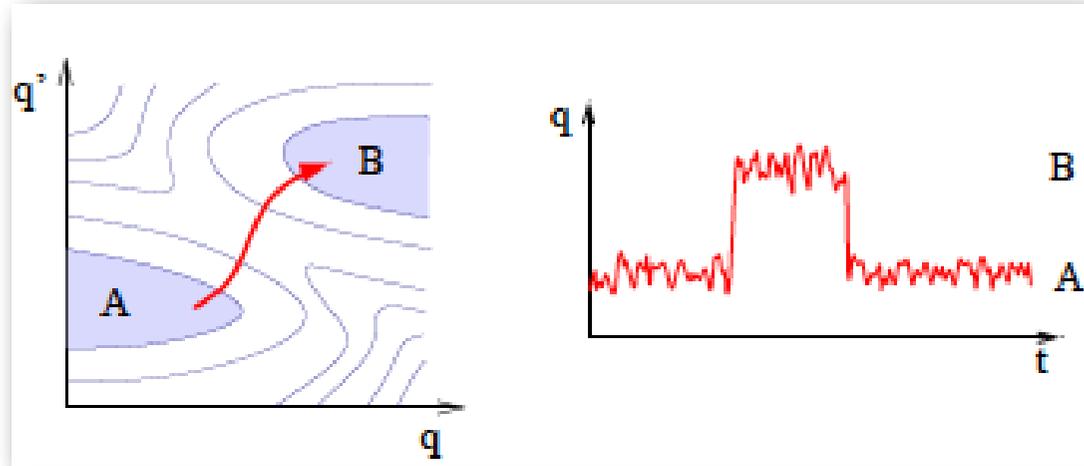
$$\begin{aligned} \mathcal{P}^{(b)}(s) &= \frac{1}{Z^{(b)}} \int dx \exp(-\beta(U(x) + V_b(s(x)))) \delta(s - s(x)) \\ &= \frac{Z}{Z^{(b)}} \exp(-\beta V_b(s)) \frac{1}{Z} \int dx \exp(-\beta U(x)) \delta(s - s(x)) \\ &= \frac{Z}{Z^{(b)}} \exp(-\beta(V_b(s) + A(s))) \end{aligned}$$

Probability in the
presence of the bias
&
un-biasing

$$A(s) = -\frac{1}{\beta} \ln \mathcal{P}^{(b)}(s) - V_b(s) - \frac{1}{\beta} \ln \left(\frac{Z}{Z^{(b)}} \right)$$

Good Coordinates for Pathways

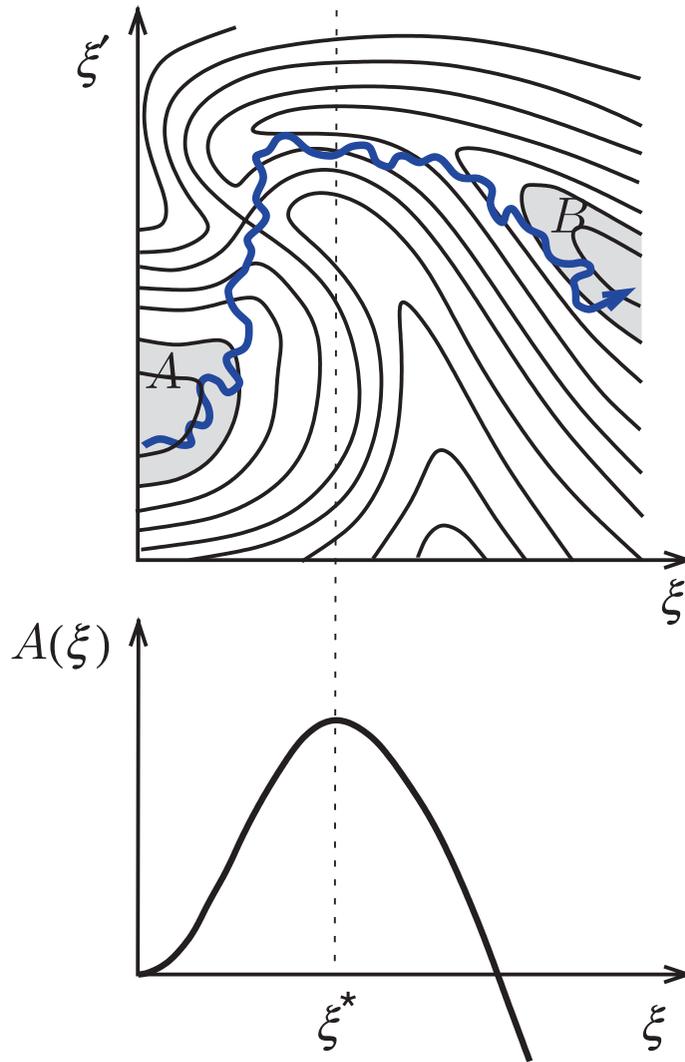
Capture the essential physics include all relevant DoF and properly describes the dynamics



q distinguishes between A and B but might fail in describing essential aspects of the transition

- Discriminate configurations of reactants and products and intermediates
- Characterisation of the mechanisms of transition
- Reversibility
- Fast equilibration of orthogonal DoF (no relevant bottlenecks)

Hypothetical 2D Free Energy Landscape



Not including
important DoF leads
to wrong
interpretation

Some simple collective variables

Derivable function of the degrees of freedom to which a given value can be assigned

● Distance

$$|\mathbf{R}_I - \mathbf{R}_J|$$

● Angle

$$\theta(\mathbf{R}_I, \mathbf{R}_J, \mathbf{R}_k)$$

● Dihedral

$$\Theta(\mathbf{R}_I, \mathbf{R}_J, \mathbf{R}_k, \mathbf{R}_L)$$

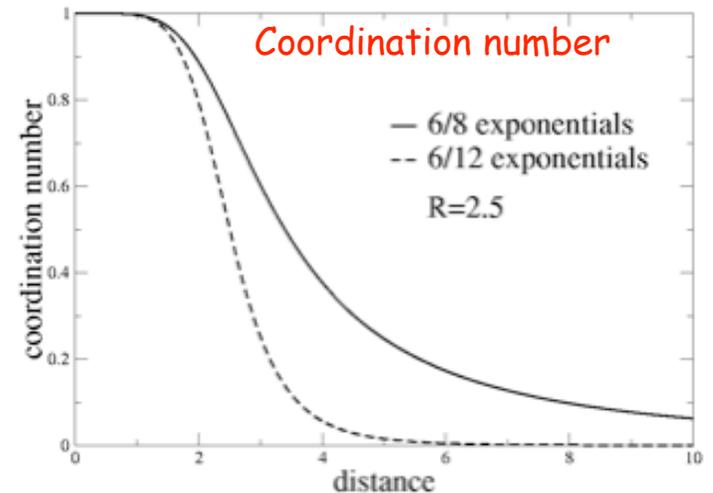
● Difference of distances $|\mathbf{R}_I - \mathbf{R}_J| - |\mathbf{R}_J - \mathbf{R}_K|$

● Generalised coordination number

$$C_{L_1 L_2} = \frac{1}{N_{L_1}} \sum_{j=1}^{N_{L_1}} \left\{ \sum_{i=1}^{N_{L_2}} \frac{1 - \left(\frac{r_{ij}}{r_0}\right)^n}{1 - \left(\frac{r_{ij}}{r_0}\right)^m} \right\}$$

● Generalised displacement

$$D_{L_1 L_2}^{[klm]} = \frac{1}{N_{L_1}} \sum_{i \in L_1} \mathbf{d}_i \cdot \hat{\mathbf{v}}_{[klm]} - \frac{1}{N_{L_2}} \sum_{j \in L_2} \mathbf{d}_j \cdot \hat{\mathbf{v}}_{[klm]}$$



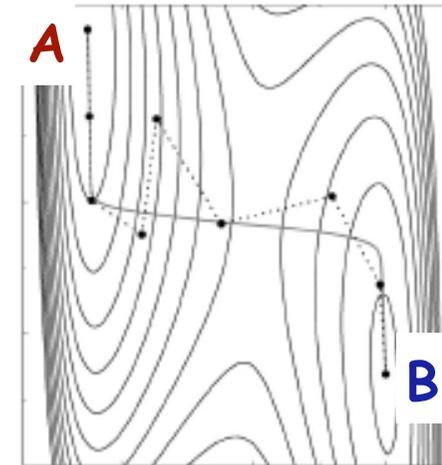
Path Collective Variables

Knowing end-points or a full approximate path (NEB)

☀ RMSD

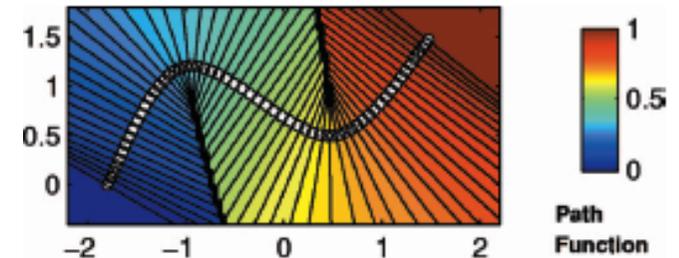
$$\mathcal{R}_k(\mathbf{R}) = \sqrt{\frac{\sum_i (\mathbf{R}_i - \mathbf{R}_i^{(k)})^2}{N}} \quad \mathcal{R}_k(\mathbf{R}) = \sqrt{\frac{\sum_j^{N_{\text{dist}}} (d_j(\mathbf{R}) - d_j^{(k)}(\mathbf{R}))^2}{N_{\text{dist}}}}$$

$$s(\mathbf{R}) = \sum_k [\mathcal{R}_k(\mathbf{R})]^{n_k} c_k$$



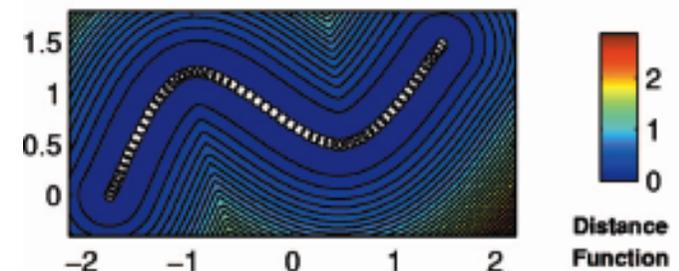
☀ Position along the path

$$s(\mathbf{R}) = \frac{1}{P-1} \frac{\sum_{k=1}^P k e^{-\lambda \|\mathbf{S}(\mathbf{R}) - \mathbf{S}(k)\|^2}}{\sum_{k=1}^P e^{-\lambda \|\mathbf{S}(\mathbf{R}) - \mathbf{S}(k)\|^2}}$$

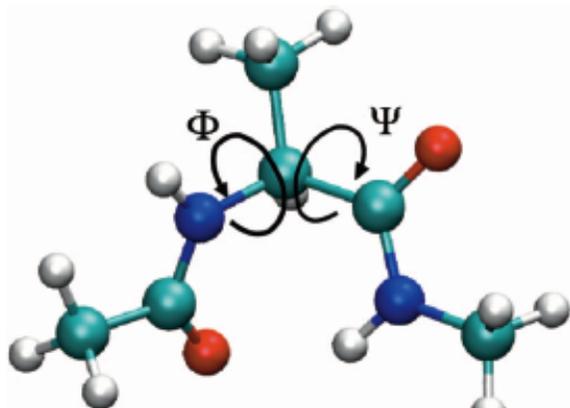


☀ Distance from the path

$$z(\mathbf{R}) = \frac{-1}{\lambda} \ln \left(\sum_{k=1}^P e^{-\lambda \|\mathbf{S}(\mathbf{R}) - \mathbf{S}(k)\|^2} \right)$$



