Multiscale Modeling

Winterschool on Theoretical Chemistry and Spectroscopy 10-14 December, 2012

> **Bernd Ensing** University of Amsterdam





Thursday, December 13, 2012

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Background

Car-Parrinello MD study of chemical reactions in water

Rare events, free energy methods, sampling methods

Hybrid multiscale MD

Photoactive proteins

Electro-chemistry, redox reactions

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metadynamics



Multiscale systems

Molecular systems may exhibit phenomena on a range of scales in time, space or energy that are coupled to each other



Potassium or proton transport channel



lon and molecule diffusion in polyelectrolyte



Enzymatic catalysis



Nanoparticle assembly on a drying surface



Signal transduction in Gcoupled sensor proteins



vesicle-membrane fusion

The Schrödinger equation cannot be solved for the entire system...

Multiscale systems

"Everything should be made as simple as possible, but not simpler."

Multiscale systems

"Everything should be made as simple as possible, but not simpler."



Multiscale methods



content

- Lecture 1: QM/MM
- Lecture 2: Density embedding
- Lecture 3: Coarse-graining
- Lecture 4: Hybrid multiscale molecular dynamics



J. Mol. Biol. (1976) 103, 227-249



Theoretical Studies of Enzymic Reactions:

Dielectric, Electrostatic and Steric Stabilization of the Carbonium Ion in the Reaction of Lysozyme

A. WARSHEL AND M. LEVITT

Medical Research Council Laboratory of' Molecular Biology Hills Road, Cambridge CB2 2QH, England

and

Department of Chemical Physics The Weixmann Institute of Science Rehovot, **Israel**

(Received 12 September 1975, and in revised form 10 February 1976)

A general method for detailed study of enzymic reactions is presented. The method considers the complete enzyme-substrate complex together with the surrounding solvent and evaluates all the different quantum mechanical and classical energy factors that can affect the reaction pathway. These factors



QM/MM pioneers

Theoretical Studies of Enzymic Reactions: Dielectric, Electrostatic and Steric Stabilization of Reaction of Lysozyme the Carbonium Ion in the A. Warshel AND M. Levitt

J. Mol. Biol. 103, 227-249 (1976)

A combined ab initio quantum mechanical and molecular mechanical method for carrying out simulations on complex molecular systems: Applications to the CH3CI + CI– exchange reaction and gas phase protonation of polyethers

U. Chandra Singh, Peter A. Kollman

J. Comput. Chem. 7, 718–730 (1986)

A combined quantum mechanical and molecular mechanical potential for molecular dynamics simulations

Martin J. Field, Paul A. Bash, Martin Karplus

J. Comput. Chem. 11, 700-733 (1990)



nature

Vol 455 30 October 2008 doi:10.1038/nature07297

LETTERS

Low-speed fracture instabilities in a brittle crystal

J. R. Kermode¹, T. Albaret², D. Sherman³, N. Bernstein⁴, P. Gumbsch^{5,6}, M. C. Payne¹, G. Csányi⁷ & A. De Vita^{8,9}

When a brittle material is loaded to the limit of its strength, it fails by the nucleation and propagation of a crack¹. The conditions for ph the uncleation and biobagation of a crack¹. The conditions to ph the uncleation and biobagation of a crack¹. The conditions for for different cleavage planes. For a given cleavage plane, it can lead to slow crack growth^{1,4,16} and anisotropy with respect to propagation to slow crack disconterval and anisotropy with respect to biobadation to slow crack disconterval. For a given cleavage blane, it can lead

J. R. Kermode¹, T. Albaret², D. Sherman², N. Bernstein^{*}, P. Gumbsch^{2,e}, M. C. Payne¹, G. Csányi⁷ & A. De Vita^{8,3}







Proton Shuttles and Phosphatase Activity in Soluble Epoxide Hydrolase





QM/MM Energy function

Coupling between the QM and MM regions



 $E_{\text{tot}} = E_{\text{QM}}(I) + E_{\text{MM}}(II)$ I: QMI: MM

 $E_{\text{tot}} = E_{\text{MM}}(I + II) + E_{\text{QM}}(I) - E_{\text{MM}}(I)$

$$E_{\text{tot}} = E_{\text{QM}}(I) + E_{\text{MM}}(II) + E_{\text{QM}-\text{MM}}(I,II)$$



Electronic structure calculation

 $H\psi = E\psi$

Semi-empirical Tight binding, AM1, PM3,...

Density Functional Theory

Hartree-Fock

MP2, CC, CI, CAS-PT2,...

Most QM/MM calculations use a semi-empirical approach for the QM part. The Car-Parrinello molecular dynamics community (CPMD, CP2K) uses DFT.



