



# TOWARDS QUANTITATIVE MODELLING OF BIOLOGICAL AND COMPLEX SYSTEMS USING POLARIZABLE FORCE FIELDS.

J.-P. Piquemal

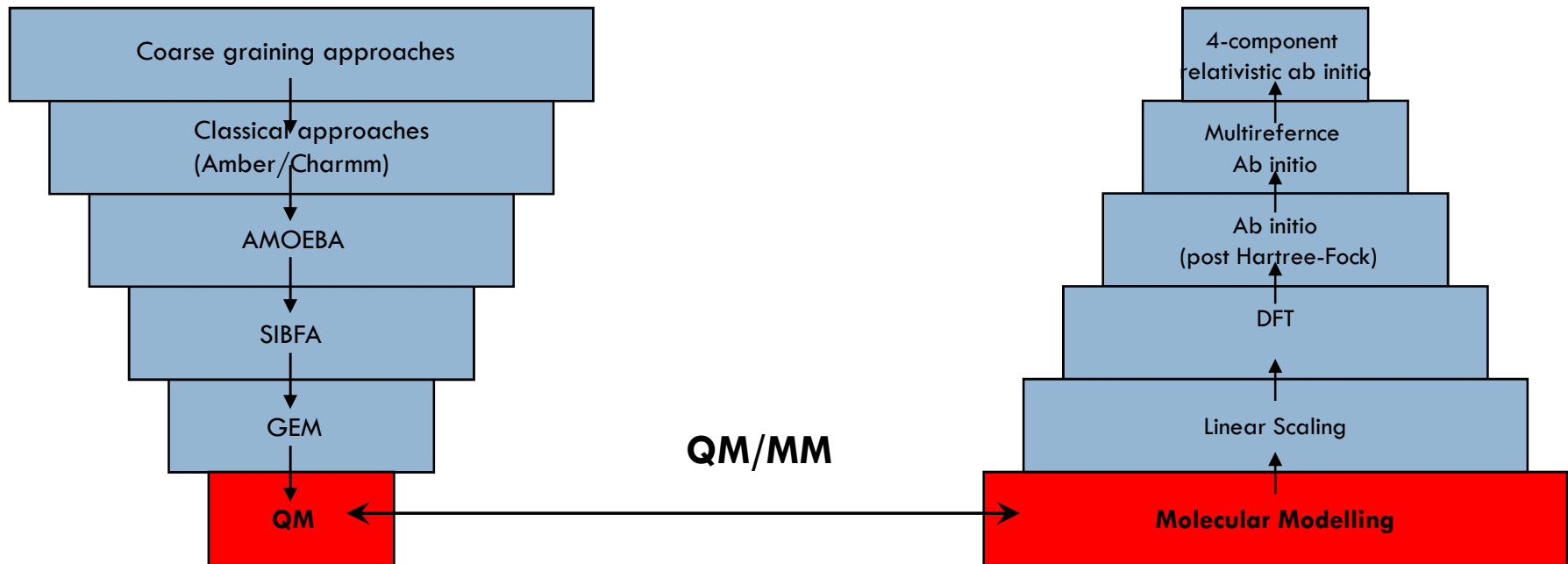
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Université Pierre et Marie Curie

10/16/2009

Ab Initio 2009, Nice, France.

## Potential energy surfaces (PES):



**MM**

**(non reactive)**

**QM**

**(access to chemical reactivity)**

**Unfortunately...our interests are:**

**metalloproteins, transition/heavy metal cations in solution, weak interactions...**

# Philosophy

## From QM computations to condensed phase molecular dynamics

Force fields advantages: fast evaluation of PES

Possible treatment of large molecules (million atoms) directly or through QM/MM

-structure

-dynamique

-interaction (docking)