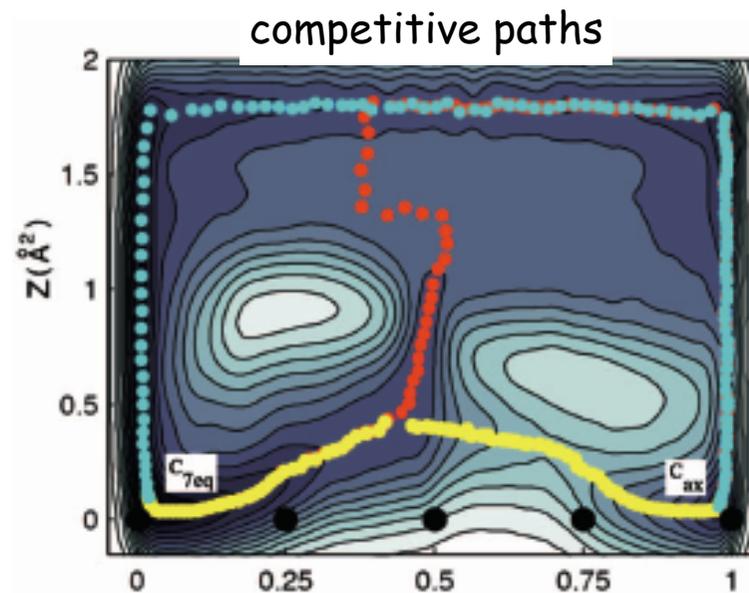
Dialanine in vacuum



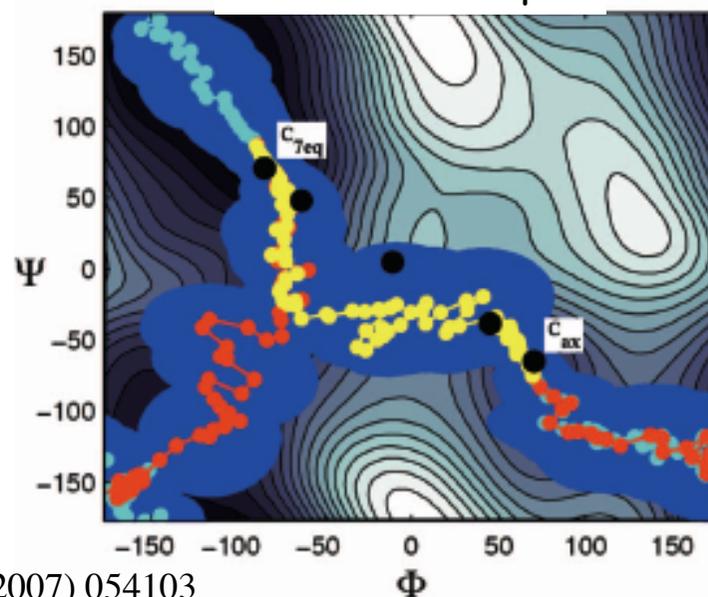
linear interpolation for the initial path

s and z in terms of coordinates

multidimensional path



Ramachandran plot



CP2K input for CV

In SUBSYS add one section per CV

```
&COLVAR
  &DISTANCE
    AXIS X
    ATOMS 1 4
  &END DISTANCE
&END COLVAR
```

```
&COLVAR
  &DISTANCE_FUNCTION
    ATOMS 4 6 6 1
    COEFFICIENT -1.00000
    # distance 1 = ( 4 - 6 )
    # distance 2 = ( 6 - 1 )
  &END DISTANCE_FUNCTION
&END COLVAR
```

```
&COLVAR
  &COORDINATION
    KINDS_FROM N
    KINDS_TO O
    R_0 [angstrom] 1.8
    NN 8
    ND 14
  &END COORDINATION
&END COLVAR
```

```
&COLVAR
  &RMSD
    &FRAME
      COORD_FILE_NAME planar.xyz
    &END
    &FRAME
      COORD_FILE_NAME cage.xyz
    &END
    SUBSET_TYPE LIST
    ATOMS 1 3 5 6 8 9
    ALIGN_FRAMES T
  &END
&END
```

Constraints and Restraints

In MOTION add one section per constraint

```
&CONSTRAINT
&COLLECTIVE
  COLVAR 1
  INTERMOLECULAR
  TARGET    5.
  TARGET_GROWTH 1.1
  TARGET_LIMIT 10.
&END COLLECTIVE
&END CONSTRAINT
```

```
&COLLECTIVE
  TARGET [deg] 0.0
  MOLECULE 1
  COLVAR 1
  &RESTRAINT
    K [kcalmol] 4.90
  &END
&END COLLECTIVE
```

Geometrical Constraints

Implicit functions of the degrees of freedom of the system

$$\sigma(\{\mathbf{R}_I\}, \mathbf{h}, \Psi) = 0 \quad \dot{\sigma}(\{\mathbf{R}_I\}, \mathbf{h}, \Psi) = 0$$

- ☀ To freeze fast degrees of freedom and increase the time step: e.g., intra-molecular bonds by distance constraints
- ☀ To explore only a sub-region of the conformational space
- ☀ As reaction coordinates : constrained dynamics and thermodynamic integration
- ☀ To prevent specific events or reactions

Lagrange formulation for simple constraints, functions of \mathbf{R}_I

$$\mathcal{L}'(\{\mathbf{R}_I\}, \{\mathbf{P}_I\}) = \mathcal{L}(\{\mathbf{R}_I\}, \{\mathbf{P}_I\}) - \sum_{\alpha} \lambda_{\alpha} \sigma(\{\mathbf{R}_I\})$$

The Lagrange multipliers ensure that positions and velocities satisfy the constraints

Shake-Rattle algorithm

Modified velocity Verlet scheme by additional **constraint forces**

- First update of velocities (first half step) and positions

$$V'_I = V_I(t) + \frac{\delta t}{2M_I} F_I(t) \quad R'_I = R_I(t) + \delta t V'_I$$

- Positions' correction by **constraint forces**

$$R_I(t + \delta t) = R'_I + \frac{\delta t^2}{2M_I} g_I^{(p)}(t)$$

- Calculation of the new forces $F_I(t+\delta t)$

- Update of velocity (second half step)

$$V_I(t + \delta t) = V'_I + \frac{\delta t}{2M_I} [F_I(t + \delta t) + g_I^{(v)}(t + \delta t)]$$

- Correction by the constraint forces

Constraint Forces

$$g_I^{(p)}(t) = - \sum_{\alpha} \lambda_{\alpha}^{(p)} \frac{\partial \sigma_{\alpha}(\{R_I\})}{\partial R_I}$$

$$e_{\alpha}(\{\lambda_{\gamma}\}) = - \sum_{\beta} \mathbf{J}_{\alpha\beta}^{-1} \sigma_{\beta}(\{\lambda_{\gamma}\}) \quad \mathbf{J}_{\alpha\beta} = \frac{\partial \sigma_{\alpha}(\{\lambda_{\gamma}\})}{\partial \lambda_{\beta}}$$

Set of non-linear equations **solved iteratively**

$$g_I^{(v)}(t) = - \sum_{\alpha} \lambda_{\alpha}^{(v)} \frac{\partial \sigma_{\alpha}(\{R_I\})}{\partial R_I}$$

$$\sum_I \frac{\partial \sigma_{\alpha}}{\partial \mathbf{R}_I} \mathbf{v}_I = \sum_I \frac{\partial \sigma_{\alpha}}{\partial \mathbf{R}_I} \cdot \mathbf{v}'_I + \sum_{\beta} \left(\sum_I \frac{\delta t^2}{2M_I} \frac{\partial \sigma_{\alpha}}{\partial \mathbf{R}_I} \frac{\partial \sigma_{\beta}}{\partial \mathbf{R}_I} \right) \lambda_{\beta}^v = 0$$

Thermodynamic Integration

$$A(\xi_1) - A(\xi_0) = \int_{\xi_0}^{\xi_1} \frac{dA}{d\xi} d\xi$$

along a one dimensional
generalized coordinate $\xi(\mathbf{x})$
Path-independent

Potential of Mean Force exerted on ξ

$$(\mathbf{x}, \mathbf{p}) \Rightarrow (\xi, q_1 \dots q_{N-1}, p^\xi \dots p_{N-1}^q)$$

generalized coordinate to simplify derivative

$$\frac{dA}{d\xi} = \frac{\int \frac{\partial \mathcal{H}}{\partial \xi} e^{-\beta \mathcal{H}} dq_1 \dots dq_{N-1} dp^\xi \dots dp_{N-1}^q}{\int e^{-\beta \mathcal{H}} dq_1 \dots dq_{N-1} dp^\xi \dots dp_{N-1}^q} = \left\langle \frac{\partial \mathcal{H}}{\partial \xi} \right\rangle_\xi$$

instantaneous force on ξ

force at ξ , averaged over
fluctuations of other DoF

$$\left\langle \frac{\partial \mathcal{H}}{\partial \xi} \right\rangle_\xi = \left\langle \frac{\partial U}{\partial \xi} - \frac{1}{\beta} \frac{\partial \ln |\mathbf{J}|}{\partial \xi} \right\rangle$$

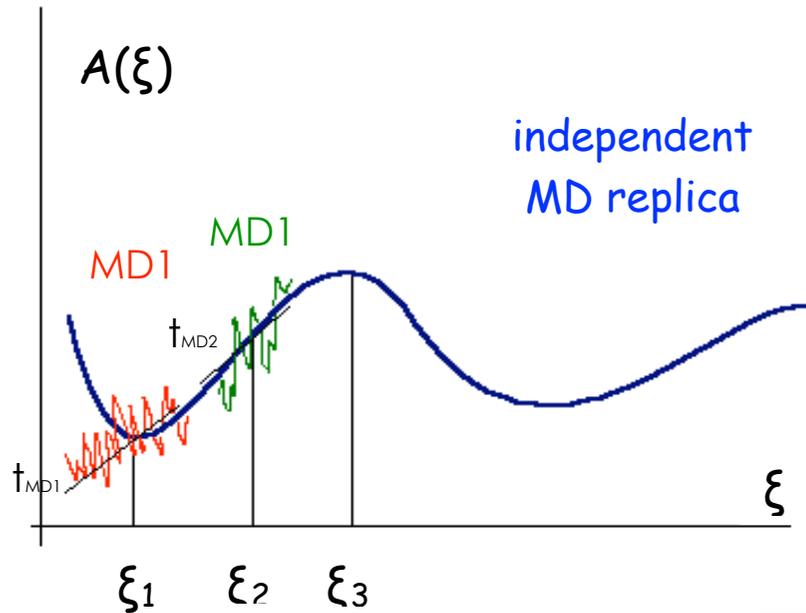
$$[\mathbf{J}]_{ij} = \frac{\partial x_i}{\partial q_j}$$

mechanical + entropic

Blue Moon Ensemble

Series of constrained MD simulations

$$\mathcal{H}^\lambda = \mathcal{H} + \lambda(\xi - \xi(\mathbf{r}))$$



$$-\lambda \nabla(\xi - \xi(\mathbf{r}))$$

mean force acting on the system to hold ξ constant

$$\dot{\xi} = 0 \quad : \quad p^\xi(\mathbf{q}, \mathbf{p}^q)$$

un-constrained $\langle \dots \rangle =$ constrained corrected $\langle \dots \rangle^F$

Fixman Potential

$$\mathcal{H}_F^\lambda = \mathcal{H}^\lambda + \frac{1}{2\beta} \ln Z_\xi$$

$$Z_\xi = \sum_i \frac{1}{m_i} \left(\frac{\partial \xi}{\partial x_i} \right)^2$$

$$\frac{dA}{d\xi} = \langle \lambda_F \rangle_{\xi \dot{\xi}}^F$$

mean force acting on ξ related to external force to hold ξ constant

...some difficulties

MD performed at fixed ξ , collecting statistics of the force acting on $\xi \Rightarrow \lambda \nabla \xi$, **constraint force**

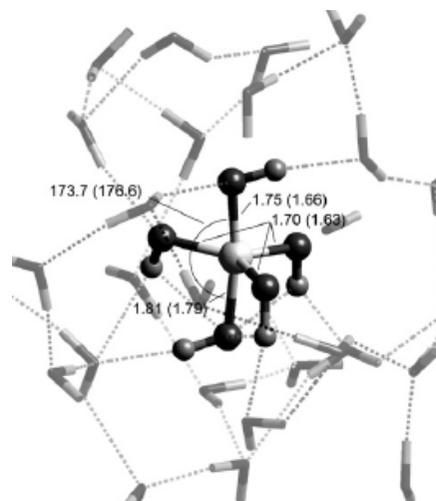
- Many estimates at ξ to reduce statistical errors
- Many windows to get accurate integrals
- May not be easy to prepare by hand the system at given ξ
- Different possible pathways: the starting configuration selects one path, but crossing is rare, $\xi(r)=\xi$ partially sampled or rate limiting
- Multidimensionality (more coordinates) too expensive

De-protonation of $P(OH)_5$

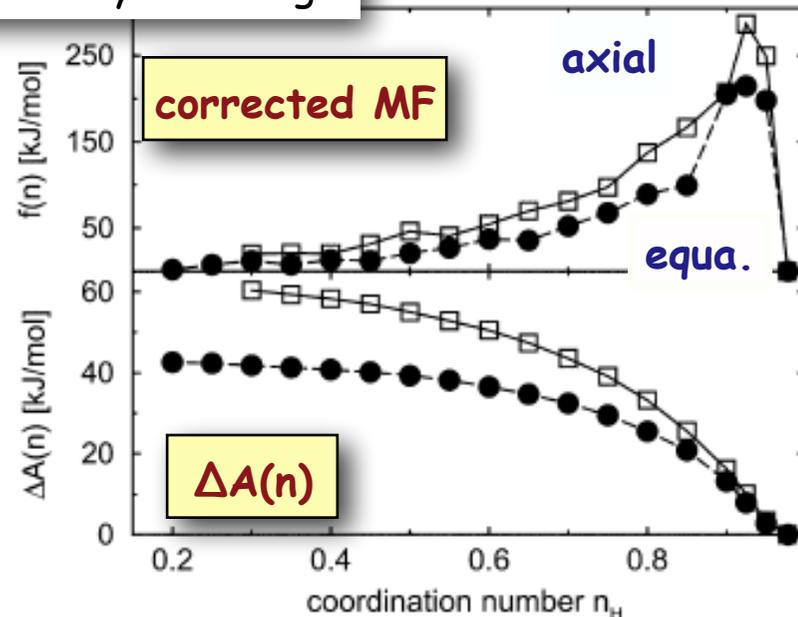
Order Parameter : coordination of a specific donor phosphorane O ($r_0=1.3 \text{ \AA}$)

equivalent PO in solvated $P(OH)_5$

Pronounced equatorphlicity of O^- in $PO_5H_4^-$



gradually reducing n



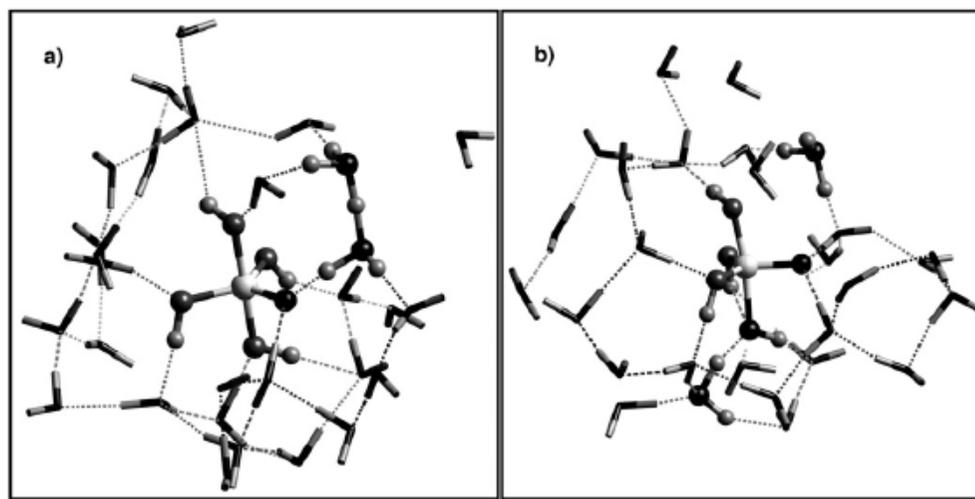
* de-protonation and formation of Zundel ion

