



Adaptive Resolution Molecular Dynamics

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School on *“Multiscale Modeling and Simulations of Hard and Soft
Materials”* JNCASR, Bangalore December 2009



Outline of Part I (a): The Method

- Concept of **adaptive**: Changing the number of degrees of freedom (DOF) on the fly
- Test system: Liquid of tetrahedral molecules
- The coupling formula on forces and the transition region
- Limitations regarding the control of the thermodynamic equilibrium
- Concept of switching as continuous change of phase space dimensionality and latent heat for equilibrium
- Why forces and not potentials
- Is the method working? Numerical tests



Outline of Part I (b): Applications

- Solvation of polymer in a liquid of tetrahedral molecules:
 - (i) The solvent properties, adaptive v.s. full atomistic
 - (ii) The polymer properties, adaptive v.s. full atomistic
- Liquid water:
 - (i) How to treat long range interactions
 - (ii) Properties of the liquid, adaptive v.s. full atomistic
 - (iii) Diffusion and position dependent thermostat



Collaborators

- **Matej Praprotnik**, Mainz
- **Kurt Kremer**, Mainz
- Simon Poblete, Mainz
- Adolfo Poma, Mainz
- Christoph Junghans, Mainz
- Cecilia Clementi, Rice University, Houston
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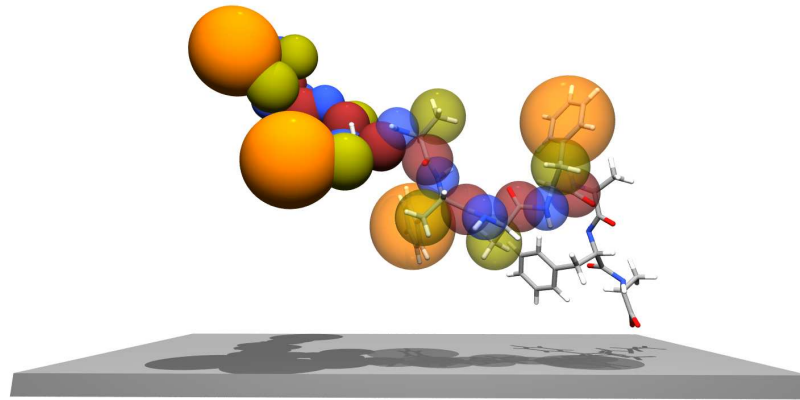
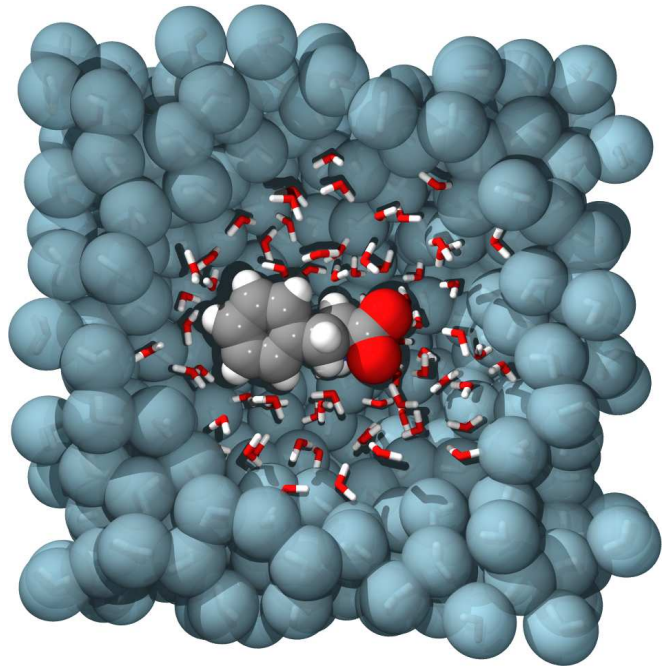


Motivations

- many problems in condensed matter are inherently **multiscale**
- **main aspect**: scales' interplay \implies simultaneous treatment of **all** the relevant scales
- **optimal computational approach**: treat only those degrees of freedom strictly required
- **different regions** may require **different resolution**



Examples





MD Simulation

All-Atom MD simulation:

- Atomistic (chemical) details
- (often) incapable to bridge the gap between a wide range of length and time scales

Coarse-Grained MD simulation:

- reduced number of degrees of freedom \implies longer length and time scales
- specific chemical details are lost

Solution:

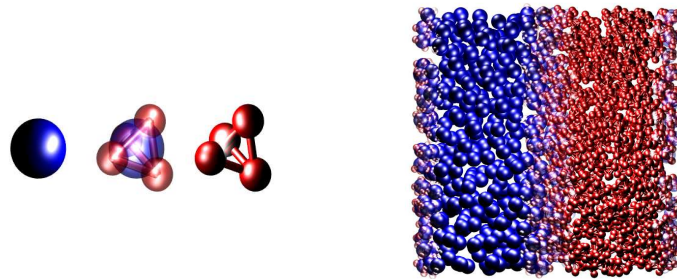
- Hybrid Adaptive MD Schemes



Method and model: General Idea

Adaptive Resolution MD Scheme

- on-the-fly interchange between atomistic and coarse-grained description **Hybrid Model**



with: $\rho_{ex} = \rho_{cg}$, $p_{ex} = p_{cg}$, $T_{ex} = T_{cg}$

M.Praprotnik, L.Delle Site and K.Kremer; Annu.Rev.Phys.Chem. 59, 545-571

(2008)

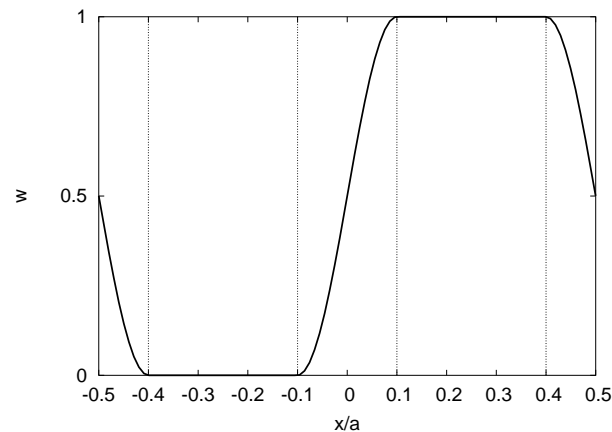


Scale Coupling

Two-stage procedure:

(a) Effective (coarse-grained) pair potential U^{cm} from the reference all-atom system.

$$(b) \mathbf{F}_{\alpha\beta} = w(X_{\alpha})w(X_{\beta})\mathbf{F}_{\alpha\beta}^{atom} + [1 - w(X_{\alpha})w(X_{\beta})]\mathbf{F}_{\alpha\beta}^{cm}$$

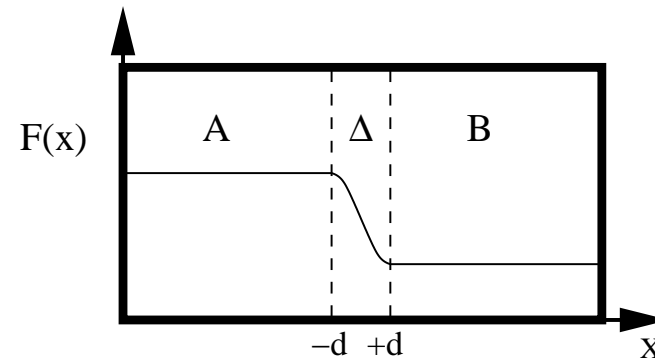


M. Praprotnik L. Delle Site and K. Kremer; J. Chem. Phys.; 123; 224106 (2005)

M. Praprotnik L. Delle Site and K. Kremer; Phys. Rev. E; 73, 066701 (2006)



How to define the switching function I



- What happens in Δ ?
- The number of DOF is $n = n(x)$ with ; $n_A = \text{const}_A$; $n_B = \text{const}_B$; and $n_\Delta = n(x)$
- The system is in equilibrium which implies:
$$\lim_{x \rightarrow d^-} \frac{\partial F_A(x)}{\partial x} = \lim_{x \rightarrow d^+} \frac{\partial F_B(x)}{\partial x} = 0$$
- If not true \rightarrow a free energy density gradient along x within the same level of resolution.