-IR spectrum

$$\begin{aligned} V_T = & \sum_{1,2\text{pairs}} 0.5 K_b (b - b_0)^2 \\ + & \sum_{\text{bond angles}} 0.5 K_\theta (\theta - \theta_0)^2 \\ + & \sum_{\text{dihedral angles}} K_\phi [ 1 + \cos(n\phi - \delta) ] \\ + & \sum_{i,j} 4\epsilon_{ij} [ (\sigma_{ij}/r)^{12} - (\sigma_{ij}/r)^6 ] \\ + & q_i q_j / \epsilon r \end{aligned}$$

**The link with QM is broken...**

**Next generation "polarizable" force fields are needed.**

# Philosophy

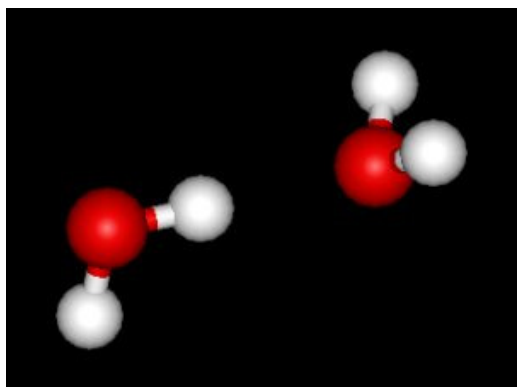
## From QM computations to condensed phase molecular dynamics

Unravelling physics at place in a complex (large) system:  
The right tool for the right problem

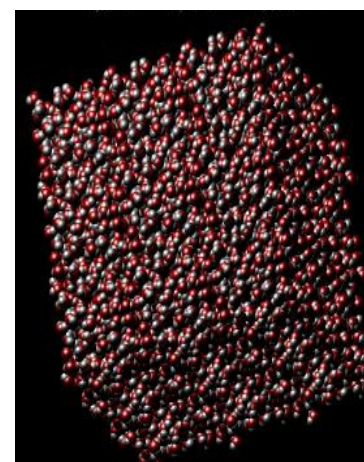
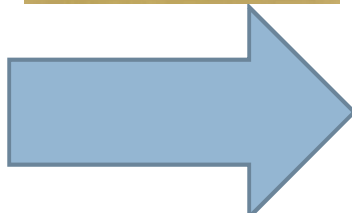
### IMPORTANCE OF QM INTERPRETATIVES TECHNIQUES

- density partition schemes
- energy decomposition analysis

Beyond QM capabilities: insights from advanced polarizable MM methods  
« Bottom-up » approach for quantitative results (general MM equations)



QM



MM or QM/MM

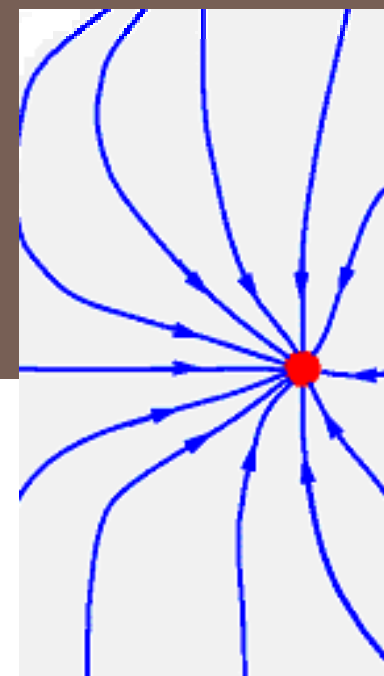
# QM interpretative techniques

## Electron Localization Function (ELF)

ELF : built upon the probability of finding an electron of a given spin in the vicinity of another same spin electron.  $0 < \text{ELF} < 1$ .

- Measure of Pauli exclusion principle associated to a local excess of kinetic energy.
- When  $\text{ELF} \rightarrow 1$ : high probability of opposite-spin pairs
- Maximal probability site are called attractors

Theory of dynamical systems:  
bassins and separatrix



\* A.D. Becke, K.E. Edgecombe, *J. Chem. Phys.* 92, 5397, 1990

\*\* B. Silvi, A. Savin, *Nature* 371, 683, 1994

# QM interpretative techniques

## ELF Topological Analysis

3D partition of the molecular space (3D grid) into chemically intuitive bassins:

Access to non atomic bassins: lone pairs,  $\pi$  systems etc...