* H_3O^+ breaks loose

* chain of proton transfers

* protonation of axial OH

* formation of H_3PO_4 and subsequent de-protonation



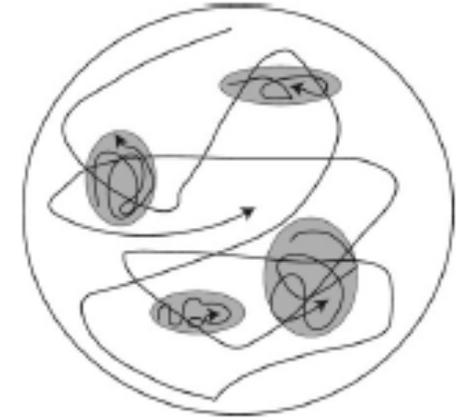
Parallel Tempering

Running **M replica** at different T
high T to explore large part of the phase space
low T to sample precisely local minima

$$T_1 < T_2 < \dots < T_M$$

Equilibration + swap attempt + rescaling of velocities

exchange ensures access to all local minima at low T

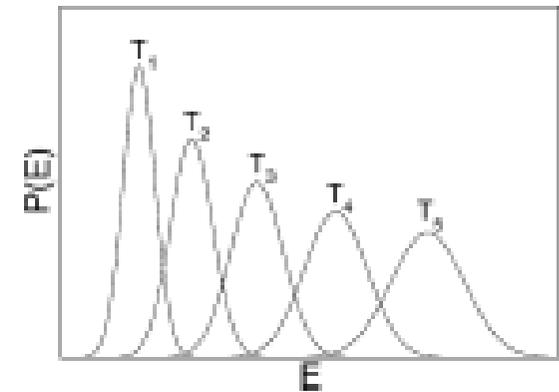


Swapping acceptance

$$\alpha_{IJ} = \min \left\{ 1, e^{-(\beta_I - \beta_J)(U(\mathbf{r}_I) - U(\mathbf{r}_J))} \right\}$$

likelihood that two replicas have overlap in phase space

Energy histogram



$$\sigma \propto (N)^{1/2}$$

M and **ΔT** strongly affect sampling efficiency and computational costs

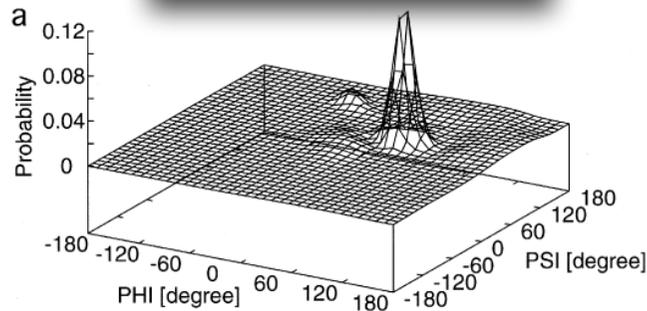
Distribution dihedral of Gly-2

Penta-peptide system in gas phase(Try-Gly-Gly-Phe-Met), all-atom AMBER

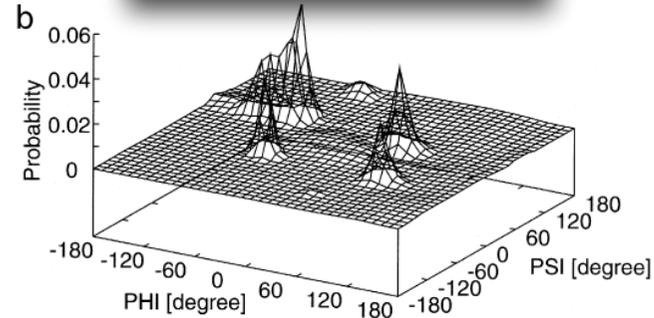
Eight temperatures
replica exchange
simulation, over 1 ns

Pair of temperatures	Acceptance ratio
200 ↔ 239 K	0.160
239 ↔ 286 K	0.149
286 ↔ 342 K	0.143
342 ↔ 409 K	0.139
409 ↔ 489 K	0.142
489 ↔ 585 K	0.146
585 ↔ 700 K	0.146

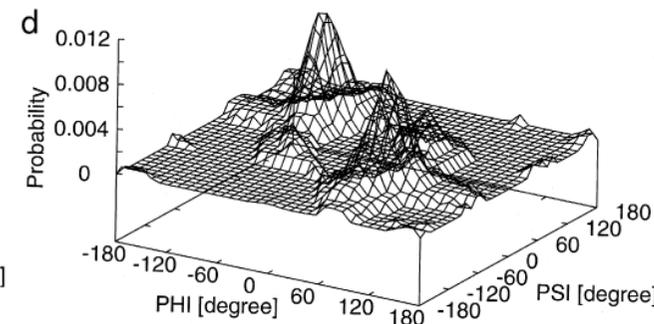
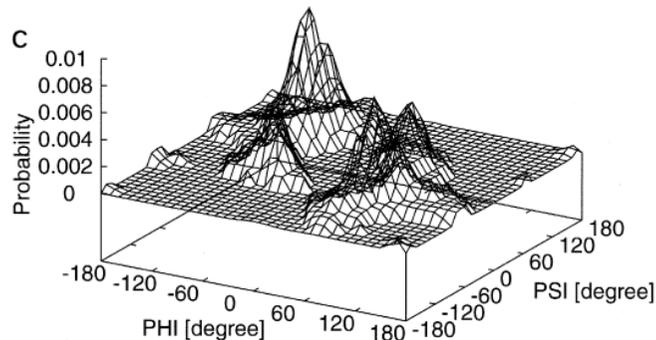
Regular NVT MD



Replica exchange



200 K



700 K

Transition Path Sampling

- Statistical, reaction-coordinate free description of pathways connecting long-lived stable states

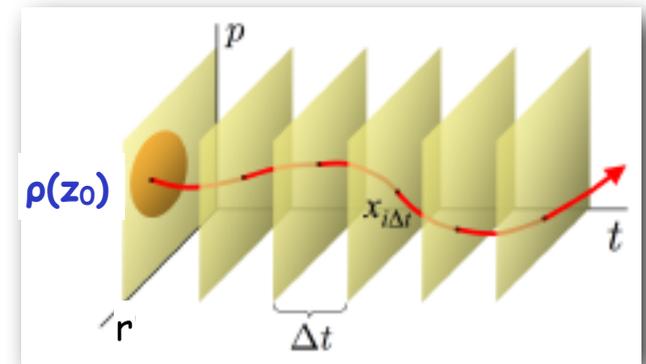
$$\mathbf{z}(\tau) \equiv \{z_0, z_{\Delta t}, \dots, z_{\tau}\}$$

- The probability distribution in the pathway ensemble depends on $\rho(z_0)$ and the dynamics

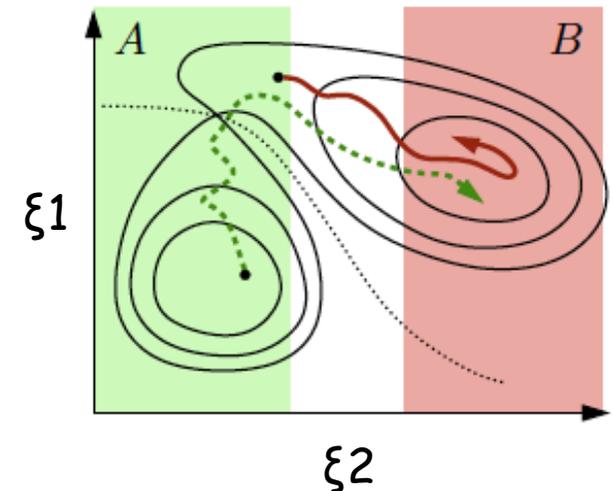
$$\mathcal{P}[z(\tau)] = \rho(z_0) \prod_{i=0}^{\tau/\Delta t - 1} p(z_{i\Delta t} \rightarrow z_{(i+1)\Delta t})$$

- Good description of initial and final states required by definition of multidimensional $\xi(\mathbf{r})$ to identify A and B

- large enough to contain equilibrium fluctuations
- no overlap with opposite basin of attraction



Markovian processes



Path generation

Transition-path-ensemble

- Focused on the sub-ensemble of pathways containing barrier-crossing events

$$\mathcal{P}_{AB}[z(\tau)] = h_A(z_0)\mathcal{P}[z(\tau)]h_B(z_\tau)/Z_{AB}(\tau)$$

- Random walk in TP-ensemble by Metropolis MC : **detailed balance criterion**

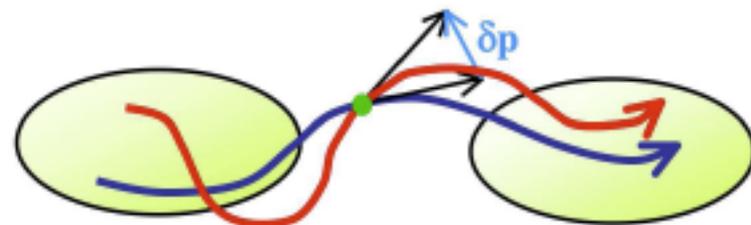
$$p_{\text{acc}}[z^n(\tau) \rightarrow z^o(\tau)] = \min \left\{ 1, \frac{\mathcal{P}_{AB}[z^n(\tau)]p_g[z^n \rightarrow z^o]}{\mathcal{P}_{AB}[z^o(\tau)]p_g[z^n \rightarrow z^o]} \right\}$$

$$p_{\text{acc}}[z^n(\tau) \rightarrow z^o(\tau)] = h_A(z_0^n)h_B(z_\tau^n) \min \left\{ 1, \frac{\rho(z_0^o)}{\rho(z_0^n)} \right\}$$

- The most probable reactive trajectories are identified and transition mechanisms are characterised

Efficiency:

- ☀ z^n and z^o as different as possible
- ☀ $p_{\text{acc}}[z^n \rightarrow z^o]$ as large as possible



Shooting



Shifting

Metadynamics

- ☀ Canonical equilibrium distribution under potential $V(\mathbf{r})$
- ☀ Choose a set of relevant **Collective Variables** $\mathbf{S}(\mathbf{r}):\{S_a(\mathbf{r})\}$, such that the process is well defined in the reduced space $\Sigma(\mathbf{S})$

$$P(\mathbf{S}) = \frac{e^{-\beta A(\mathbf{S})}}{\int d\mathbf{S} e^{-\beta A(\mathbf{S})}} \quad A(\mathbf{S}) = -\frac{1}{\beta} \ln \left(\int d\mathbf{r} e^{-\beta V(\mathbf{r})} \delta(\mathbf{S} - \mathbf{S}(\mathbf{r})) \right)$$

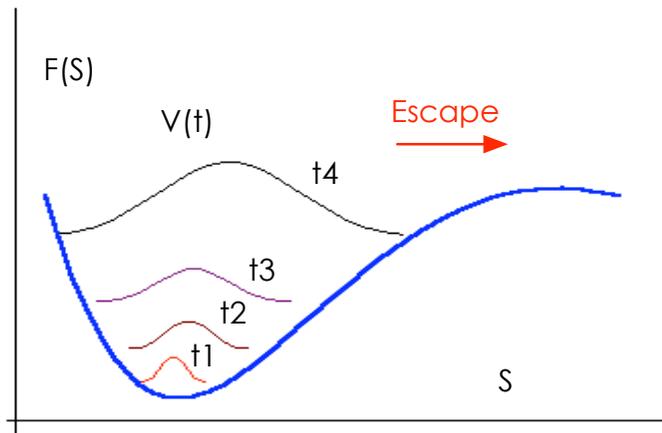
- ☀ Perform MD and re-map each micro-state by projecting the trajectory into the configuration space $\Sigma(\mathbf{S})$: **meta-trajectory** $\mathbf{S}(\mathbf{r}(t))$
- ☀ Enhance the exploration by adding a **penalty potential** that discourages the system to visit already explored macro-states

$$V_{\text{MTD}}(\mathbf{S}(\mathbf{r}), t) = \sum_{t'=\tau_G, 2\tau_G, \dots} W_{t'} e^{-\frac{[\mathbf{S}(\mathbf{r}) - \mathbf{S}(\mathbf{r}_G(t'))]^2}{2\Delta\mathbf{S}^2}}$$

- ☀ New probability distribution generated under the action of $V+V_{\text{MTD}}$

History Dependent Potential

Non Markovian Coarse-grained MD



Fill-up by successive contributions
 Keeping track of the explored space
 Direct estimate of FES topology