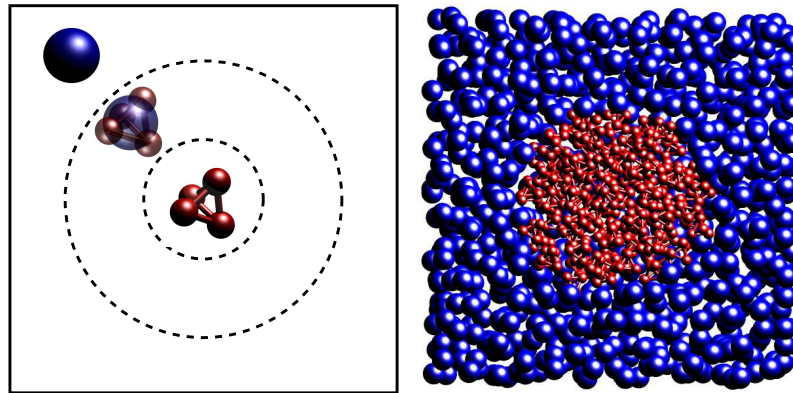


How to define the switching function II

- $\lim_{x \rightarrow d^-} \frac{\partial F_A(x)}{\partial x} = \lim_{x \rightarrow d^+} \frac{\partial F_B(x)}{\partial x} = 0$ can be shown to be equivalent to
 $\lim_{x \rightarrow d^-} \frac{\partial n_A(x)}{\partial x} = \lim_{x \rightarrow d^+} \frac{\partial n_B(x)}{\partial x} = 0$
- $w(x)$ is such that $w(x) = 1; \forall x \in A$ and $w(x) = 0; \forall x \in B$, with
 $\lim_{x \rightarrow d^+} \frac{\partial w(x)}{\partial x} = \lim_{x \rightarrow d^-} \frac{\partial w(x)}{\partial x} = 0$.
- In accordance with the equation above, we require that the weighting function $w(x)$ is continuous up to the first derivative and that goes monotonically from the value one to zero in the region Δ .



Spherical Geometry



Spherical Symmetry:

$$\mathbf{F}_{\alpha\beta} = w(R_\alpha)w(R_\beta)\mathbf{F}_{\alpha\beta}^{atom} + [1 - w(R_\alpha)w(R_\beta)]\mathbf{F}_{\alpha\beta}^{cm}$$

C.Junghans, M.Praprotnik and L.Delle Site, in **Multiscale Simulation Methods in Molecular Sciences**, NIC Series Volume 42 (2009)



Change of Resolution

- Molecule from **Atom** → **cg**: vanishing **contributions** of switching DOFs
- $w(x)$ smoothly "*freezes*" the **dynamical evolution** of such DOFs and their **contribution to the interactions**
- (vice versa) $w(x)$ smoothly "*reactivates*" their dynamics and their contributions to the interactions

by construction the **third Newton's Law** is preserved (**crucial in MD**).

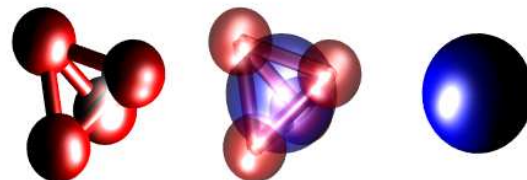


Technical aspects of the test molecule

A tetrahedral molecule has a defined spatial orientation and $3N = 12$

DOFs:

- 3 translational
- 3 rotational
- $3N - 6 = 6$ vibrational





Molecular Interactions in the Test System

Shifted 12-6 Lennard-Jones potential:

$$U_{LJ}^{atom}(r_{i\alpha j\beta}) = \begin{cases} 4\epsilon \left[\left(\frac{\sigma}{r_{i\alpha j\beta}} \right)^{12} - \left(\frac{\sigma}{r_{i\alpha j\beta}} \right)^6 + \frac{1}{4} \right]; & r_{i\alpha j\beta} \leq 2^{1/6}\sigma \\ 0; & r_{i\alpha j\beta} > 2^{1/6}\sigma \end{cases}$$

FENE potential:

$$U_{FENE}^{atom}(r_{i\alpha j\alpha}) = \begin{cases} -\frac{1}{2}kR_0^2 \ln \left[1 - \left(\frac{r_{i\alpha j\alpha}}{R_0} \right)^2 \right]; & r_{i\alpha j\alpha} \leq R_0 \\ \infty; & r_{i\alpha j\alpha} > R_0 \end{cases}$$



Effective pair potential

- In the limit of $\rho \rightarrow 0$:

$$U^{eff} = F(r) = -K_B T \ln g(r) \quad (1)$$

- For $\rho > 0$ we use the above relation as the initial approximation of an **iteration scheme**:

$$U_{i+1}^{eff}(r) = U_i^{eff}(r) + k_B T \ln \frac{g_i(r)}{g_{target}(r)} \quad (2)$$

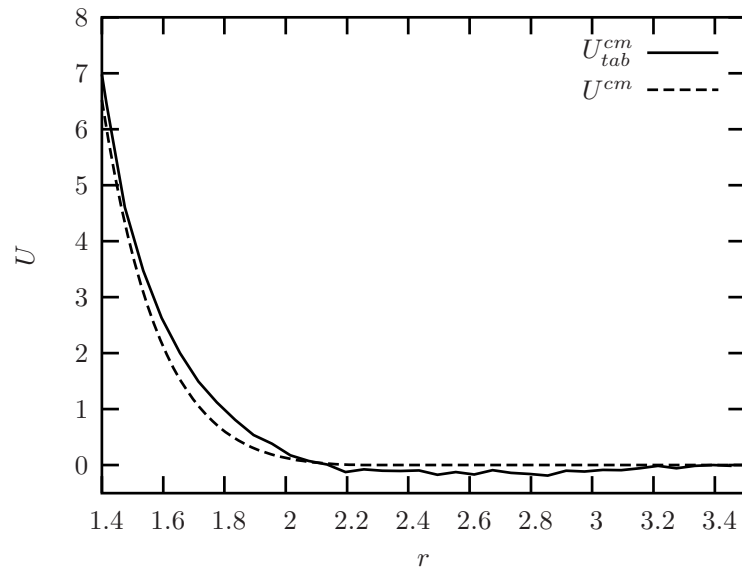
Pressure correction:

$$\Delta V_{lin} = A \left(1 - \frac{r}{r_{cut}}\right). \quad (3)$$

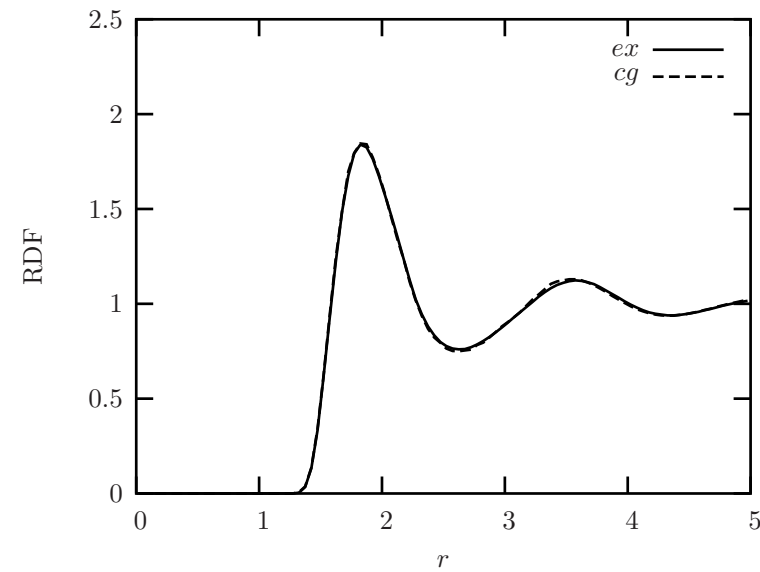
D. Reith, M. Pütz, F. Müller-Plathe, J. Comput. Chem., **24**, 1624 (2003).



effective pair potentials



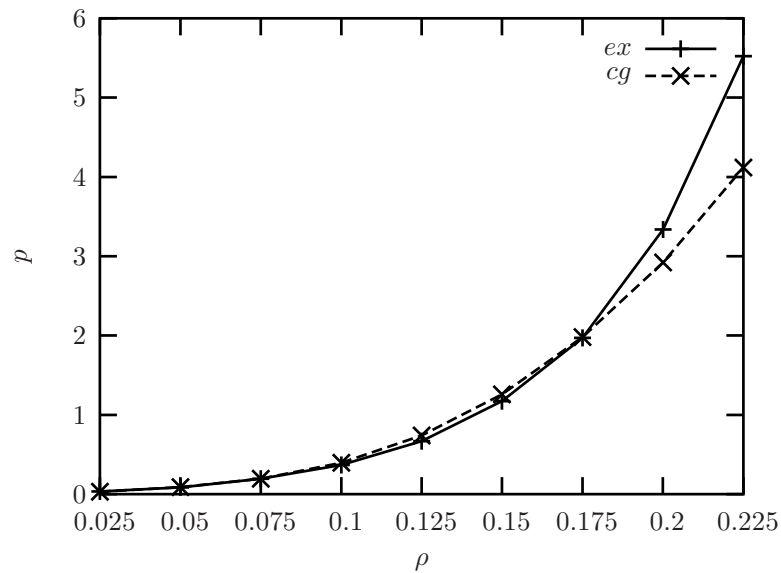
(g) Tabulated potential.