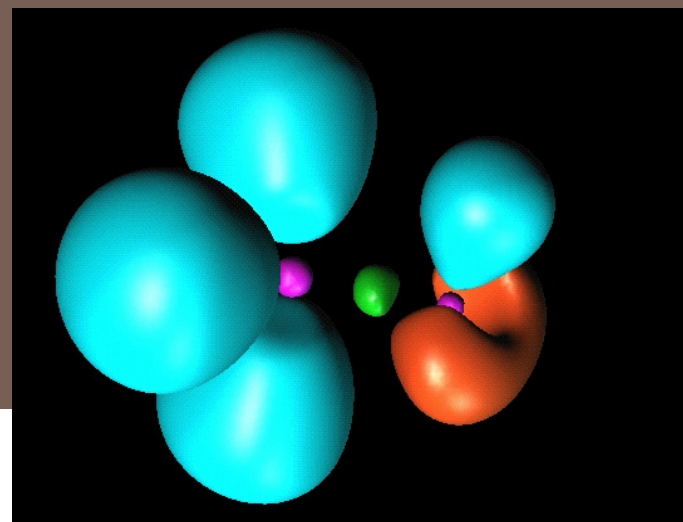
Several type of bassins exist:

Core (violet)

Disynaptic (green) : covalent bond

Protonated Disynaptic (blue) : C-H bond

Monosynaptic (red) : Lewis-type lone pair



*Valence Shell Electron Pair Repulsion (VSEPR)  
from Gillespie.*

# QM interpretative techniques

## Distributed Electrostatic Moments from ELF partition (DEMEP)

Beyond traditional charge analysis: computation of local electrostatic properties on atomic and non atomic centers. Introduction of the  $M_1$  (dipole) and  $M_2$  (quadrupole) quantities

$$M_0(\Omega) = - \int_{\Omega} \rho(\mathbf{r}) d\tau$$

$$M_{1,x}(\Omega) = - \int_{\Omega} (x - X_c) \rho(\mathbf{r}) d\tau$$

$$M_{1,y}(\Omega) = - \int_{\Omega} (y - Y_c) \rho(\mathbf{r}) d\tau$$

$$M_{1,z}(\Omega) = - \int_{\Omega} (z - Z_c) \rho(\mathbf{r}) d\tau$$

$$M_{2,zz}(\Omega) = - \frac{1}{2} \int_{\Omega} (3(z - Z_c)^2 - r^2) \rho(\mathbf{r}) d\tau$$

$$M_{2,x^2-y^2}(\Omega) = - \frac{\sqrt{3}}{2} \int_{\Omega} [(x - X_c)^2 - (y - Y_c)^2] \rho(\mathbf{r}) d\tau$$

$$M_{2,xy}(\Omega) = - \sqrt{3} \int_{\Omega} (x - X_c)(y - Y_c) \rho(\mathbf{r}) d\tau$$

$$M_{2,xz}(\Omega) = - \sqrt{3} \int_{\Omega} (x - X_c)(z - Z_c) \rho(\mathbf{r}) d\tau$$

$$M_{2,yz}(\Omega) = - \sqrt{3} \int_{\Omega} (y - Y_c)(z - Z_c) \rho(\mathbf{r}) d\tau$$