Eliminate metastability and reconstruct $A(\mathbf{S})$ within $\Sigma(\mathbf{S})$

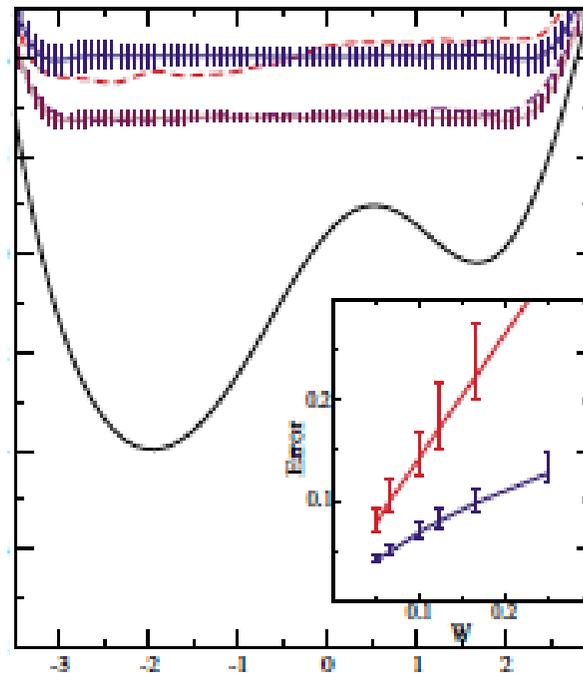
$$A_G(\mathbf{S}, t) = -V_{\text{MTD}}(\mathbf{S}(\mathbf{r}), t)$$

Flattening of free energy surface

$$W/\tau_G \rightarrow 0$$

$$P(\mathbf{S}) \propto e^{-\beta[A(\mathbf{S}) - A_G(\mathbf{S}, t)]}$$

$$\delta t_{\text{MD}} \ll \tau_G \ll \tau_s$$

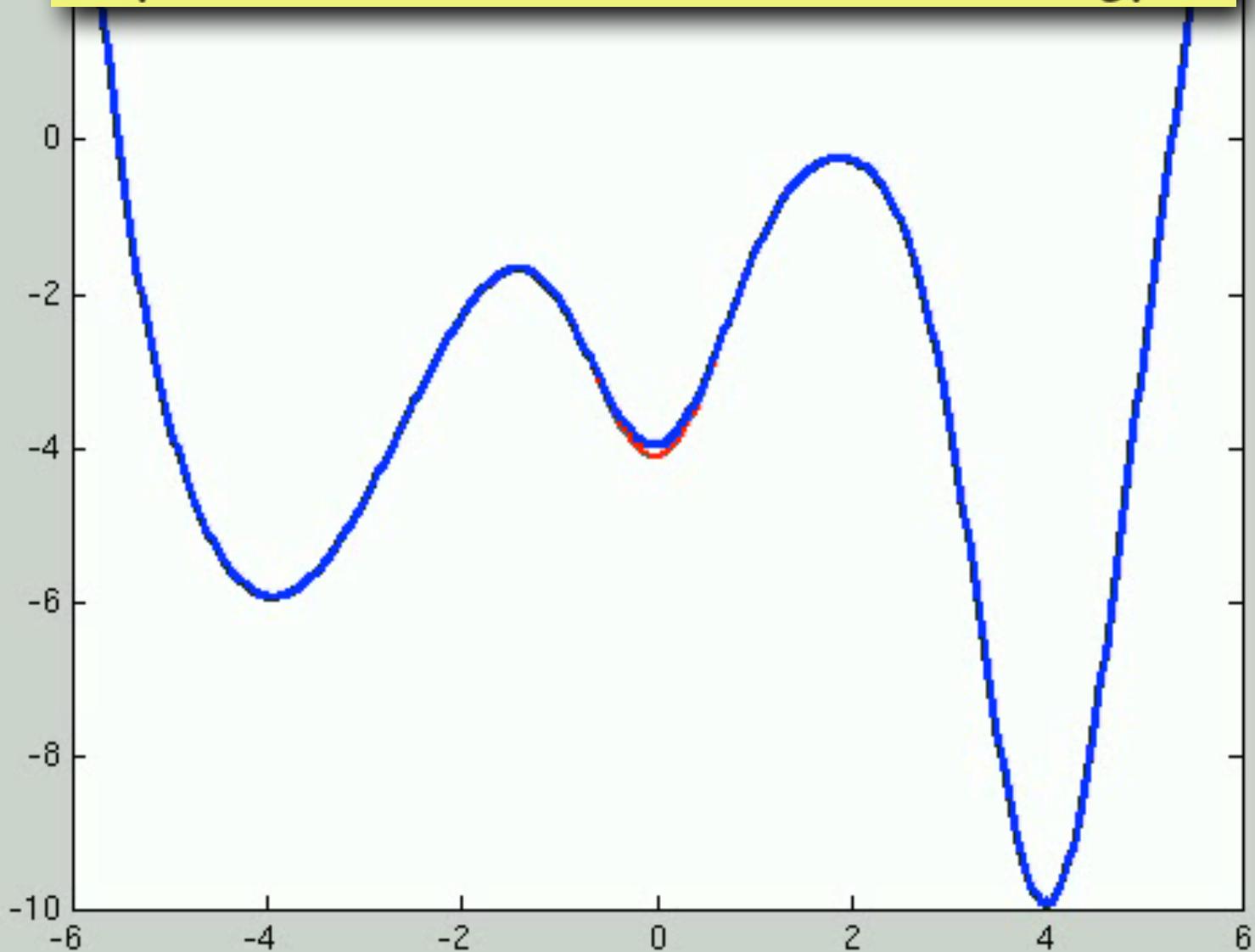


$$\delta A(\mathbf{S}) = A(\mathbf{S}) - A_G(\mathbf{S}, t)$$

$$\langle \delta A(\mathbf{S}) \rangle$$

2D FES

Histogram of a trajectory generated by $V+V_{\text{MTD}}$ provides a direct estimate of free energy



Extended Lagrangian MTD

Enforcing adiabatic separation, $\tau_s \gg$ other time scales and memory effects

Introduction of auxiliary variables $\mathbf{s} : \{s_\alpha\}$ one for each $S_\alpha(\mathbf{r})$
with large enough M

$$\mathcal{L} = K - V(\mathbf{r}) + \sum_{\alpha} \frac{1}{2} M_{\alpha} \dot{s}_{\alpha} - \sum_{\alpha} \frac{1}{2} k_{\alpha} (s_{\alpha} - S_{\alpha}(\mathbf{r}))^2 - V_G(\mathbf{s}, t)$$

thermalization coupling to system DoF sampling enhancement

$$V_G(\mathbf{s}, t) = \sum_{t' < t} W_{t'} e^{-\frac{(\mathbf{s} - \mathbf{s}_G(t'))^2}{2\Delta s^2}}$$

For large t and slow deposition rate, V_G approximates the free energy and the meta-trajectory $\mathbf{s}(t)$ follows the MEP

$$A_{\mathbf{k}}(\mathbf{s}) = -\frac{1}{\beta} \ln \left(\int d\mathbf{r} e^{-\beta [V(\mathbf{r}) + \frac{1}{2} \sum_{\alpha} k_{\alpha} (s_{\alpha} - S_{\alpha}(\mathbf{r}))^2]} \right)$$

$$\lim_{\mathbf{k} \rightarrow \infty} A_{\mathbf{k}}(\mathbf{s}) = A(\mathbf{s})$$

Equations of Motion

Smooth and continuous meta-trajectories that follow the minimum energy pathway are obtained by a modified Velocity Verlet algorithm

By the assumption that S_α is function only of \mathbf{R}_I

Force on the meta-variable

$$f_\alpha(t) = k_\alpha(S_\alpha(\{\mathbf{R}_I(t)\}) - s_\alpha(t)) - \frac{\partial}{\partial s_\alpha} V_{\text{MTD}}(\mathbf{s}, t)$$

V_{harm} and V_{MTD} generate opposite contributions

Modified force on ions due to the coupling with the dynamics of the meta-variables

$$f_I(t) = f_I^{(0)}(t) - k_\alpha(S_\alpha(\{\mathbf{R}_I(t)\}) - s_\alpha(t)) \frac{\partial S_\alpha}{\partial R_I}$$

The Metadynamics Lagrangian generates fictitious dynamics describing the most probable pathways

Gaussian anisotropic shape

Different CVs, anisotropic $A(s)$, different diffusion coefficients

$$\tau_{s_\alpha} = L_\alpha / D_\alpha$$

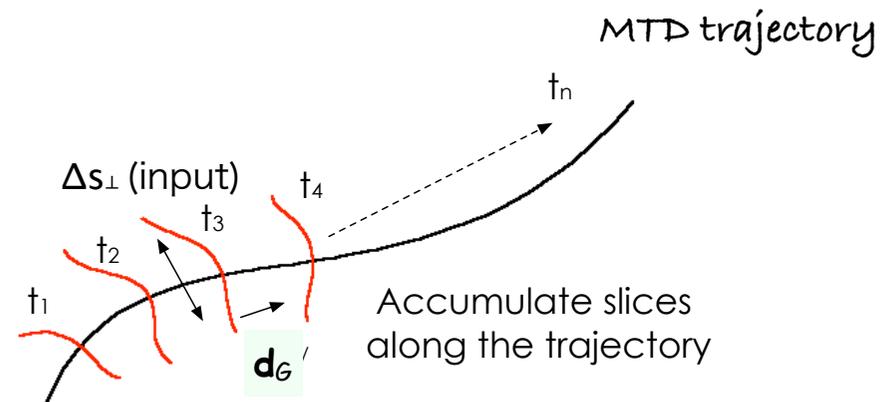
$$\sigma_\alpha = s_\alpha / L_\alpha$$

$$V_G(\mathbf{s}, t) = \sum_{t' < t} W_{t'} e^{-\sum_\alpha \frac{(s_\alpha - s_\alpha(t'))^2}{2(\Delta s_\alpha)^2}}$$

$$V_G(\mathbf{s}, t) = \sum_{t' < t} W_{t'} e^{-\frac{(\mathbf{s} - \mathbf{s}_G(t'))^2}{2\Delta s^2}}$$

$$\times e^{-\frac{[(\mathbf{s} - \mathbf{s}_G(t')) \cdot \mathbf{d}_G(t')]^2}{2\|\mathbf{d}_G(t')\|^4}}$$

re-shaped Gaussian tube



Accuracy of FES profile

Dynamics generating the equilibrium distribution associated with $A(\mathbf{s}) - A_G(\mathbf{s}, t)$

Averaging over many independent trajectories

$$\epsilon(\mathbf{s}, t) = \sqrt{\langle (A_G(\mathbf{s}, t) - A(\mathbf{s}) - \langle A_G(\mathbf{s}, t) - A(\mathbf{s}) \rangle)^2 \rangle}$$

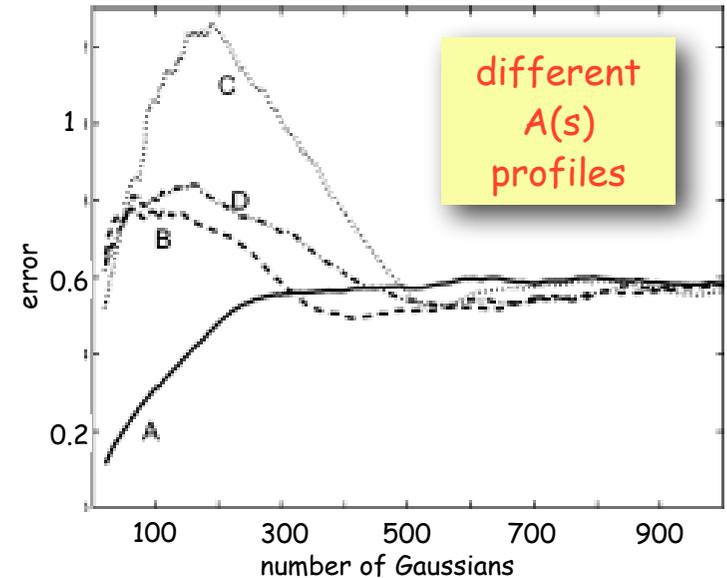
$$\bar{\epsilon}(t) = \frac{\int_{\Omega} ds \epsilon(\mathbf{s}, t)}{\int_{\Omega} ds}$$

Empirical error estimate

$$\bar{\epsilon} = C(d) \sqrt{\frac{\mathcal{V}_{\Omega} \|\Delta \mathbf{s}\| W}{\|\mathbf{D}\| \tau_G \beta}}$$

$$t_{\text{tot}} = \tau_G \frac{\int_{\Omega: A(\mathbf{s}) < A_{\text{max}}} ds (A_{\text{max}} - A(\mathbf{s}))}{(2\pi)^{d/2} W \prod_{\alpha} \Delta s_{\alpha}}$$

$$\bar{\epsilon} \propto \sqrt{\frac{\|\tau_{\mathbf{s}}\| \bar{A}}{t_{\text{tot}} \beta} \frac{\mathcal{V}_{\Omega}}{\prod_{\alpha} \Delta s_{\alpha}}}$$