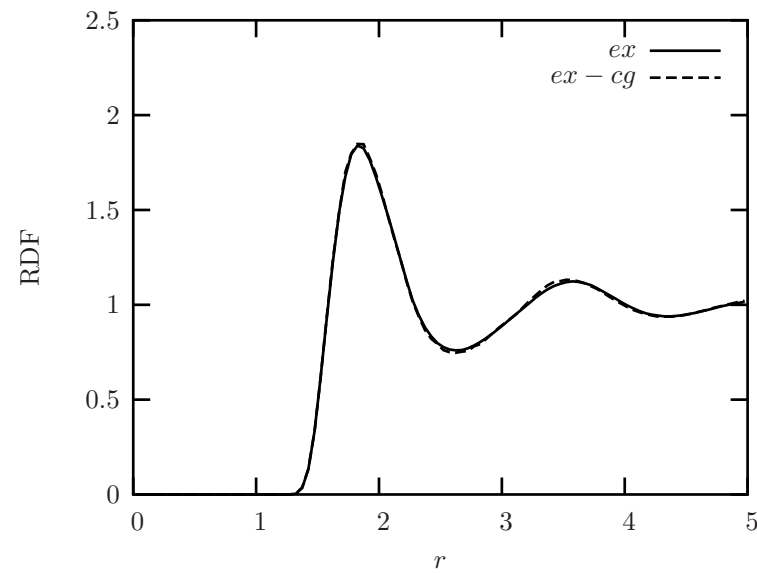
(h) RDF_{cm} .



Equation of state



(i) Equation of state for $\rho = 0.175$ and $T = 1$.



(j) RDF_{cm} .

M.Praprotnik, L.Delle Site and K.Kremer, Phys.Rev.E 73, 066701 (2006).



Control of the thermodynamical equilibrium (I)

- Adaptive force **alone** cannot assure **thermodynamic equilibrium**
- it is not conservative \implies **potential energy** cannot be written explicitly
- **BUT:** statistical mechanics principles of **varying resolution** \implies tools for thermodynamical equilibrium



Control of the thermodynamical equilibrium (II)

- Free energy is an extensive quantity and thus proportional to the number of DOF
- **preferential tendency** of one species to migrate into the other region
- **or equivalently** $\mu_{atom} \neq \mu_{cg}$
- **unphysical**: artifact of the different representations.



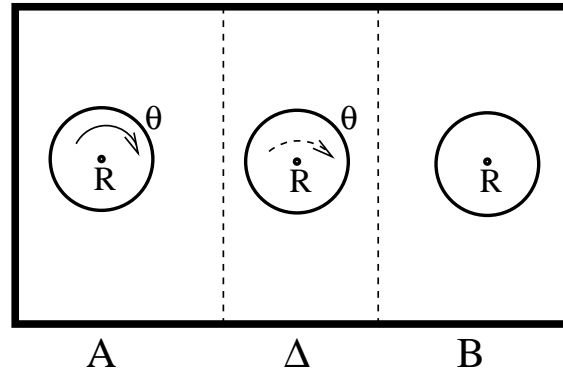
Change of resolution and corresponding latent heat

- The change of resolution: change of the number of *active/thermalized* DOF \implies latent heat ϕ
- $\mu_{atom} = \mu_{cg} + \phi$
- Explicit form of ϕ unknown
- **numerical:** couple the system to a position dependent thermostat
- How to define the temperature in Δ



Temperature in the region Δ

- Equipartition Theorem: $T_A = \frac{2\langle K_A \rangle}{n_A}$; $T_B = \frac{2\langle K_B \rangle}{n_B}$
- in Δ : $T_\Delta = \frac{2\langle K_\Delta \rangle}{n_\Delta}$?; What is $\langle K_\Delta \rangle$?
- 2-dimensional example:



- $K_A = \frac{1}{2}[\dot{\mathbf{R}}^2 + p_\Theta^2] = \frac{1}{2}[\dot{R}_x^2 + \dot{R}_y^2 + \dot{\Theta}^2]$; $K_B = \frac{1}{2}\dot{\mathbf{R}}^2 = \frac{1}{2}[\dot{R}_x^2 + \dot{R}_y^2]$
- R_x , R_y , and Θ , contribute each with $T/2$ to the kinetic energy
- Hence, $T_A = \frac{2\langle K_A \rangle}{3}$, $T_B = \frac{2\langle K_B \rangle}{2}$



Adaptive resolution via a continuous change of the phase space dimensionality

- **Statistical calculation of physical quantities:**
 - (a) a DOF **fully counts** in the atomistic region ($\int \dots d\theta$)
 - (b) it **does not count at all** in the coarse-grained region (no integration over θ)
- In Δ ($0 < w(x) < 1$): switching **equivalent to** continuously change **the dimensionality of the phase space**
- **mathematical formalism:** fractional calculus \implies
$$dV_w = d^w q \Gamma(w/2) / 2\pi^{w/2} \Gamma(w) = |q|^{w-1} dq / \Gamma(w) = dq^w / w \Gamma(w)$$
(for a fixed value of w , and generic DOF q)



Temperature in the region Δ : The extension of the Equipartition Theorem to non integer DOF

- **partition function** for the fractional quadratic DOF Θ :

$$\exp(-\beta F_w) = C \int \exp(-\beta p_\Theta^2/2) dV_w =$$
$$2C \int_0^\infty \exp(-\beta p_\Theta^2/2) |p_\Theta|^{w-1} \frac{dp_\Theta}{\Gamma(w)} = \frac{2^{w/2} C \Gamma(w/2)}{\Gamma(w)} \beta^{-w/2} \sim \beta^{-w/2}$$

- **fractional analog of the Equipartition Theorem:**

$$\langle K_w \rangle = \frac{d(\beta F_w)}{d\beta} = \frac{w}{2\beta} = \frac{wT}{2}$$

$\langle K_w \rangle$: average kinetic energy per fractional quadratic DOF with the parametric weight w

- In equilibrium $T_A = T_B = T_\Delta = T$ and thus: $n_\Delta = 2 + w$ (proof of consistency)



Fractional DOF: A more general view

- A generic statistical average of a quantity A in the canonical ensemble:

$$\langle A \rangle = \frac{\int_0^\infty e^{-\beta H(q)} A(q) dq}{\int_0^\infty e^{-\beta H(q)} dq}. \quad (4)$$

- with a fractional degrees α :

$$\langle A \rangle_\alpha = \frac{\int_0^\infty e^{-\beta H(q)} q^{\alpha-1} A(q) dq}{\int_0^\infty e^{-\beta H(q)} q^{\alpha-1} dq}. \quad (5)$$

- If $H(q) = |q|^m; m > 0$ and $A(q) = C_n |q|^n; n > 0$ then:

$$\langle A \rangle_\alpha = \frac{C_n \int_0^\infty e^{-\beta |q|^m} q^{\alpha+n-1} dq}{\int_0^\infty e^{-\beta |q|^m} q^{\alpha-1} dq}. \quad (6)$$

$$\langle A \rangle_\alpha = \frac{\Gamma(\frac{\alpha+n}{m})}{\Gamma(\frac{\alpha}{m})} \beta^{-n/m}. \quad (7)$$

- if $m = n \rightarrow \langle H(q) \rangle_\alpha$:

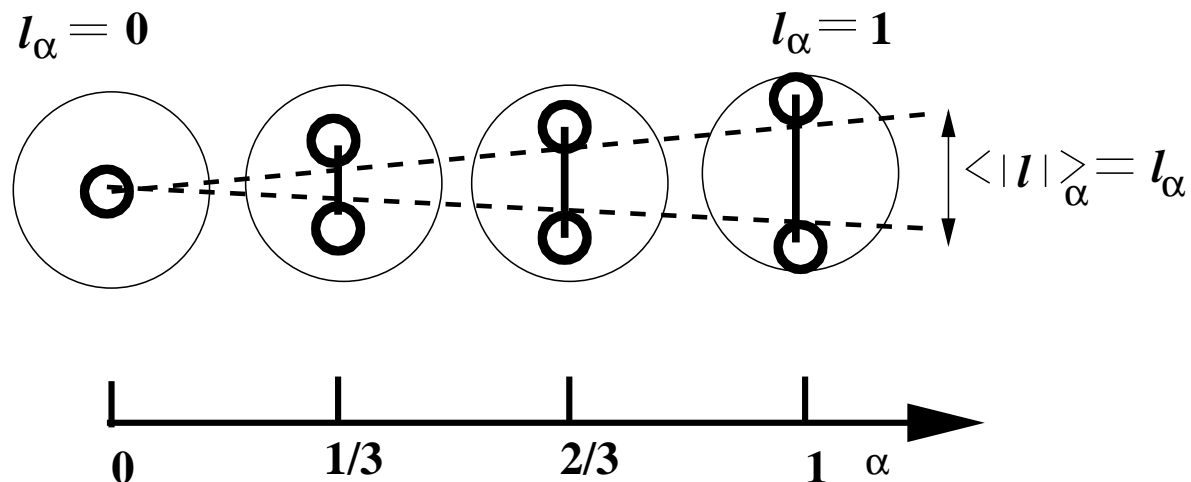
$$\langle H(q) \rangle_\alpha = \frac{\alpha}{n} \beta^{-1} \text{ that is } \langle H(q) \rangle_\alpha = \alpha \langle H(q) \rangle$$