Density Functional Theory

Kohn-Sham

$$E_0(\mathbf{R}) = \min_{\Psi_i} E_{\mathrm{KS}}[\Psi_i, \mathbf{R_i}]$$

$$E_{\rm KS}[\Psi_i, \mathbf{R_i}] = \frac{1}{2} \sum_i \int d\mathbf{r} \, \psi_i^*(\mathbf{r}) \nabla^2 \psi_i(\mathbf{r}) + \int d\mathbf{r} \, V_N(\mathbf{r}) \rho(\mathbf{r}) \\ + \frac{1}{2} \int d\mathbf{r} d\mathbf{r}' \, \frac{\rho(\mathbf{r})\rho(\mathbf{r}')}{\mathbf{r} - \mathbf{r}'} + E_{\rm xc}[\rho(\mathbf{r})]$$

$$\rho(\mathbf{r}) = 2\sum_{i} \psi_{i}^{*}(\mathbf{r})\psi_{i}(\mathbf{r})$$
$$\int d\mathbf{r} \,\psi_{i}^{*}(\mathbf{r})\psi_{j}(\mathbf{r}) = \delta_{ij}$$

MM

Molecular Mechanics

- Empirical Forcefield based on pair-wise potential functions
- Commonly used in classical molecular dynamics simulations
- MM2, AMBER, CHARMM, UFF, CFF, CVFF, GROMOS, OPLS,...

$$\begin{split} V_{\rm MM}(\mathbf{R}) &= \frac{1}{2} \sum_{N_b} k_b (b-b_0)^2 + \frac{1}{2} \sum_{N_a} k_a (a-a_0)^2 & \text{bonds + bends} \\ &+ \sum_{N_t} k_t [1 + \cos(mt-t_0)] + \frac{1}{2} \sum_{N_o} k_o (o-o_0)^2 & \text{dihedral + out-of-plane} \\ &+ \sum_{I} \sum_{J>I} \left(\frac{C_{12,IJ}}{R_{IJ}^{12}} - \frac{C_{6,IJ}}{R_{IJ}^6} + \frac{q_I q_J}{R_{IJ}} \right) & \text{Vanderwaals + electrostatics} \end{split}$$

... + Charge-Dipole Dipole-Dipole

Coupling QM and MM

- Mechanical embedding only: bonds, angles, VdW
- Electrostatic embedding only: point charges-
- Electrostatic + mechanical
- QM system feels MM environment
- MM environment feels QM system via MM interactions
- MM environment feels QM system in a self-consistent manner

Thursday, December 13, 2012

Electrostatic coupling is needed to include environment interaction in the QM system.

- polarization of electron density



- can

$$V_{\rm QM/MM} = \sum_{i \in \rm MM} q_i \int \frac{\rho_{\rm QM}(r)}{|r - r_i|} dr$$

- MM charges interact with QM charges
 - QM charges are obtained from density
 - but how? population analysis not rigorous
 - more efficient but more approximate

$$V_{\text{QM/MM}} = \sum_{i \in \text{MM}} \sum_{j \in \text{QM}} \frac{q_i q_j(\rho_{\text{QM}})}{|r_j - r_i|}$$





Electrostatic coupling

Electrostatic coupling is needed to include environment interaction in the QM system.

- polarization of electron density



No MM Pauli repulsion

- charge spilling into MM part
 - charge delocalization
 - MM pseudopotentials

MM multipole expansion

Polarization is one-way

- polarizable forcefield

Periodic boundary conditions?

- Ewald
- Reaction Field

QM/MM partitioning

Cutting through chemical bonds: QM/MM bonds Example: peptide chain



QM/MM partitioning

Cutting through chemical bonds: QM/MM bonds Example: peptide chain



QM/MM partitioning

Cutting through chemical bonds: QM/MM bonds Example: peptide chain





Add link atoms

- only seen by QM atoms?
- H, CH3, halogen atom
- Electronegativity of link atom should match MM part
- What about spectroscopy, excited states?

Or use frozen orbitals ("frozen bonds")

Improving MM

Force matching, fitting, learning on the fly,...



QM/QM

Frozen density embedding

T. A. Wesolowski, A. Warshel, *J. Phys. Chem.* **97**, 8050 (1993) T. A. Wesolowski, In Computational Chemistry: Reviews of Current Trends, Vol. 10; Leszczynski, J., Ed.; World Scientific: Singapore, 2006.