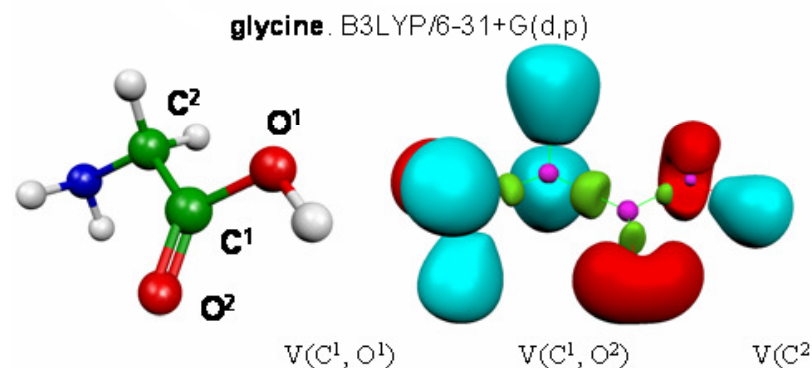
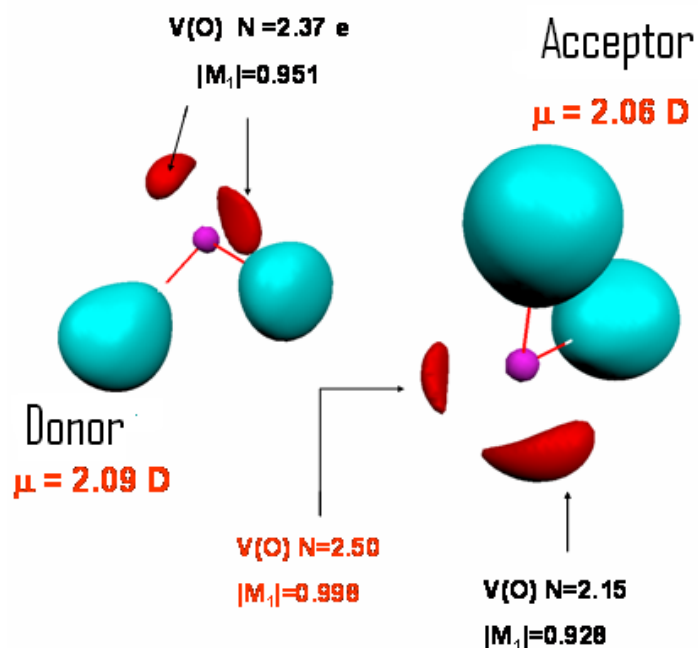
$$|\mathbf{M}(\Omega)| = \sqrt{\sum_i M_i(\Omega)^2}$$