The average molecular bond: An example of adaptive representation

Consider the case of the average length of a diatomic molecule

- $\langle |l| \rangle_\alpha = \frac{\int_0^\infty l^\alpha e^{-\beta l^2} dl}{\int_0^\infty e^{-\beta l^2} dl}$
- $\langle |l| \rangle_\alpha = \frac{\Gamma(\frac{1}{2})\Gamma(\frac{\alpha+1}{2})}{\Gamma(\frac{\alpha}{2})} \langle |l| \rangle$





Theoretical Basis: Conclusions

- **(1)** Changing Resolution \rightarrow Latent Heat \rightarrow position dependent
Thermostat
- **(2)** The temperature in the switching region, T_{Δ} , can be obtained
by **extending the equipartition theorem to non integer
dimensions**

M.Praprotnik, K.Kremer and L.Delle Site, Phys.Rev.E, 75, 017701 (2007);

M.Praprotnik, K.Kremer and L.Delle Site, J.Phys.A:Math.Th.40, F281 (2007).

- **Still to address** : Why **forces** and not **potentials**



Coupling the atomistic and mesoscopic scales via a POTENTIAL approach

Why the forces and not the Potentials?

- Coupling potentials:

$$U_{\alpha\beta} = w(x_\alpha)w(x_\beta)U_{\alpha\beta}^{atom} + [1 - w(x_\alpha)w(x_\beta)]U_{\alpha\beta}^{cg}$$

- Consequence: $\mathbf{F}_{drift} = U_{atom} \frac{\partial w}{\partial x} + U_{cg} \frac{\partial w}{\partial x}$

(a) if considered *physical* \longrightarrow **violation of Third Newton's law**

M.Praprotnik, K.Kremer and L.Delle Site, J.Phys.A:Math.Th.40, F281 (2007)

(b) if removed \longrightarrow mathematical inconsistency

L.Delle Site, Phys.Rev.E, 76, 047701 (2007)



Violation of the Third Newton's law I

- if: $V_{\alpha\beta} = w(x_\alpha)w(x_\beta)V_{\alpha\beta}^{atom} + [1 - w(x_\alpha)w(x_\beta)]V_{\alpha\beta}^{cg}$
- Then: $\mathbf{F}_{\alpha\beta} = -\frac{\partial V_{\alpha\beta}}{\partial \mathbf{r}_\alpha}$
- $-\frac{\partial V_{\alpha\beta}}{\partial \mathbf{r}_\alpha} = w(x_\alpha)w(x_\beta)\mathbf{F}_{\alpha\beta}^{atom} + [1 - w(x_\alpha)w(x_\beta)]\mathbf{F}_{\alpha\beta}^{cg} + \left(\frac{\partial w(x_\alpha)}{\partial x_\alpha}w(x_\beta) \left[V_{\alpha\beta}^{cg} - V_{\alpha\beta}^{atom}\right], 0, 0\right)$
- The force acting on the molecule β is: $\mathbf{F}_{\beta\alpha} = -\frac{\partial V_{\alpha\beta}}{\partial \mathbf{r}_\beta}$
- $-\frac{\partial V_{\alpha\beta}}{\partial \mathbf{r}_\beta} = w(x_\alpha)w(x_\beta)\mathbf{F}_{\beta\alpha}^{atom} + [1 - w(x_\alpha)w(x_\beta)]\mathbf{F}_{\beta\alpha}^{cg} + \left(\frac{\partial w(x_\beta)}{\partial x_\beta}w(x_\alpha) \left[V_{\alpha\beta}^{cg} - V_{\alpha\beta}^{atom}\right], 0, 0\right)$
- where $\frac{\partial}{\partial \mathbf{r}} = \left(\frac{\partial}{\partial x}, \frac{\partial}{\partial y}, \frac{\partial}{\partial z}\right)$, $\mathbf{F}_{\alpha\beta}^{atom} = -\frac{\partial V_{\alpha\beta}^{atom}}{\partial \mathbf{r}_\alpha}$, and $\mathbf{F}_{\alpha\beta}^{cg} = -\frac{\partial V_{\alpha\beta}^{cg}}{\partial \mathbf{r}_\alpha}$.



Violation of the Third Newton's law II

- Using $\mathbf{F}_{\beta\alpha}^{atom} = -\mathbf{F}_{\alpha\beta}^{atom}$ and $\mathbf{F}_{\beta\alpha}^{cg} = -\mathbf{F}_{\alpha\beta}^{cg}$ we obtain:
- $-\mathbf{F}_{\beta\alpha} = w(x_\alpha)w(x_\beta)\mathbf{F}_{\alpha\beta}^{atom} + [1 - w(x_\alpha)w(x_\beta)]\mathbf{F}_{\alpha\beta}^{cg} - \left(\frac{\partial w(x_\beta)}{\partial x_\beta} w(x_\alpha) \left[V_{\alpha\beta}^{cg} - V_{\alpha\beta}^{atom} \right], 0, 0 \right)$
- In order that the force satisfies Newton's Third Law:
$$\frac{\partial w(x_\alpha)}{\partial x_\alpha} w(x_\beta) = -\frac{\partial w(x_\beta)}{\partial x_\beta} w(x_\alpha)$$
- implying that: $\frac{1}{w(x_\alpha)} \frac{\partial w(x_\alpha)}{\partial x_\alpha} = -\frac{1}{w(x_\beta)} \frac{\partial w(x_\beta)}{\partial x_\beta}$
- whose solution is $w(x_\alpha)$ and $w(x_\beta)$ are constants!



The Mathematical Problem I

- A generalization to the potentials using two generic weighting functions $f(x)$ and $g(x)$ writes: $U^{coupling} = f(X_\alpha, X_\beta)U_{cg} + g(X_\alpha, X_\beta)U_{atom}$

- $\mathbf{F}_{drift} = 0 \rightarrow$

$$\begin{aligned} U_{cg} \frac{\partial f(X_\alpha, X_\beta)}{\partial X_\alpha} + U_{atom} \frac{\partial g(X_\alpha, X_\beta)}{\partial X_\alpha} &= 0 \\ U_{cg} \frac{\partial f(X_\alpha, X_\beta)}{\partial X_\beta} + U_{atom} \frac{\partial g(X_\alpha, X_\beta)}{\partial X_\beta} &= 0. \end{aligned} \quad (8)$$

equations of first order, each 2 boundary conditions (**system overdetermined**)

- If generalized: $U^{coupling} = f(X_\alpha, X_\beta)U_{cg} + g(X_\alpha, X_\beta)U_{atom} + \Phi: \rightarrow$ overdetermination shifted from f and g to Φ .



The Mathematical Problem II

Above all:

- $\int [U_{cg} \frac{\partial f(x)}{x} + U_{atom} \frac{\partial g(x)}{x} U_{atom}] dx + \Phi_0 = \Phi(x)$
- cannot be a solution, since one **must** have either
 $\Phi(x_0) = \Phi_0 = 0$ (atomistic region) or
 $\Phi(x_1) = \Phi_0 = U_{atom} - U_{cg}$ (coarse-grained region)
but not both!
- Overdetermination of $\Phi \rightarrow$ Conserved Energy does not correspond to dynamical equations



Does the method work?

Comparison of the results of AdResS with those of full atomistic simulations:

- Global and local structures are reproduced
- There is no net flux across the transition region
- The diffusion occurs in a correct way
- Method successfully reproduces statistical properties of the model liquid.

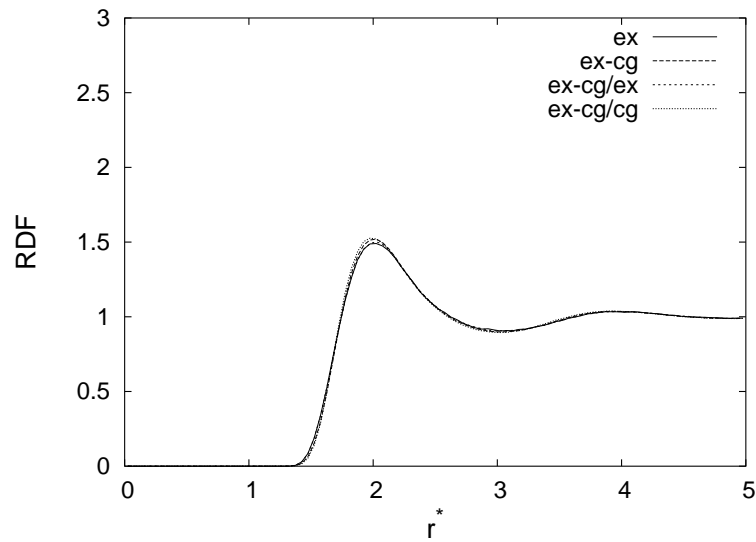
see in particular:

medium dense liquid: M.Praprotnik, L.Delle Site and K.Kremer, J.Chem.Phys.123,224106 (2005).