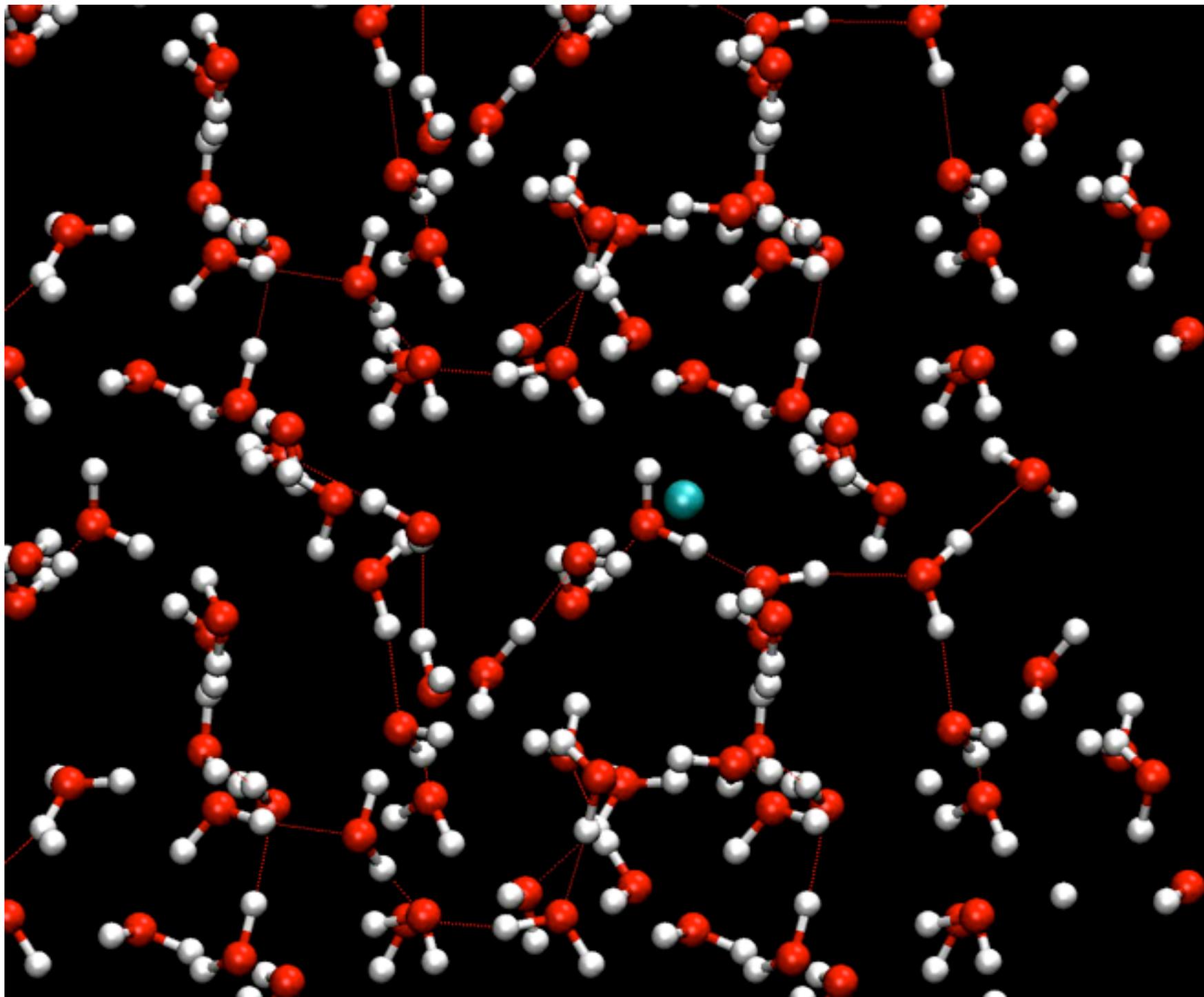
too large Δs would smear out $A(\mathbf{s})$ details : $\Delta s/L < 0.1$

Only relevant time scale is $\tau_{\mathbf{s}}$

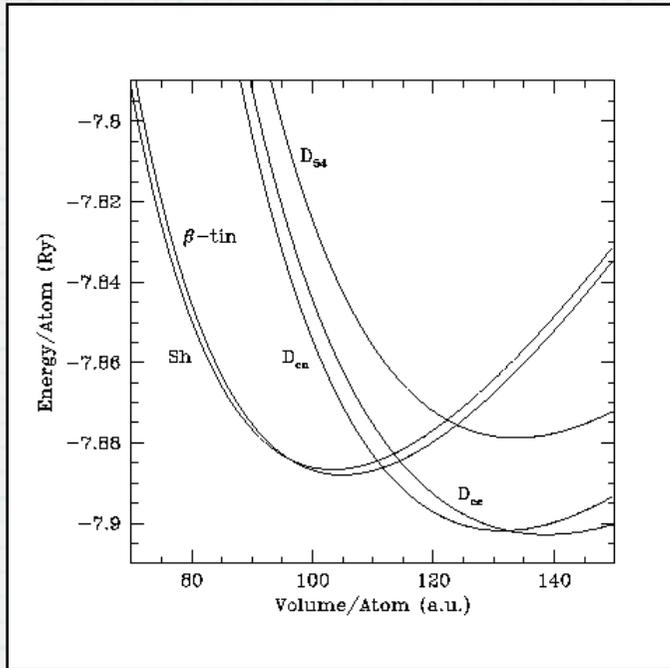
the error depends on τ_G/W
small Gaussians more frequently
is better

To be considered ...

- The selected CV must **discriminate among the relevant states** (reactants, products, TS)
- The **number of hills** required to fill the well is proportional to $1/(\Delta)^{N_{CV}}$
- The sampling of large variations of the CV over almost **flat energy regions** is expensive: **diffusive behavior**
- **MTD is not the true dynamics**. Reaction rates are derived a posteriori from the estimated FES
- The analysis of the trajectory is needed to isolate the TS
- With proper choices of CV and parameters, the MTD trajectory describes the **most probable pathway** taking into account also possible kinetic effects (larger and shallower channels are preferred)
- The accuracy in the evaluation of the FES depends on hills' shape and size, and on the deposition rate. The **ideal coverage** $V_G(\{S\}) = -A(\{S\})$ (**flat surface**)



Phase Transitions



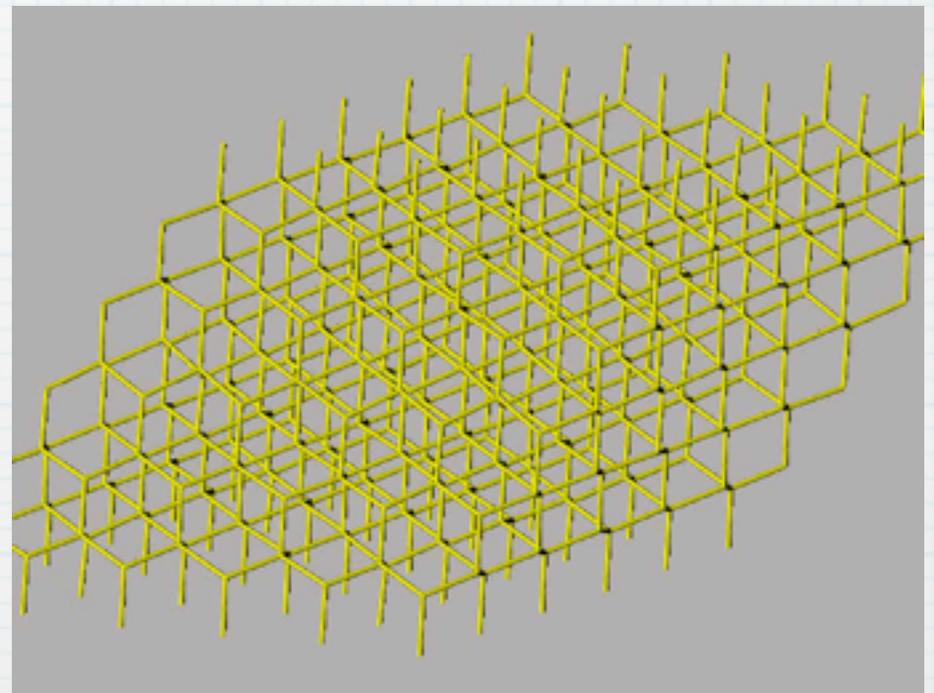
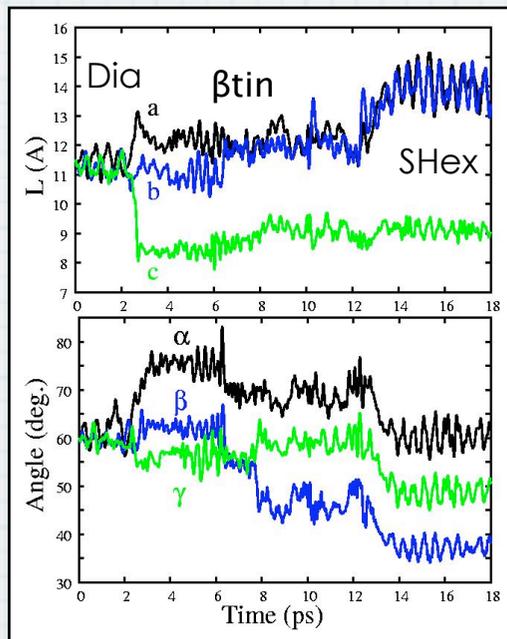
Crystalline Silicon can assume different lattice structures. Phase transitions induced by external pressure are known from experiment

Metadynamics is able to explore **all the accessible metastable states**, without requiring any over-pressurization

Collective variables: the 3 vectors that define the simulation box

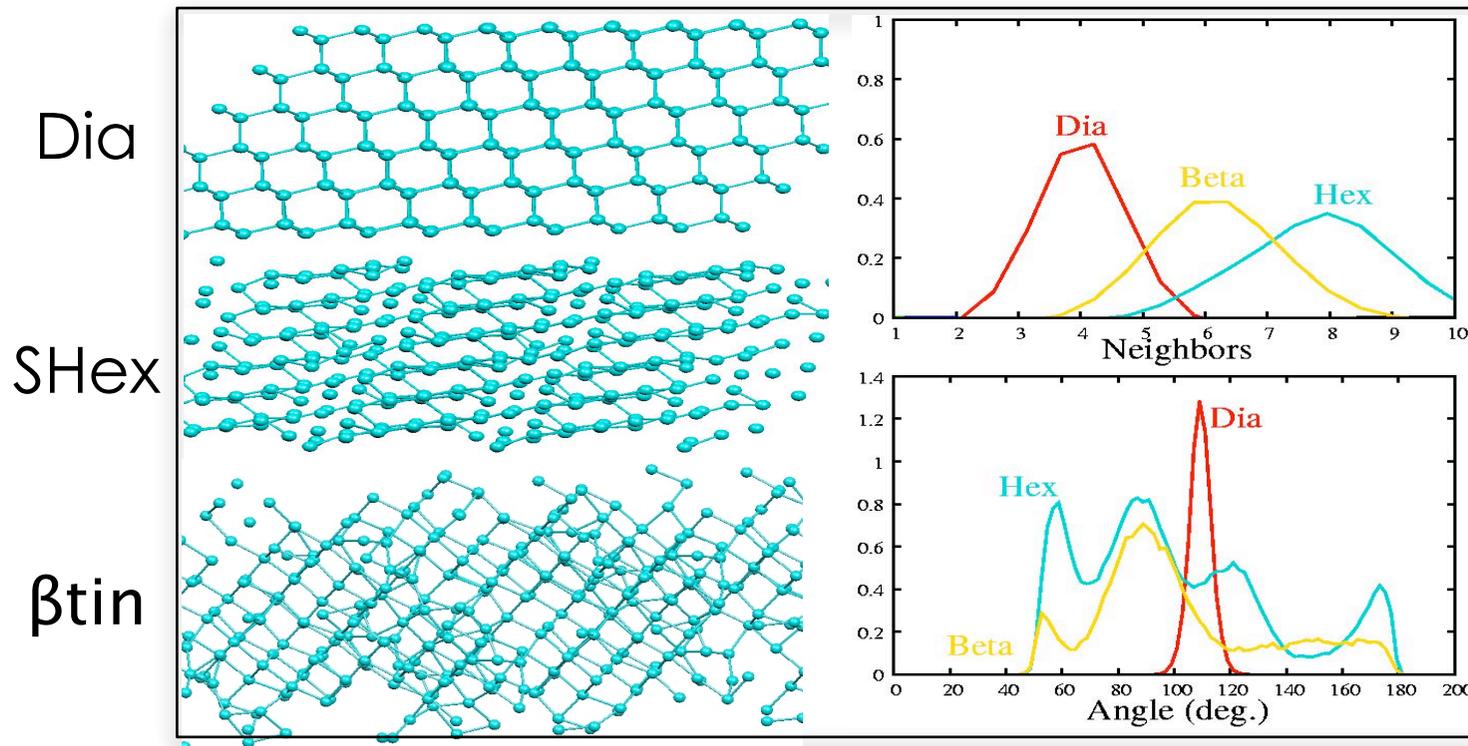
$$\mathbf{h} = [\mathbf{a}_1, \mathbf{a}_2, \mathbf{a}_3]$$