# QM interpretative techniques

## Applications of DEMEP (1)

### Intermolecular Interactions

### Intramolecular Interactions.

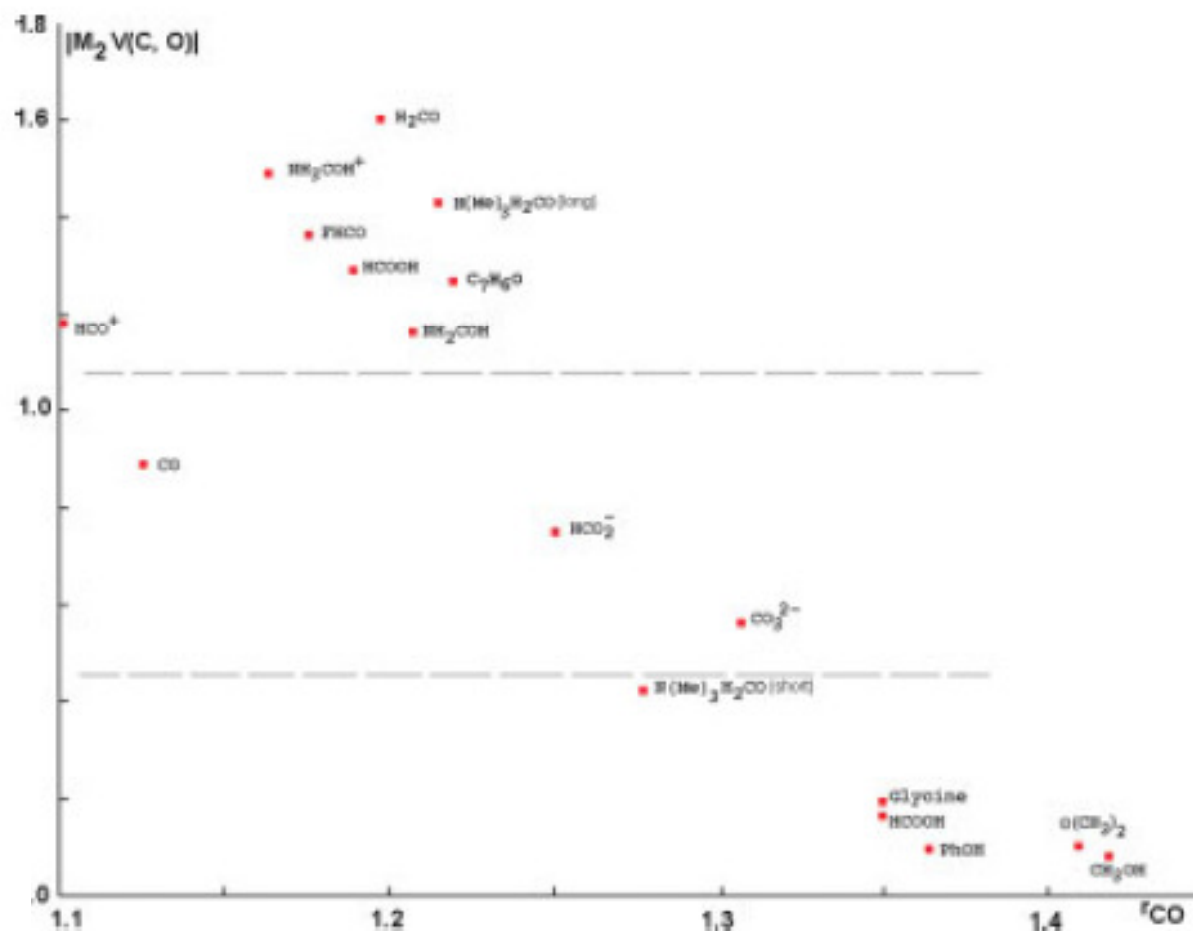
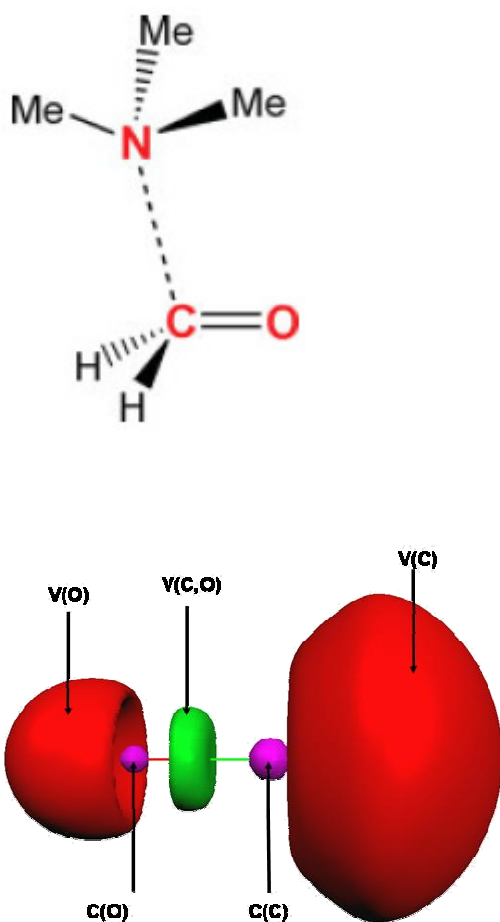


Acide Aminé	$V(C^1, O^1)$		$V(C^1, O^2)$		$V(C^2, N)$		$V(N)$		$ \mu $	$V(O^2)$	
	$ M_1 $	$ M_2 $	$ M_1 $	$ M_2 $	$ M_1 $	$ M_2 $	$ M_1 $	$ M_2 $		$ M_1 $	$ M_2 $
Glycine <sup>ab</sup>	0.255	1.249	0.052	0.257	0.183	0.166	0.952	0.125	11.6	3.308	2.446
Valine <sup>a</sup>	0.281	1.241	0.059	0.256	0.177	0.173	0.931	0.167	11.5	3.296	2.417
Tyrosine <sup>a</sup>	0.270	1.210	0.059	0.268	0.177	0.172	0.929	0.161	11,5	3.286	2.454