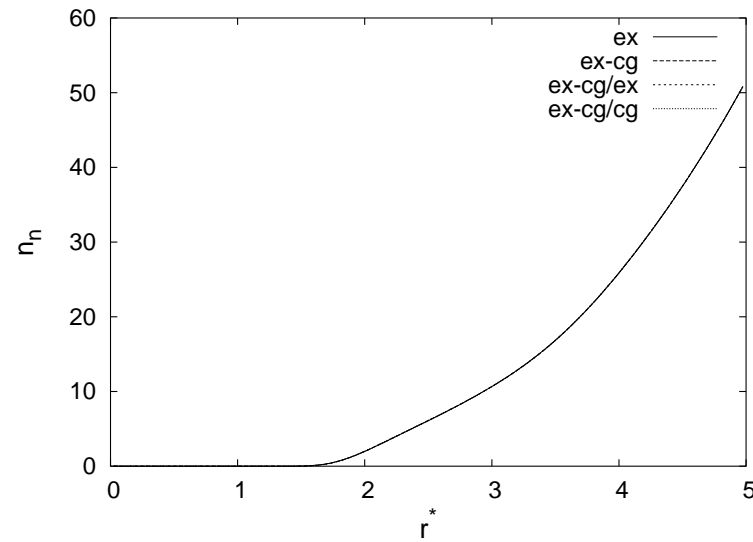
dense liquid: M.Praprotnik, L.Delle Site and K.Kremer, Phys.Rev.E, 73, 066701 (2006).



Results for a medium dense liquid



(k) RDF_{cm} .

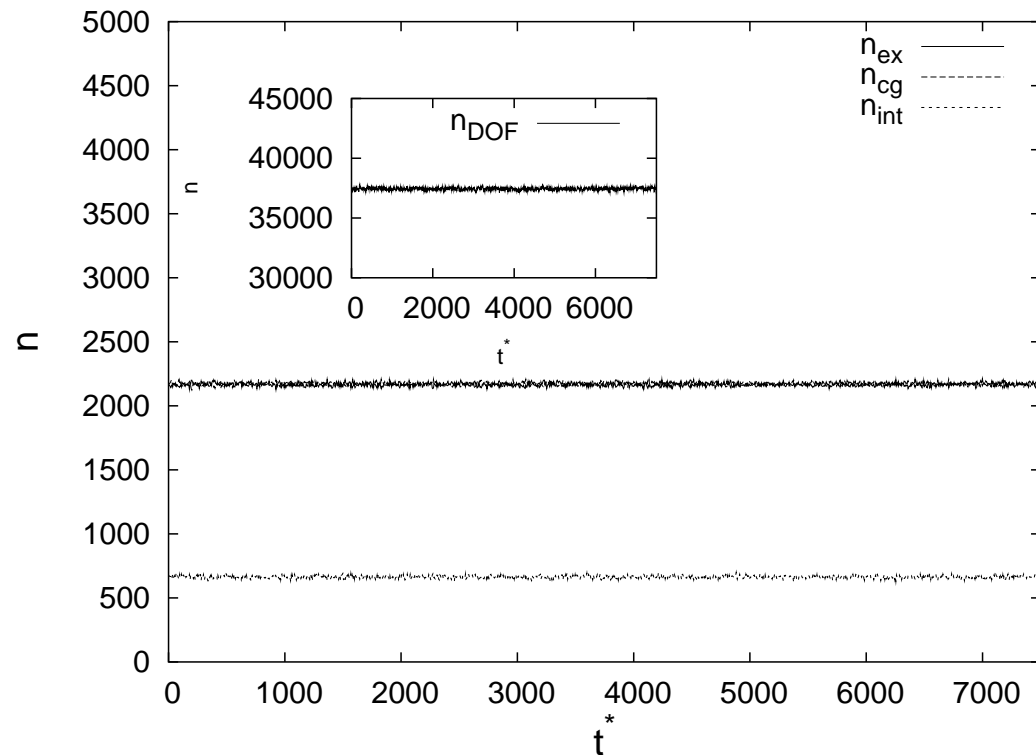


(l) Number of neighbors.

Global and local structure are properly reproduced



Number of DOFs

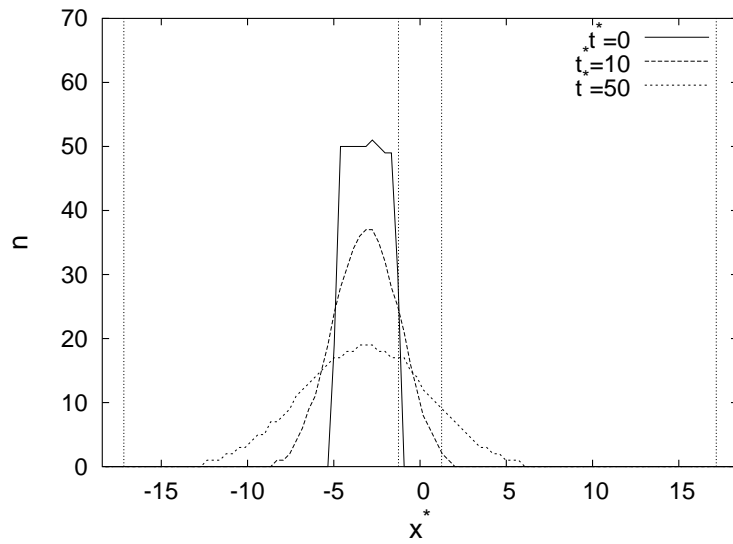


$$n_{DOS} = 3n[4w + (1 - w)]$$

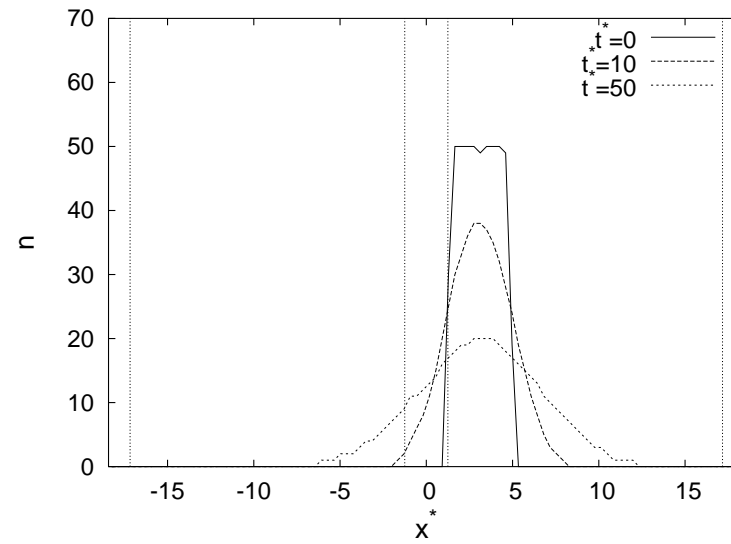
Numerical proof of the "Zero Flux"



Diffusion across the transition regime



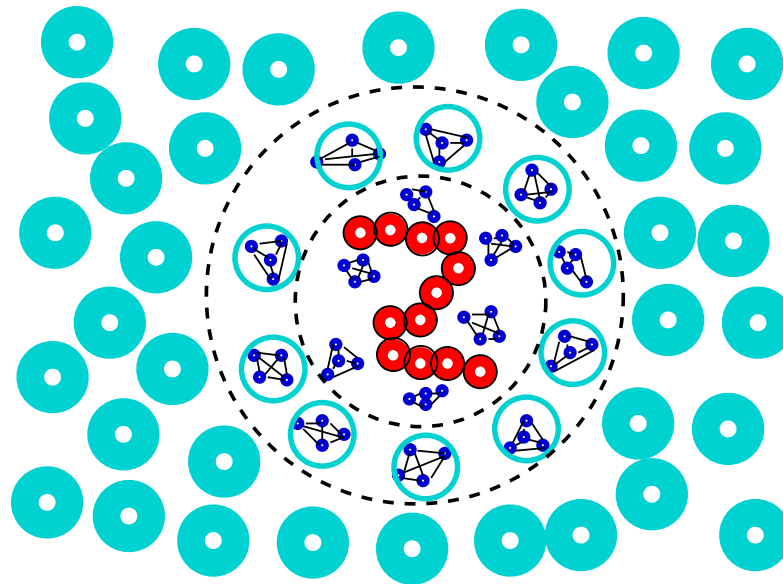
(m) Coarse-grained side.