Evolution of the cell parameters during the metadynamics run



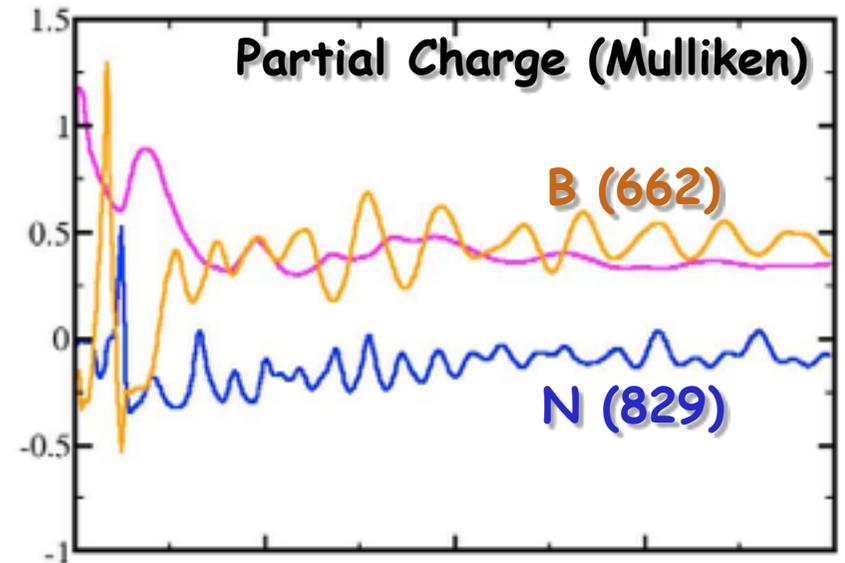
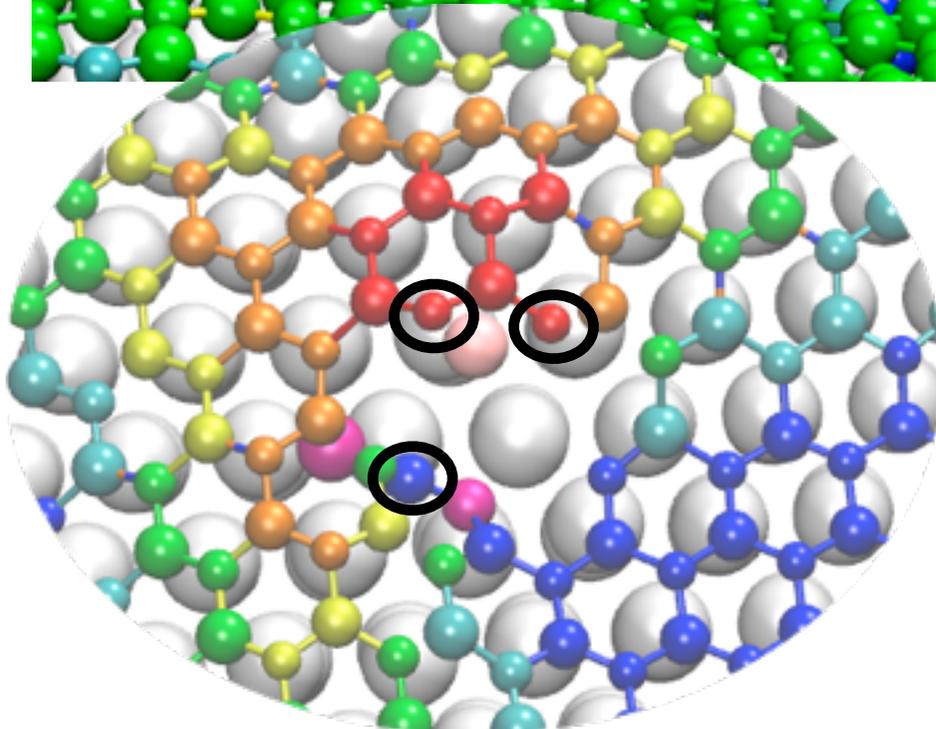
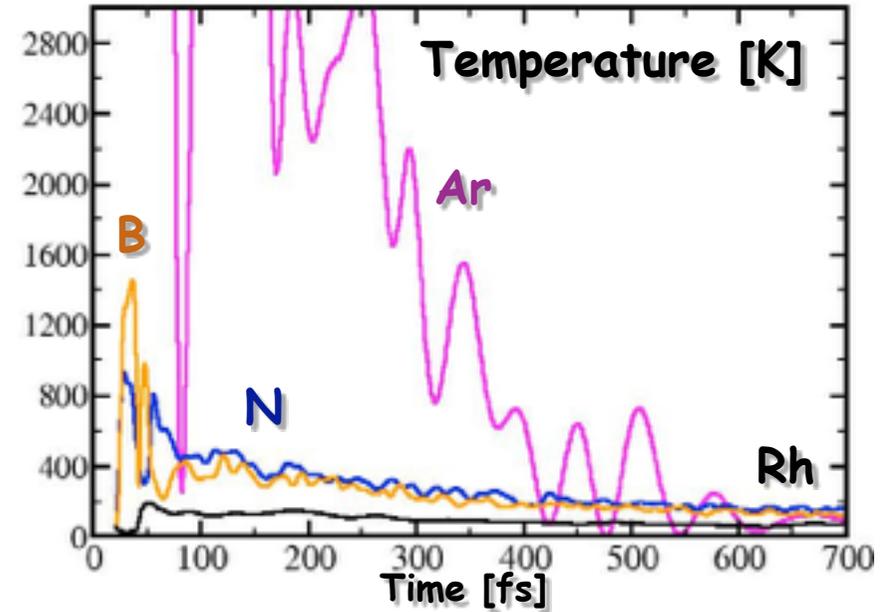
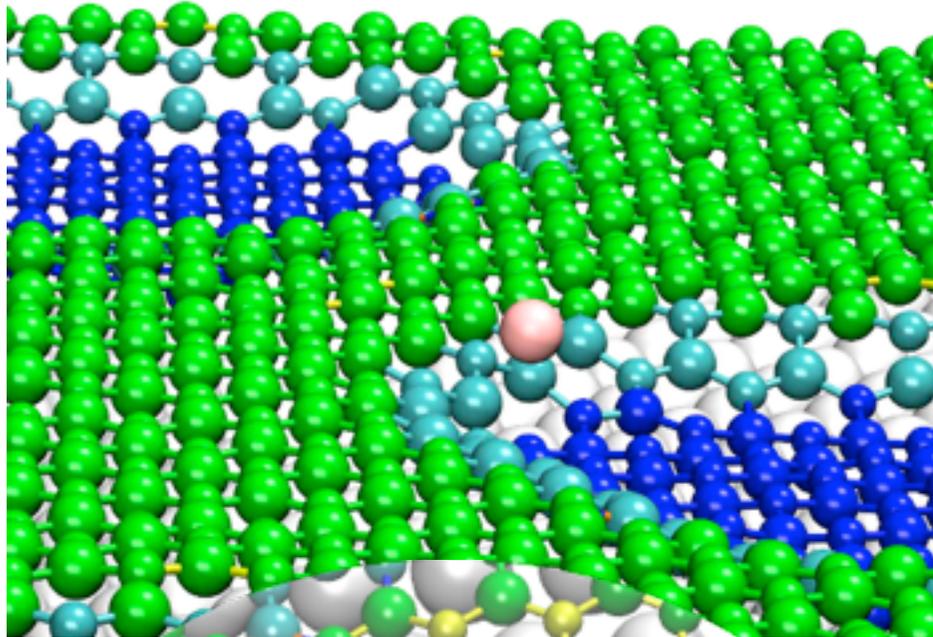
Silicon: Diamond, β -tin, Simple Hexagonal

Structural analysis by radial correlation function



Implantation Process

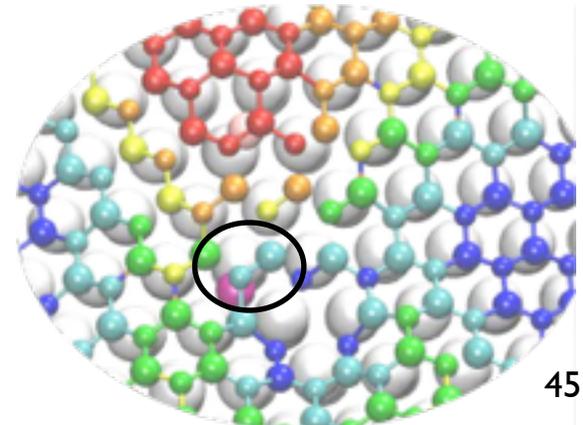
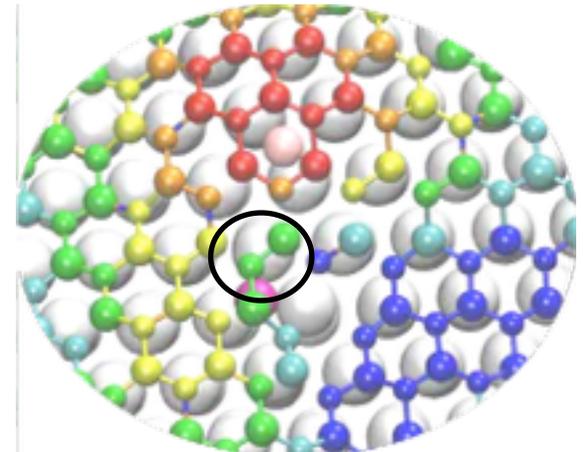
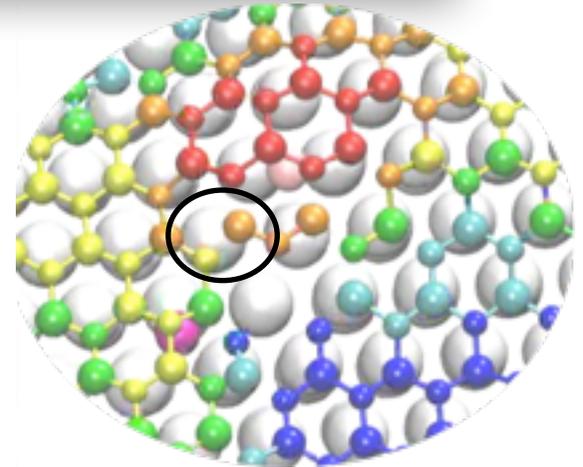
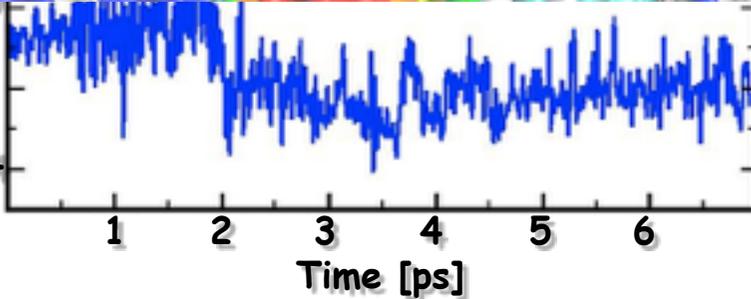
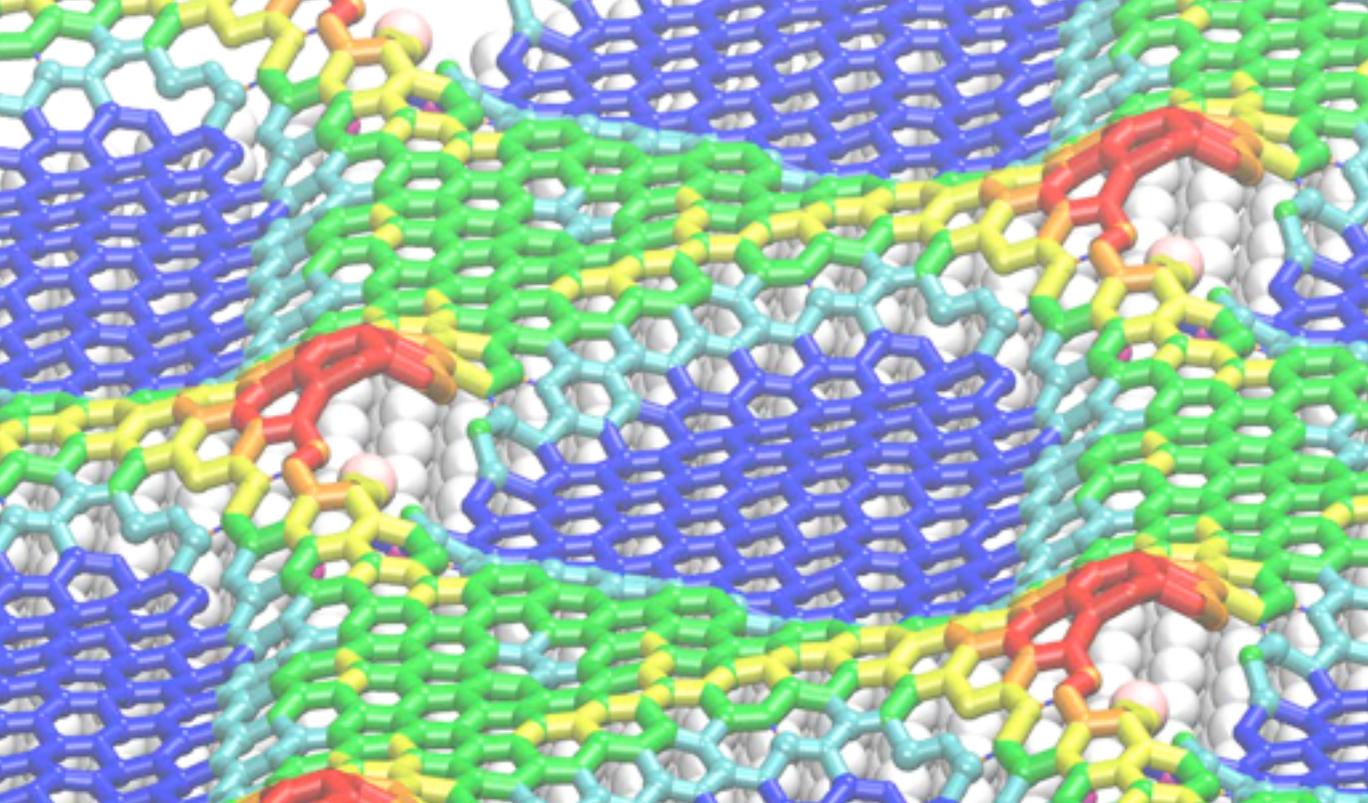
Ar+ at 50 eV striking the rim