Christoph R. Jacob, Johannes Neugebauer, Lucas Visscher *J. Comput. Chem.* **29** 1011–1018 (2008)



The total density is a sum of the densities ρ_I and ρ_{II} in different regions or subsystems

Density ρ_{II} of the environment is kept fixed The total density $\rho_{tot} = \rho_I + \rho_{II}$ is obtained by optimizing ρ_I

KS-like equation with embedding term due to ρ_{II}

$$\begin{bmatrix} -\frac{\nabla^2}{2} + V_{\text{eff}}^{\text{KS}}[\rho_{\text{I}}](\boldsymbol{r}) + V_{\text{eff}}^{\text{emb}}[\rho_{\text{I}}, \rho_{\text{II}}](\boldsymbol{r}) \end{bmatrix}$$
$$\phi_i^{(\text{I})}(\boldsymbol{r}) = \varepsilon_i \phi_i^{(\text{I})}(\boldsymbol{r}), \quad i = 1, \dots, N_{\text{I}}.$$

$$\begin{split} V_{\text{emb}}^{\text{eff}}[\rho_{\text{I}},\rho_{\text{II}}](\boldsymbol{r}) &= \sum_{A_{\text{II}}} -\frac{Z_{A_{\text{II}}}}{|\boldsymbol{r}-\boldsymbol{R}_{A_{\text{II}}}|} + \int \frac{\rho_{\text{II}}(\boldsymbol{r}')}{|\boldsymbol{r}'-\boldsymbol{r}|} \, \mathrm{d}\boldsymbol{r}' \\ &+ \frac{\delta E_{\text{xc}}[\rho]}{\delta \rho} \Big|_{\rho=\rho_{\text{tot}}} - \frac{\delta E_{\text{xc}}[\rho]}{\delta \rho} \Big|_{\rho=\rho_{\text{I}}} \\ &+ \frac{\delta T_{s}[\rho]}{\delta \rho} \Big|_{\rho=\rho_{\text{tot}}} - \frac{\delta T_{s}[\rho]}{\delta \rho} \Big|_{\rho=\rho_{\text{I}}}, \end{split}$$

Embedding potential

- approximate kinetic energy

functional (Thomas-Fermi, PW91k)

- accurate for weakly bound systems

non-additive kinetic energy functional

$$T_s^{\text{nadd}}[\rho_{\mathrm{I}}, \rho_{\mathrm{II}}] = T_s[\rho_{\mathrm{I}} + \rho_{\mathrm{II}}] - T_s[\rho_{\mathrm{I}}] - T_s[\rho_{\mathrm{II}}]$$

If $\rho_{tot} - \rho_{II}$ is positive definite and non-interacting v_s representable, the solution give exact ground-state.



Acetonitrile and 15, 25, 40 water molecules



Christoph R. Jacob, Johannes Neugebauer, Lucas Visscher *J. Comput. Chem.* **29** 1011–1018 (2008)



Example application: NMR shifts for Acetonitrile-Water clusters

Table 1. Solvent Shifts $\Delta \sigma$ of the Nitrogen NMR Shielding in Acetonitrile–Water Clusters with 15, 25, and 40 Water Molecules, Respectively.

Isolated	15 H ₂ O Δσ (ppm)		25 H ₂ O Δσ (ppm)		40 H ₂ O $\Delta \sigma$ (ppm)	
	0.0		0.0		0.0	
SumFrag	12.6		11.9		11.8	
1 H ₂ O relaxed	12.9	+0.3	12.3	+0.4	11.9	+0.1
2 H ₂ O relaxed	13.3	+0.4	12.7	+0.4	12.4	+0.5
3 H ₂ O relaxed	13.7	+0.4	13.3	+0.6	13.1	+0.7
8 H ₂ O relaxed	14.6	+0.9	13.9	+0.6	14.0	+0.9
13 H ₂ O relaxed	15.3	+0.7	14.7	+0.8	14.7	+0.7
23 H ₂ O relaxed			15.2	+0.5	14.6	-0.1
38 H ₂ O relaxed					14.6	0.0
Supermolecule	14.0		13.1		12.5	

In the FDE calculations, the closest two water molecules have been included in the nonfrozen subsystem, for the remaining frozen fragments different approximations have been employed. For comparison, also the results of a conventional, supermolecular calculation are given. See text for details.



Example application: NMR shifts for Acetonitrile-Water clusters

Table 2. Wall Clock Time (in Minutes) Required for the Calculation of the Nitrogen NMR Shielding in Acetonitrile–Water Clusters with 15, 25, and 40 Water Molecules, Respectively, on 8 Dual Processor Nodes of an Intel Xeon 3.4 GHz Cluster, Using Different Approximations for the Frozen Density (see text for details).