(n) Explicit side.



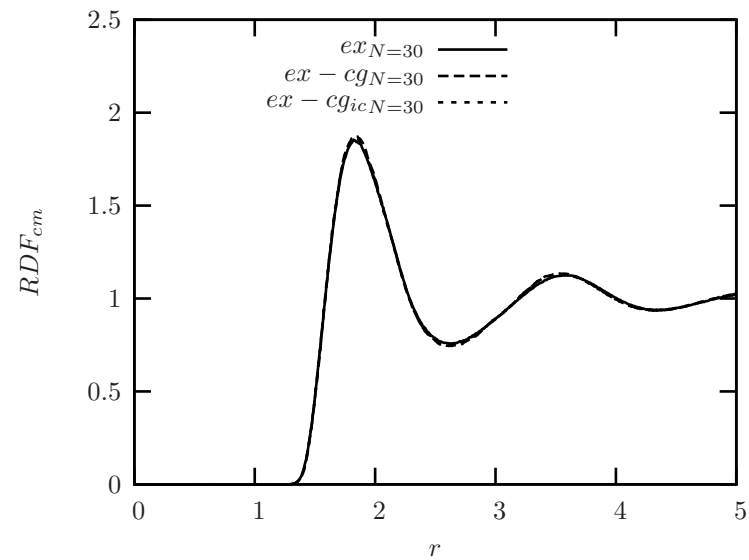
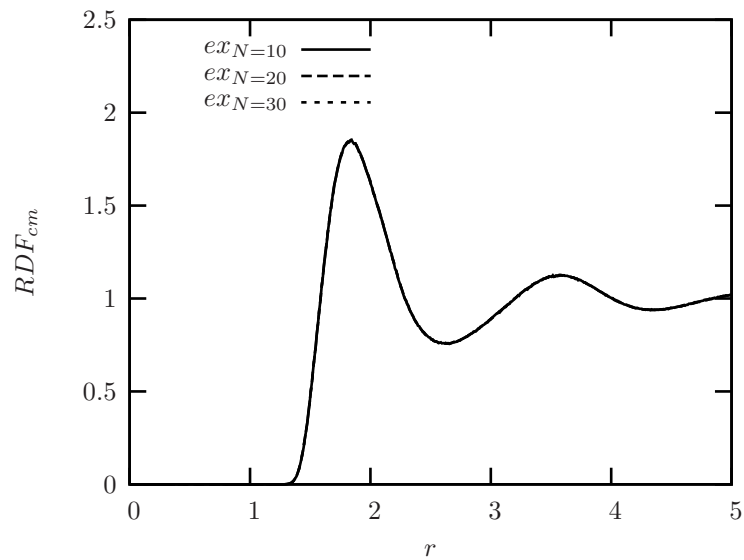
Further application: Solvating a Polymer



A generic Polymer solvated by a Tetrahedron Liquid

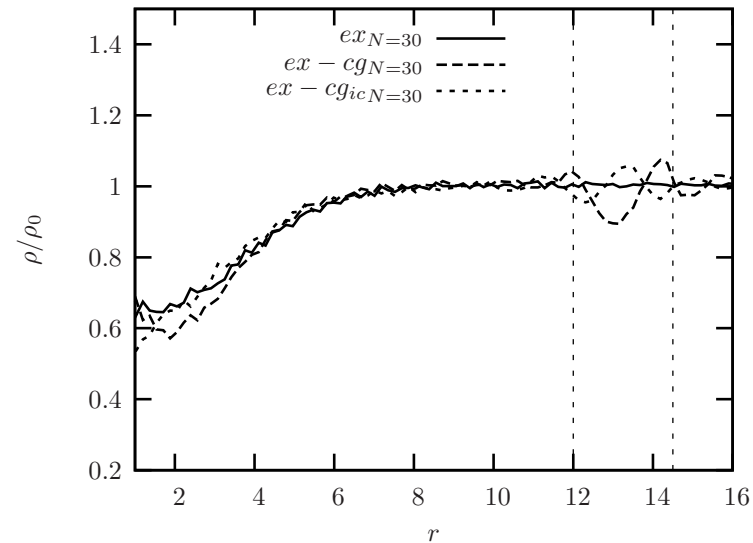
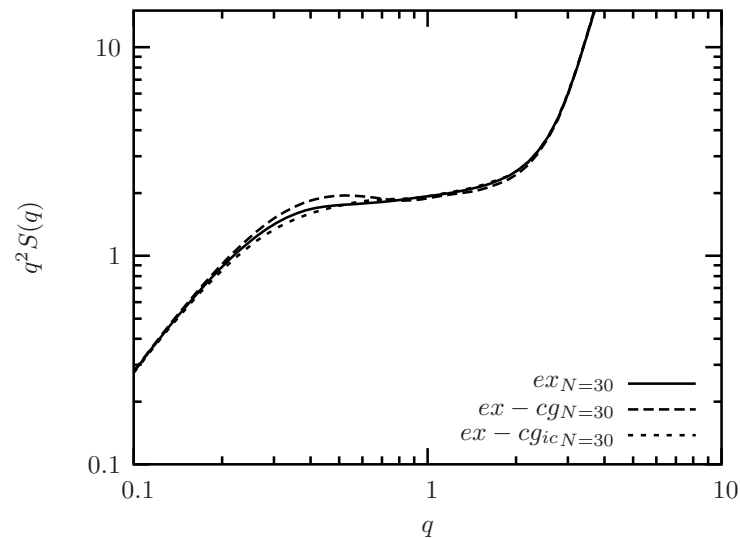


Static properties of the solvent





Static properties of the polymer

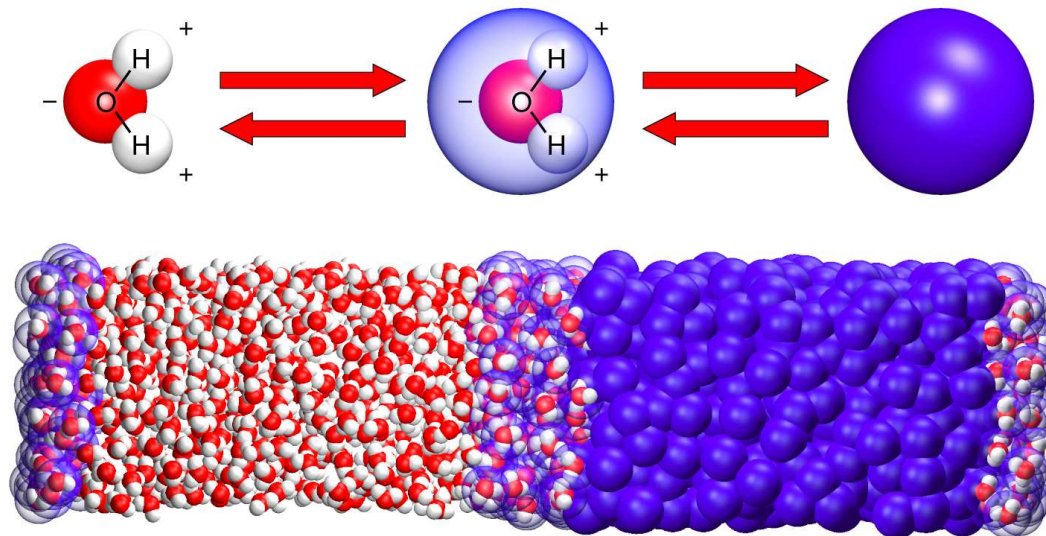


$$S(q) = \frac{1}{N} \left\langle \sum_{ij} \exp(i\mathbf{q} \cdot (\mathbf{r}_i - \mathbf{r}_j)) \right\rangle$$

M.Praprotnik, L.Delle Site and K.Kremer, J.Chem.Phys.126 134902 (2007).



Liquid Water



The simulation speed-up is $\sim 17 - 20$ compared to atomistic simulations
M.Praprotnik, S. Matysiak, L. Delle Site, K. Kremer, C. Clementi, J. Phys.:
Condens.Matter, 19, 292201, 2007.