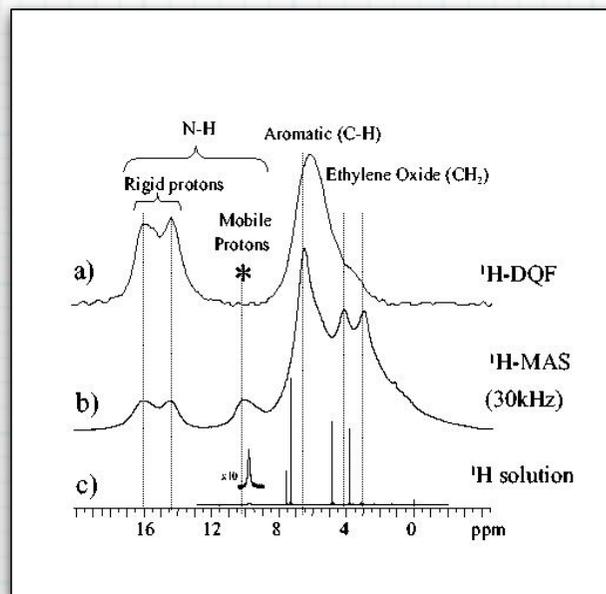
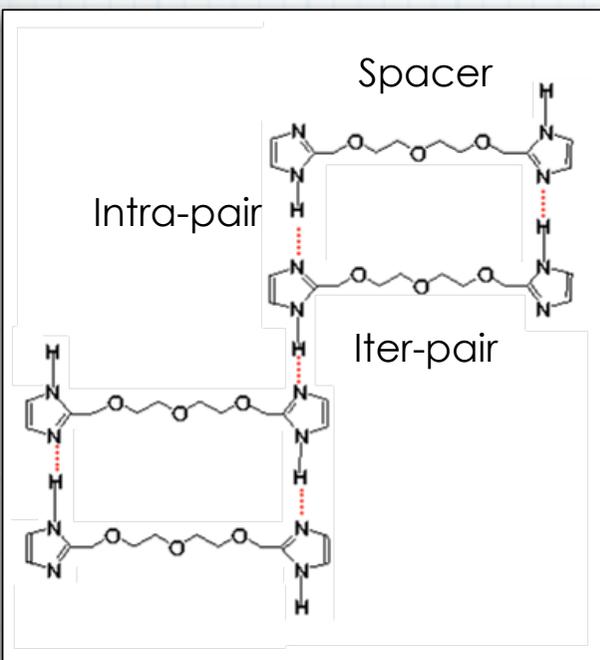
hBN Reconstruction

Metadynamics to accelerate the simulation of the healing process

Coordinate



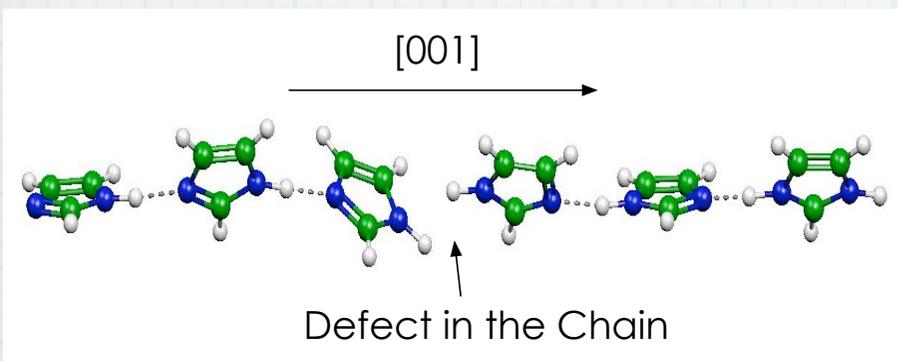
Imidazole-2EO: proton conductor



Good conductivity in doped samples

Dynamical disorder of local hydrogen-bond patterns

Fast microscopic rearrangements

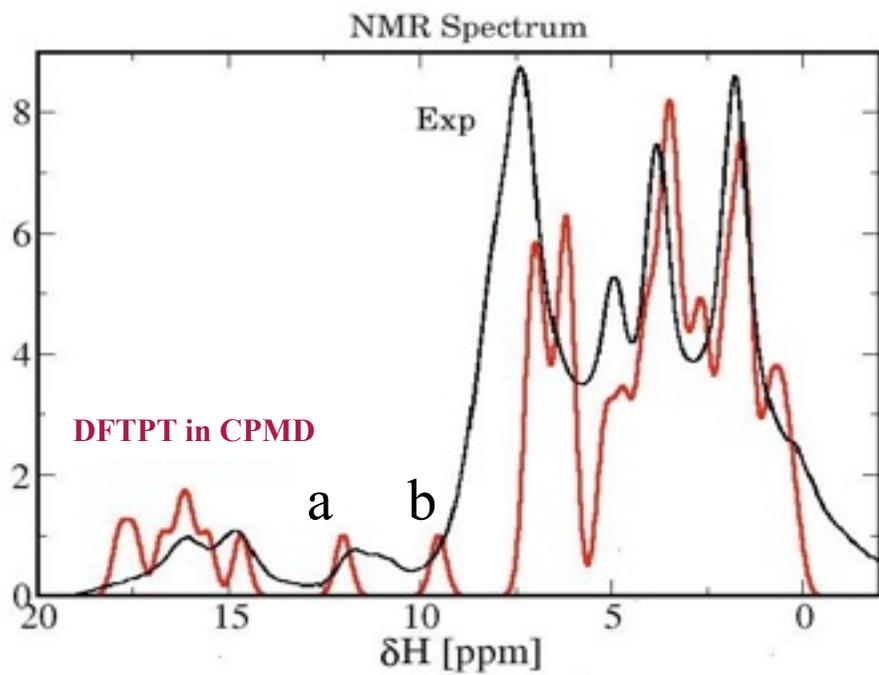


Generalized relative displacement

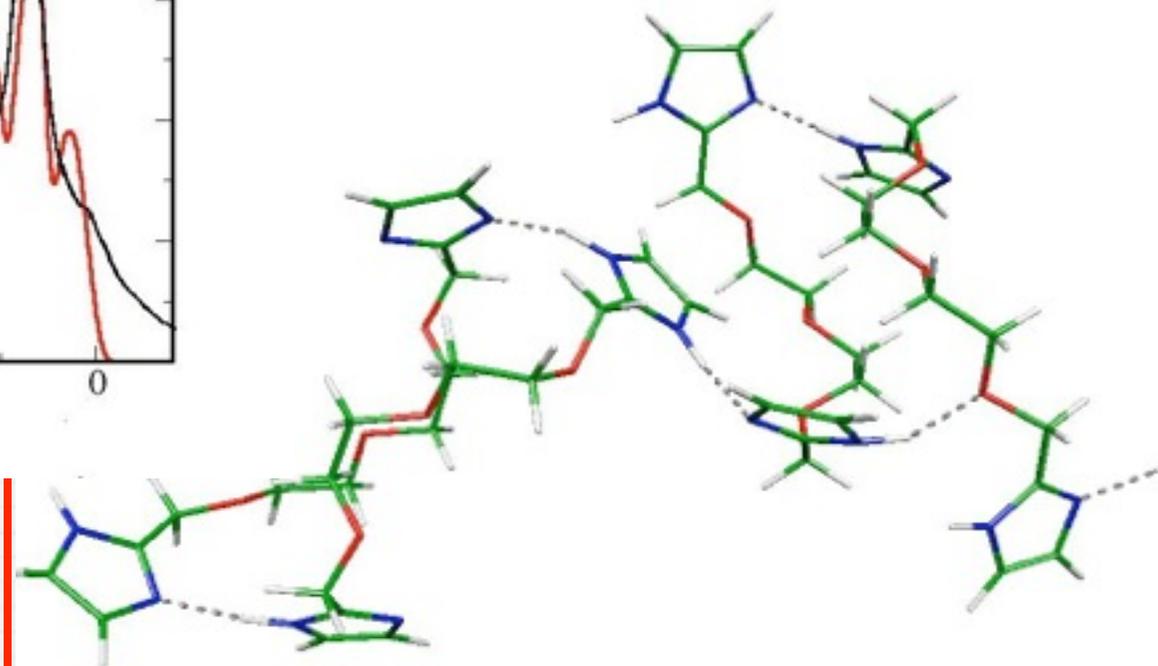
$$D_{H,N}^{[001]}(\{\mathbf{R}_I\}) = \frac{1}{N_H} \sum_{i \in H} \mathbf{d}_i \cdot \hat{\mathbf{v}}_{[001]} - \frac{1}{N_N} \sum_{j \in N} \mathbf{d}_j \cdot \hat{\mathbf{v}}_{[001]}$$

Coordination numbers and/or rotation angles

Excess-proton

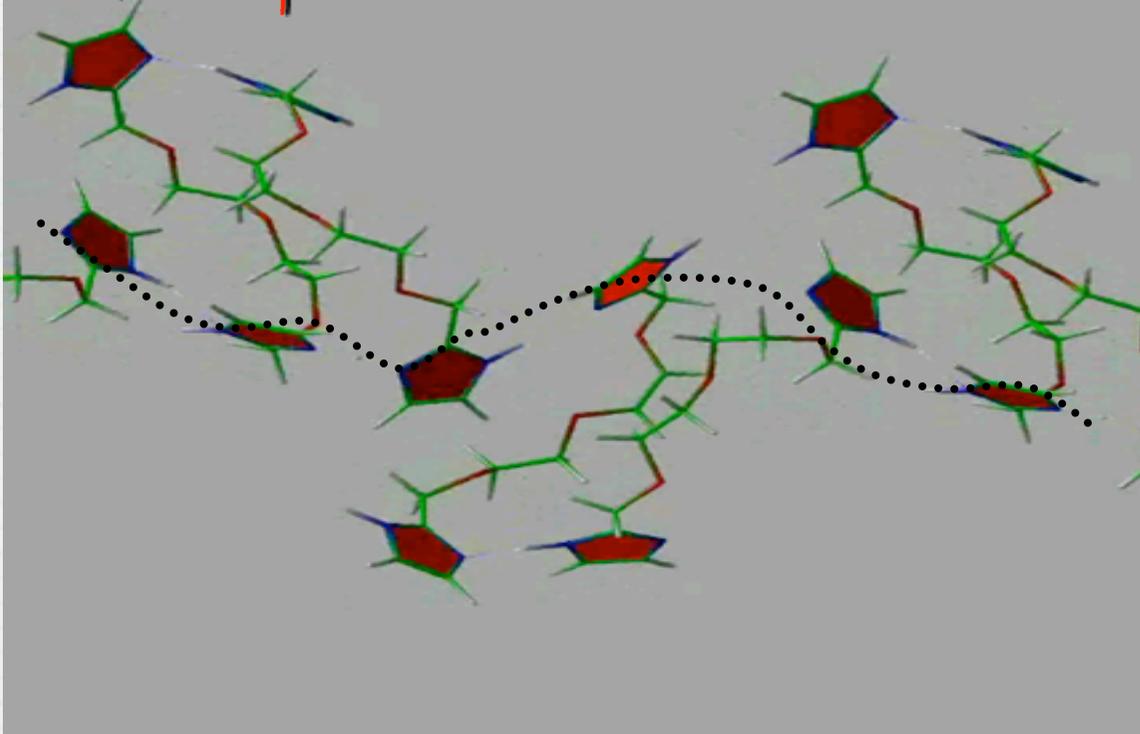


H-Doped Optimized Structure



Structure Diffusion

Excess proton: structural distortion

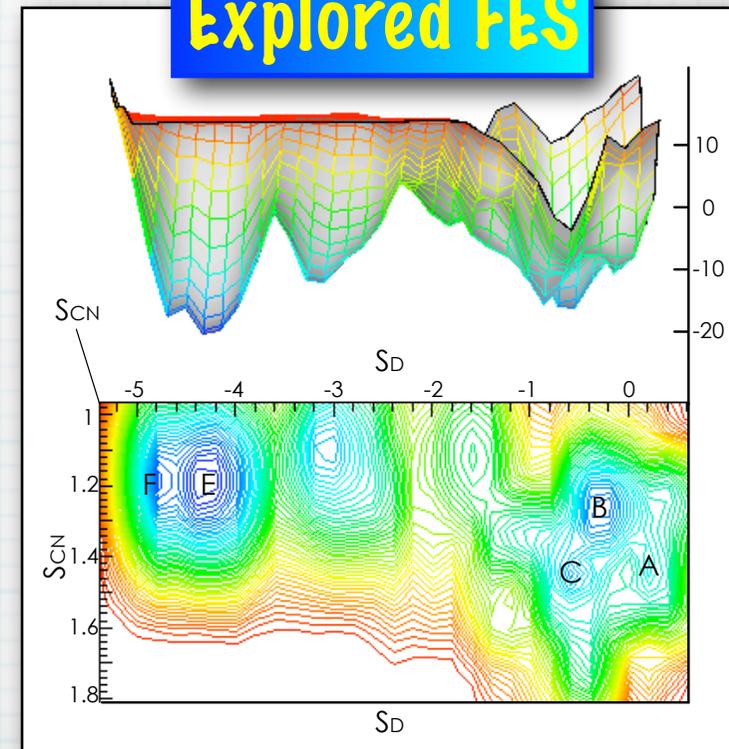


- Fluctuating hydrogen-bond network: structure diffusion
- Mobility induced by excess protons
- Monitoring defect diffusion by activating **5 Collective Variables**