	$15 H_2O$	$25 \text{ H}_2\text{O}$	$40 \text{ H}_2\text{O}$
Isolated	0.4	0.4	0.4
SumFrag	1.7	2.1	2.9
1 H2O relaxed	2.6	3.1	4.3
2 H ₂ O relaxed	2.8	3.6	5.0
3 H ₂ O relaxed	3.0	3.8	5.0
8 H ₂ O relaxed	3.6	5.2	6.2
13 H ₂ O relaxed	4.0	5.3	7.2
23 H ₂ O relaxed		7.1	9.4
38 H ₂ O relaxed			12.8
Supermolecule	8.5	29.8	103.5

Polarizable QM/MM

Discreet reaction field model

Solvent is represented by charges q_s and polarizabilities α_s places at positions \mathbf{r}_s

Kohn-Sham like equations

$$\begin{bmatrix} -\frac{\nabla^2}{2} + V_{\text{eff}}^{\text{KS}}[\rho](\mathbf{r}) + V^{\text{DRF}}[\rho](\mathbf{r}) \end{bmatrix} \phi_i(\mathbf{r}) = \varepsilon_i \phi_i(\mathbf{r})$$

$$i = 1, \dots, N.$$

with a Coulomb term due to permanent charges and a many-body polarization term:

$$V^{\text{DRF}}[\rho](\mathbf{r}) = V^{\text{el}}(\mathbf{r}) + V^{\text{pol}}[\rho](\mathbf{r})$$
$$= \sum_{s} \frac{q_s}{|\mathbf{r} - \mathbf{R}_s|} + \sum_{s} \boldsymbol{\mu}_s^{\text{ind}} \cdot \frac{(\mathbf{r} - \mathbf{R}_s)}{|\mathbf{r} - \mathbf{R}_s|^3}$$

The MM induced atomic dipole:

$$\mu_s^{\text{ind}} = \alpha_s \left[\mathbf{F}_s^{\text{init}} + \sum_{t,t \neq s} \boldsymbol{T}_{st}^{(2)} \boldsymbol{\mu}_t^{\text{ind}} \right]$$

Г

with **T** the screened dipole interaction tensor of interaction between sites. Field **F** at site s is due to QM charge distribution, QM nuclei, MM charges, MM dipoles. Self-consistent procedure!

B. T. Thole and P. Th. van Duijnen, Theor. Chim. Acta 55 307-318 (1980)

Т

Implicit solvent models

Polarizable Continuum Model (C-PCM, D-PCM)

COSMO: a new approach to dielectric screening in solvents with explicit expressions for the screening energy and its gradient S. Miertus, E. Scrocco, J. Tomasi, *Chem. Phys.* **55**, 117 (1981). R. Cammi, J. Tomasi, *J. Comput. Chem.* **16**, 1449 (1995).

Continuum Solvation model (COSMO)

COSMO: a new approach to dielectric screening in solvents with explicit expressions for the screening energy and its gradient A. Klamt and G. Schüürmann

J. Chem. Soc., Perkin Trans. 2, 1993, 799-805



Reviews:

- C. J. Cramer and D. G. Truhlar, *Chem. Rev.* **99**, 2161-2200 (1999)
- J. Tomasi, B. Mennucci and R. Cammi, Chem. Rev. 105, 2999-3094 (2005)

Implicit solvent models

Solute molecule in continuous solvent

Based on Onsager's reaction field method

- Solute is treated explicitly at QM level
- Only electrostatic solvent-solute interactions
- Solute is placed in a void cavity
 - shape of cavity is important; different methods
 - should contain most of solute electron density
 - should exclude the solvent
 - e.g. iso-density surface
 - or iso-energy surface (with He/Ar atom), SAS
 - or simply van der Waals radii

- Solvent is represented by continuous dielectric medium

- Charge distribution of the solute, inside the cavity, polarizes the dielectric continuum, which in turn polarizes the solute charge distribution

 $-\nabla^2 V(\vec{r}) = 4\pi \rho_{\rm M}(\vec{r})$ within C

$$-\epsilon \nabla^2 V(\vec{r}) = 0$$
 outside C



Summary QM/MM

Many algorithms and implementations available

QSite, Jaguar program, Schrödinger QM: Hartree Fock, DFT, MP2

GROMACS-CPMD (Rothlisberger) QM: DFT/plane-waves

QMMM Minnesota, (Truhlar) QM package: GAMESS, Gaussian, ORCA + MM package: TINKER

ChemShell TCL interface (Daresbury) Combines: GAMESS-UK, DL_POLY, MNDO97, TURBOMOLE, CHARMM, GULP, Gaussian94

ONION, IMOMM (Morokuma) also in ADF: DFT + Amber/Sybyl forcefields

QM/MM can not (yet) be used as a black box method

- many parameters
- quality of QM model, MM model
- quality of QM/MM coupling
- be careful: for every approach a situation can be concocted such that it will fail