Reaction field method for Electrostatics

The electrostatic forces interactions are described using the Reaction field (RF) method:

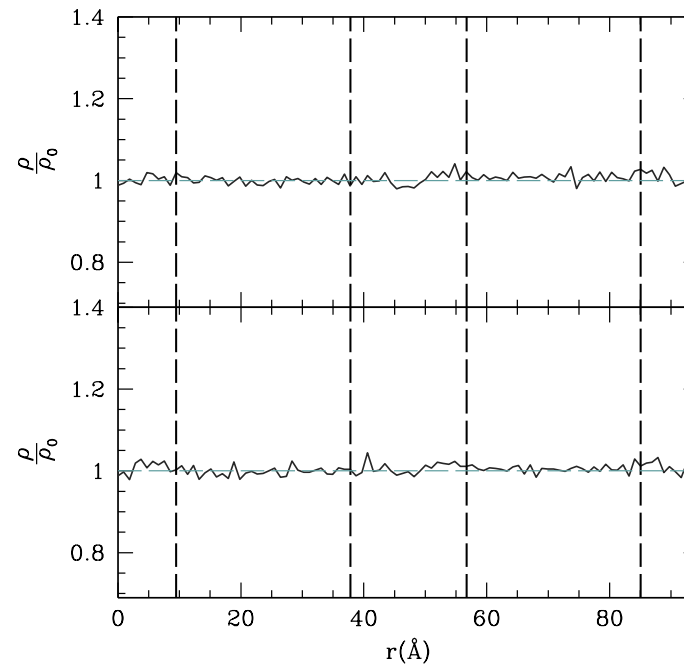
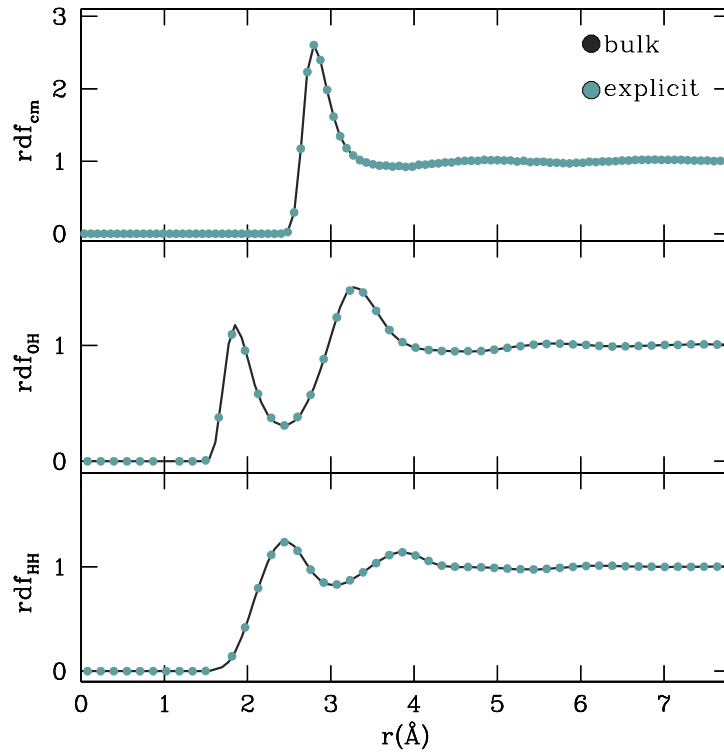
$$\mathbf{F}_{C_{i\alpha j\beta}}^{atom}(\mathbf{r}_{i\alpha j\beta}) = \frac{e_{i\alpha} e_{j\beta}}{4\pi\epsilon_0} \left[\frac{1}{r_{i\alpha j\beta}^3} - \frac{1}{R_c^3} \frac{2(\epsilon_{RF} - 1)}{1 + 2\epsilon_{RF}} \right] \mathbf{r}_{i\alpha j\beta}.$$

The RF is suitable to be used with AdResS because:

- it is pairwise
- like AdResS it must also be applied with a thermostat



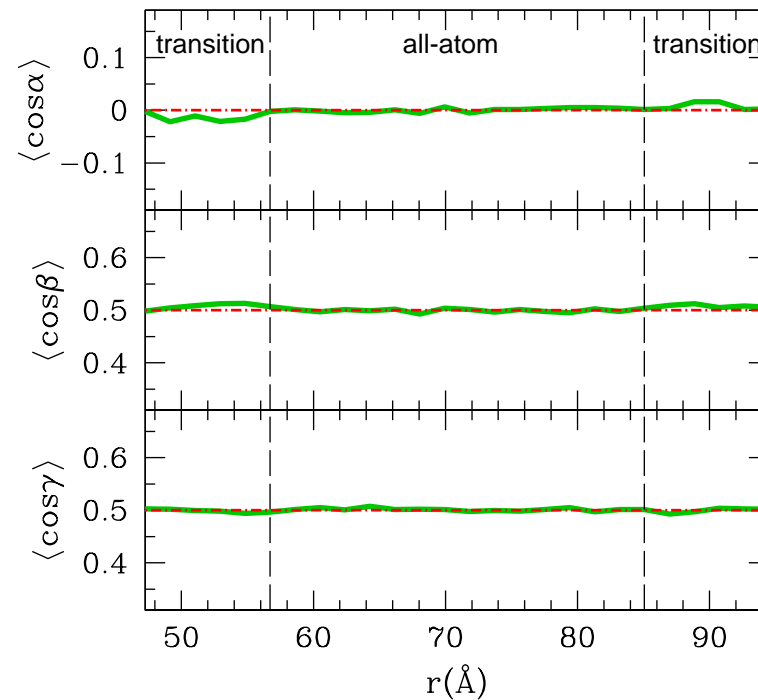
Results for water



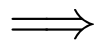
S.Matysiak, C.Clementi, M.Praprotnik, K.Kremer and L.Delle Site;
J.Chem.Phys.128, 024503 (2008).



Interface effect of the cg water



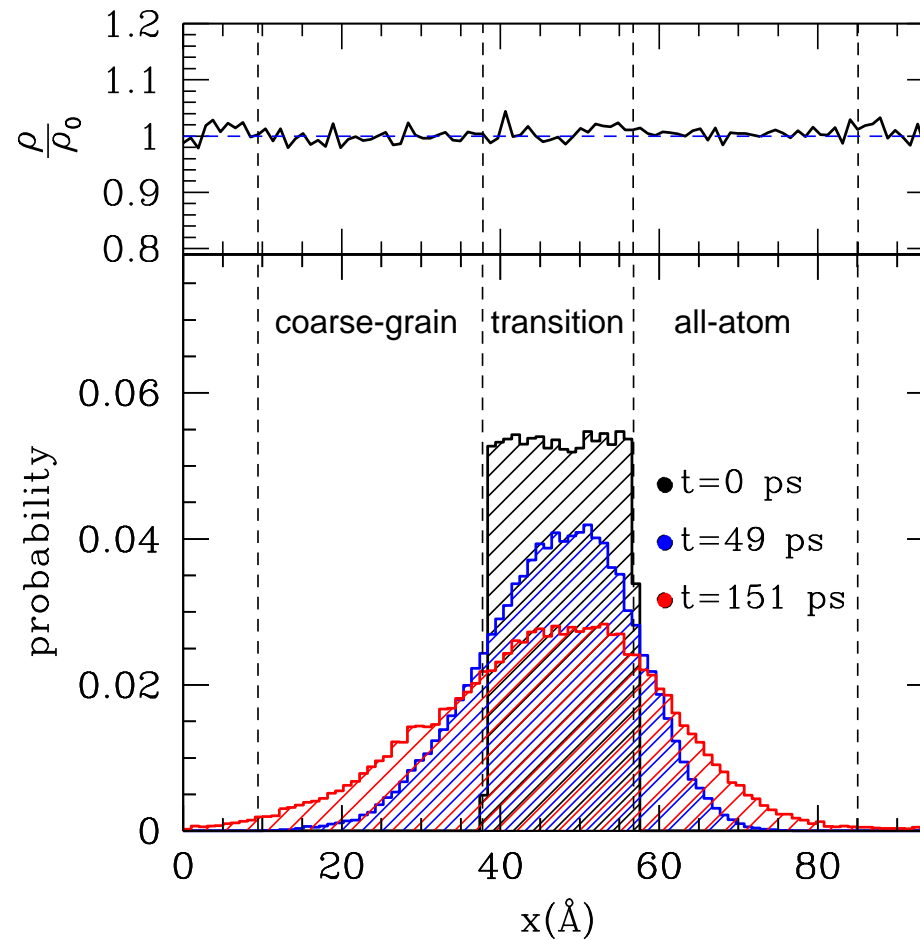
The transition regime neutralizes the interface effect of the cg water



the structure of water in the explicit regime is the same as in the bulk.