- Facile intermolecular proton hopping
- Proton transferred over 4 molecules
- Complex pathway with several intermediates
- Rearrangement of the h-bond network
- Stabilizing role of O...H interactions

M Iannuzzi, Parrinello, PRL, **93**, 025901(2004);
M. Iannuzzi, JCP, **124**, 124710 (2006)

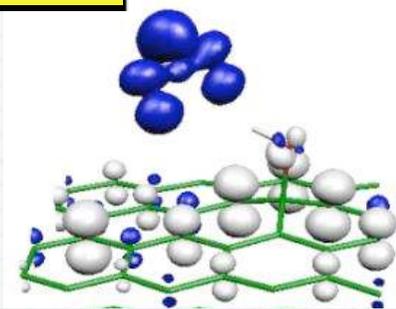
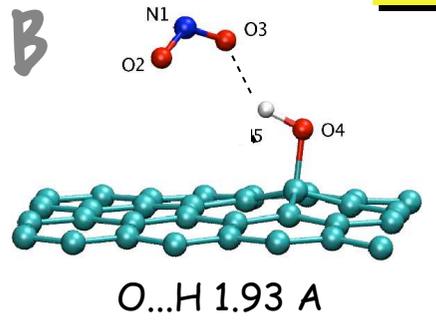
Explored FES



On Pristine Graphite

Three CN as CVs : N vs O, O vs C, N vs C

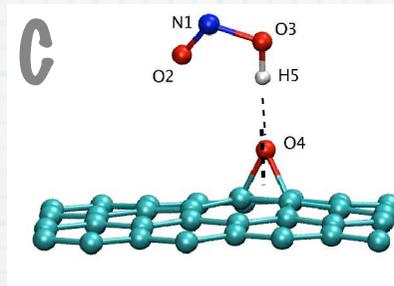
hydroxilation



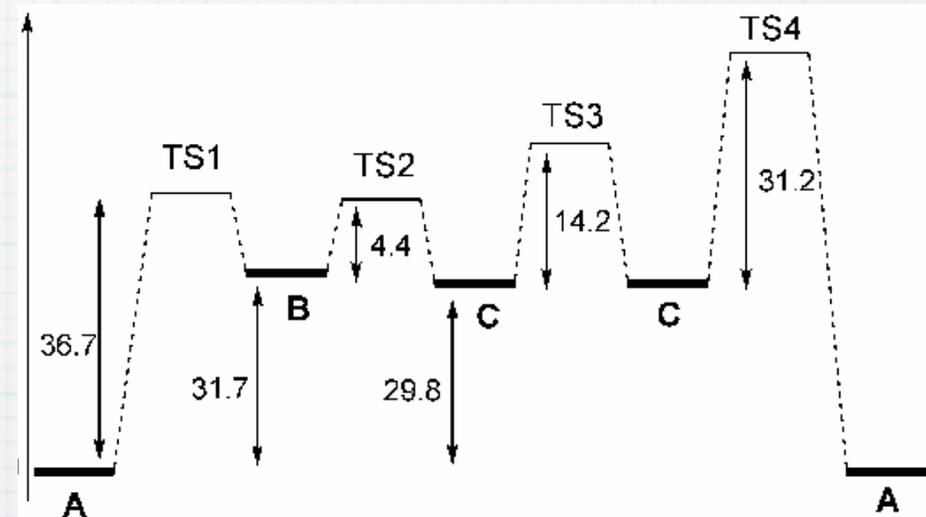
The first event is OH dissociation and hydroxilation of the surface (~35 kcal/mol), with formation of bi-radical system. System stabilized by the formation of H-bond and delocalization of the unpaired electron transferred to the surface. sp³ hybridization of C'

H transfer

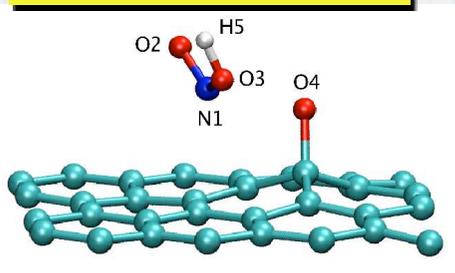
reactive NO₂ system is slightly stabilized; closed shell, only 1.9 kcal/mol more stable than open shell.
Epoxide structure migrates over the surface



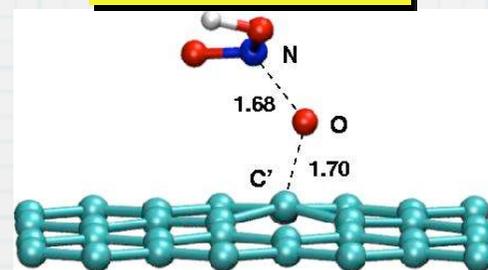
MEP energy profile



TS Epoxide migration



TS back transfer



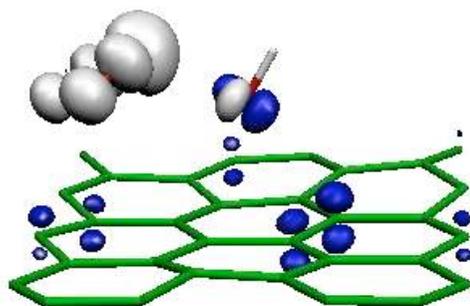
The direct epoxidation is a much less probable process

Graphite's role

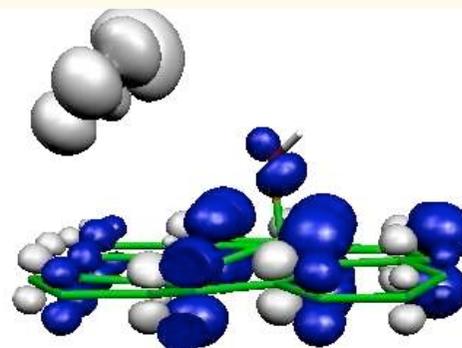
The OH-graphite interaction starts before the dissociation is complete

Early interaction between the incipient radical and the surface.
N-OH 2.1 Å
OH-graphite 2.5 Å
interaction with π el.

MD snapshot during the transition



MD snapshot at dissociation completed



As soon as the OH radical is adsorbed on the surface, its unpaired electron is delocalized among the atoms of the graphite layer.

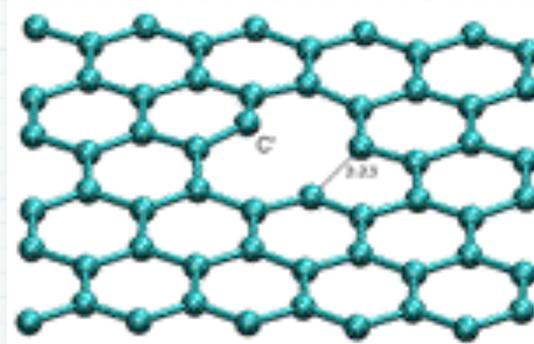
The dissociation barrier is 12 kcal/mol lower than in gas phase

Graphite catalyzes the dissociation of nitric acid
if the partial pressure is such as to guarantee a high probability of having HNO₃ in the vicinity of the surface.

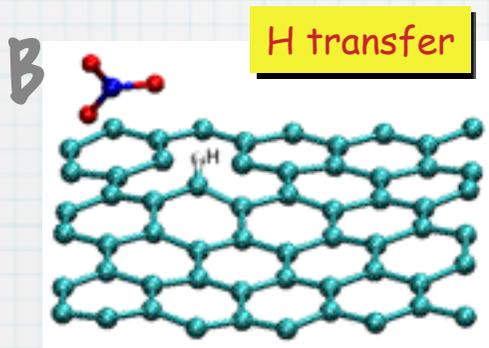
On Defective Graphite : vacancy

Simplest type of defects, predominant in carbon layers

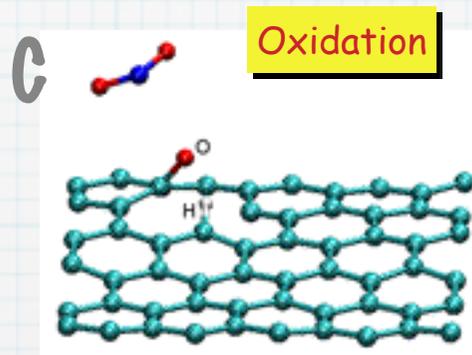
σ type sp^2 dangling bonds at the unsaturated C'



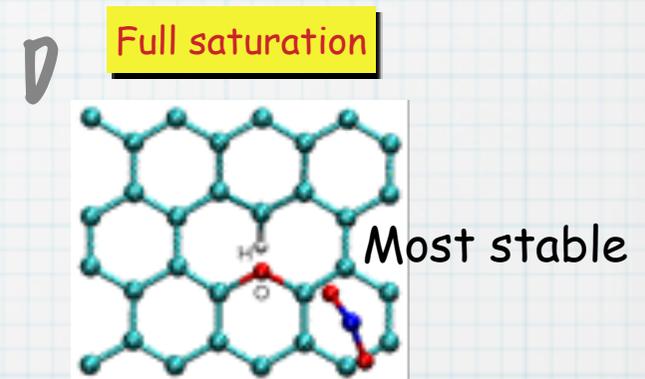
4 CN as CVs:
N \rightarrow O; O \rightarrow C;
N \rightarrow C; H \rightarrow C



unpaired H el. saturates σ type dangling,
 π spin density delocalized



C=O, unpaired π el. engaged
no π spin density

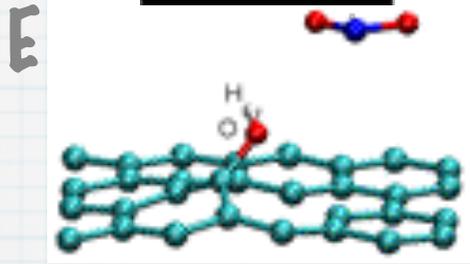


Ether group

Important effects of translational entropy on the free energy barriers

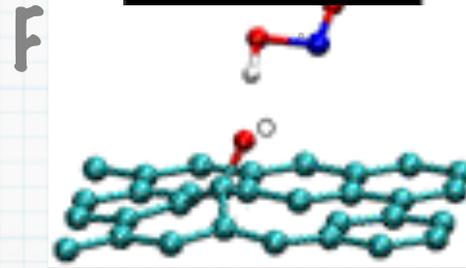
Alternative Paths

Hydroxilation



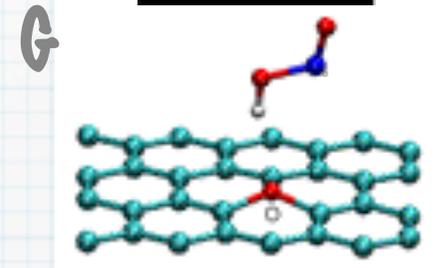
C-O σ type dangling,
 π spin density delocalized

Direct oxidation



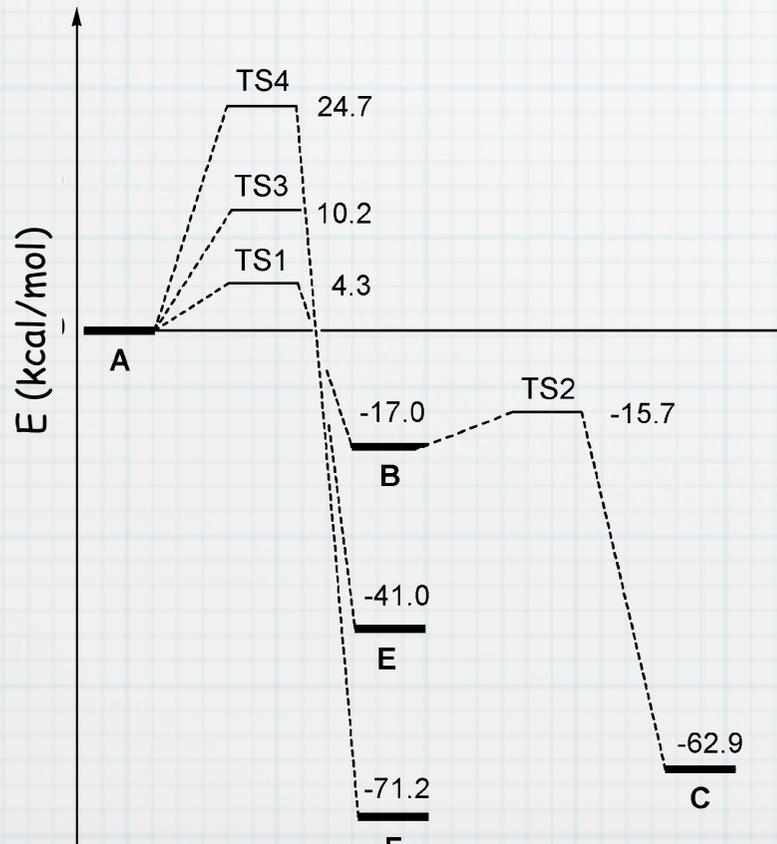
weak hydrogen bond
(1.88Å); closed shell

Replacement



sp3 character, not
symmetric, electropositive
O, no H-bond

the direct oxidation leads
to more stable
configurations but requires
higher energy barriers



The estimated activation energies
are much smaller than what
obtained in gas phase as well as on
pristine graphite

MEP: ABC