# QM interpretative techniques

## Applications of DEMEP (2)



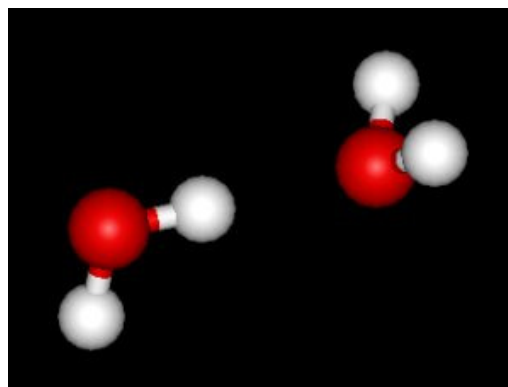
# QM interpretative techniques

## Energy Decomposition Analysis (EDA)

$$\Delta E = E_c + E_{\text{exch-rep}} + E_{\text{pol}} + E_{\text{CT}} + E_{\text{disp}}$$

$E_{\text{induction}}$

$\Delta E$ : Interaction Energy between Fragments.



# QM interpretative techniques

## EDA (1)

Perturbative techniques: high level QM (CCSD), non orthogonal orbitals.

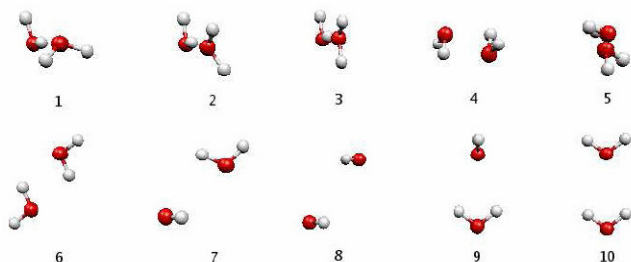
$$\Delta E = E_c + E_{\text{exch-rep}} + \underbrace{E_{\text{pol}} + E_{\text{CT}}}_{E_{\text{induction}}} + E_{\text{disp}}$$

Symmetry Adapted Perturbation Theory (SAPT): K. Szalewicz et al.

- Reference computation for first order contributions and dispersion(\*)
- Bad convergence of induction (métal etc...). Requires 3rd order corrections (tests on water dimer \*\*\*).

(\*)Piquemal, Cisneros, Gresh, Reinhardt, Darden, J Chem. Phys. 124, 104101, 2006.

(\*\*) P. Reinhardt., J-P. Piquemal, Int. J. Quant. Chem. Chem. 29, 1440, 2009



n	SCFluke(2)	SCFluke,r(2)	SCFluke(3)	SCFluke,r(3)	$\Delta E(\text{HF})$
1	2.43	2.63	3.71	3.91	3.53
2	2.25	2.43	3.32	3.50	3.16
3	2.31	2.48	3.34	3.51	3.18
4	2.11	2.24	2.84	2.97	2.62
5	2.02	2.14	2.62	2.73	2.44
6	2.06	2.16	2.60	2.71	2.44
7	1.80	1.85	2.23	2.29	2.04
8	0.66	0.68	1.79	1.81	0.71
9	1.72	1.78	2.25	2.31	2.00
10	1.39	1.43	1.69	1.73	1.53