Diffusion across the transition regime





Position dependent Langevin thermostat

The Langevin equation with a position dependent coefficient $\Gamma(x)$ can be written as:

$$m_i dv_i/dt = F_i - m_i \Gamma(x) v_i + R_i(x, t) \quad (9)$$

where $R_i(x, t)$ is:

$$\langle R_i(x, t) \rangle = 0, \quad (10)$$

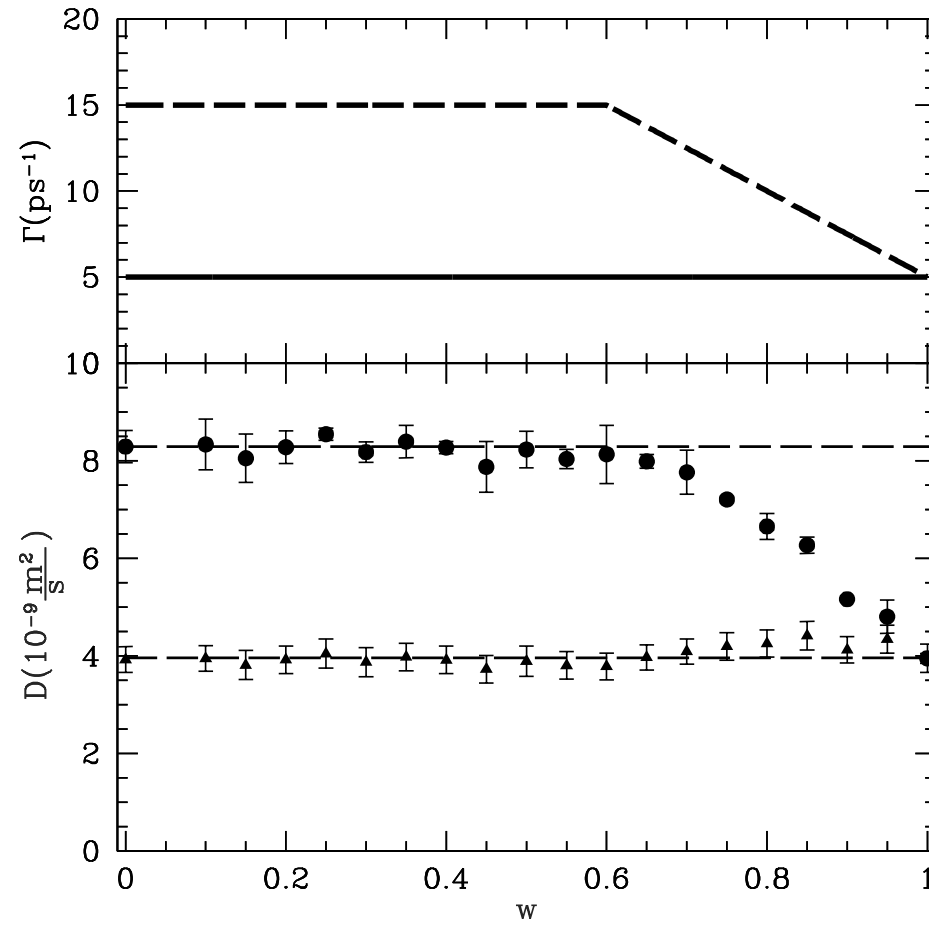
$$\langle R_i(x, t_1) R_j(x, t_2) \rangle = 2\Gamma(x) m_i kT \delta(t_1 - t_2) \delta_{ij} \quad (11)$$

$$\Gamma(x) = \begin{cases} \Gamma_{cg} & \text{if } x \leq 0.6 \\ \alpha x + \beta & \text{if } 0.6 < x \leq 1.0 \end{cases} \quad (12)$$

This choice provides a simple interpolation between the two limit values of $\Gamma(0.6) = \Gamma(0) = \Gamma_{coarse-grained}$ and $\Gamma(1) = \Gamma_{all-atom}$

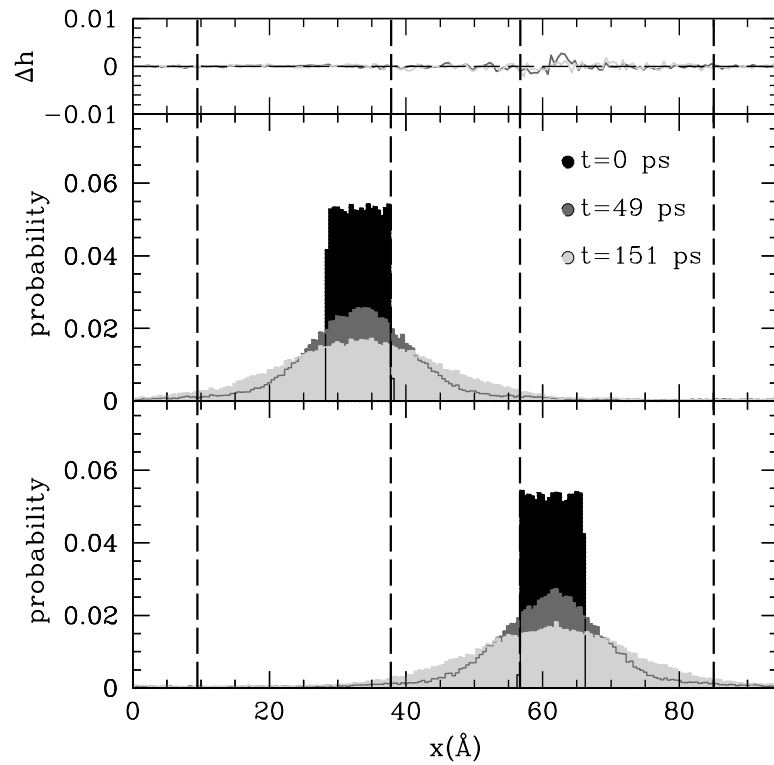


Diffusion across the simulation box I

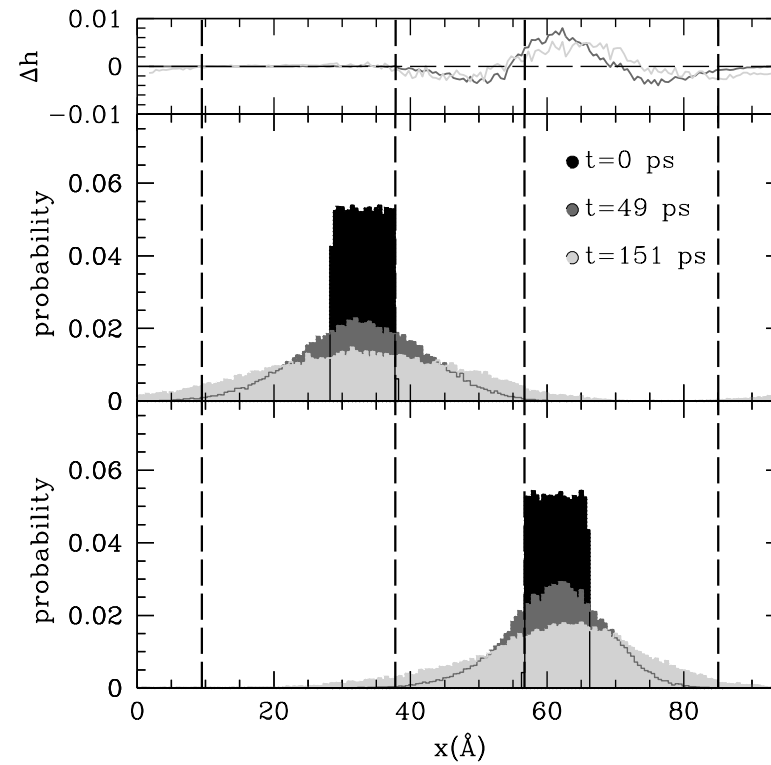




Diffusion coefficient across the simulation box II.



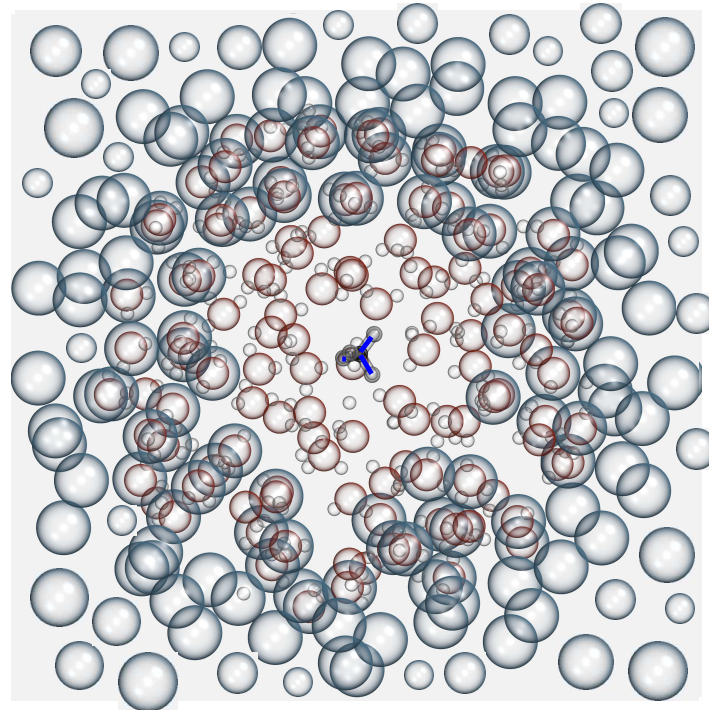
(v) Position dependent thermostat



(w) Regular thermostat



Next: Hydrophobic solvation via adaptive simulation



What is the range of influence of the hydrogen bonding network of the bulk in the formation of the hydration structure around the solute?