# SCFluke(2) = E100pol + E100ex + E200Ind + E200ExInd

# SCFluke,r(2) = E100pol + E100ex + E200Ind,r + E200ExInd,r

# SCFluke(3) = E100pol + E100ex + E200Ind + E200ExInd + E300Ind(1+E200ExInd/E200Ind)

# SCFluke,r(3) = E100pol + E100ex + E200Ind,r + E200ExInd,r + E300Ind(1+E200ExInd/E200Ind)

Method/system	1	2	3	4	5	6	7	8	9	10
LMP2	-4.46	-3.95	-3.90	-3.75	-3.75	-3.49	-2.82	-1.05	-2.59	-1.84
F12-MP2	-4.87	-4.38	-4.35	-4.16	-3.94	-3.92	-3.10	-1.16	-2.89	-2.06
F12-CCSD	-4.62	-4.15	-4.13	-3.92	-3.69	-3.67	-3.02	-1.18	-2.83	-2.06
F12-CCSD(T)	-4.84	-4.33	-4.30	-4.13	-3.89	-3.86	-3.19	-1.28	-2.97	-2.15
SAPT (hybrid)	-4.85	-4.39	-4.37	-4.11	-3.91	-3.90	-3.17	-1.28	-3.02	-2.22
SAPT (Paris)	-5.23	-4.72	-4.70	-4.46	-4.21	-4.17	-3.41	-1.37	-3.34	-2.42

# QM interpretative techniques

## EDA (2)

$$\Delta E = E_c + E_{\text{exch-rep}} + E_{\text{pol}} + E_{\text{CT}}$$

- Variational approaches (orthogonal orbitals) :

Constrained Space Orbital Variations (CSOV) (Bagus et al. 1984)

- correlation through DFT (\*) or MCSCF (open-shell )
- correct induction (including exchange effects); charge transfer separation between donation-backdonation (\*) - charge transfer increases at the DFT level !!!!
- Application to transition and heavy metals via pseudopotentials (\*\*)
- Error bars on induction : improved charge transfer/polarization separation (\*\*\*)

**- Beyond 2 molecules: Localized DFT approach. (\*\*\*\*)**

(\*) Piquemal Marquez, Parisel, Giessner Prettre, J. Comput. Chem. 26, 1052, **2005**.

(\*\*) Gourlaouen Piquemal, Saue, Parisel, J. Comput. Chem., 27, 142, **2006**.

(\*\*\*) Piquemal, Bagus, **2009**.

(\*\*\*\*) P. Reinhardt., J-P. Piquemal, A. Savin, J. Chem. Theo. Comput. 4, 2020, **2008**

instance reference [19]). The scheme starts from a set of given (guess) orbitals and solves iteratively the Hartree-Fock equations via the steps:

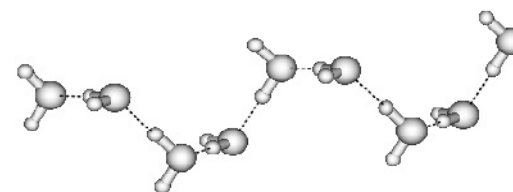
1. Symmetric (Löwdin) orthogonalization of the orbitals via  $\mathbf{S}^{-1/2}$
2. Construction of the Fock matrix
3. Calculation of the total energy
4. Construction and search for the lowest eigenvalue of an approximate Singles-CI matrix
5. Use in first order of the CI coefficients to correct the occupied and virtual molecular orbitals
6. Return to step 1

For constructing intrinsically localized DFT molecular orbitals, we take up an old idea,<sup>15</sup> further developed and published by Daudey<sup>16</sup> and still cited recently,<sup>2,17,18</sup> without the need to have delocalized canonical orbitals. The central point is that in Hartree-Fock theory a CI of singly excited Slater determinants lowers the total energy toward a set of Hartree-Fock orbitals, reaching convergence when satisfying Brillouin's theorem: interactions between occupied and virtual orbitals via the Fock matrix vanish. The singles-CI matrix is locally concentrated, as excitations on fragments with large spatial separation are much less important than those within close neighborhood. That permits the local correction of the monomer orbitals, disregarding the actual extent of the complete system. No canonical (completely delocalized) molecular orbitals are created or needed.

In step 3, a criterion of convergence may be introduced to terminate the iterations. Two other points should be mentioned: instead of taking the correct Singles-CI matrix, we may resort to a simpler one, omitting single bi-electronic integrals and using only Fock-matrix elements as:

$$\langle \Phi_i^a | \mathbf{H} | \Phi_i^b \rangle \approx F_{ab} \delta_{ij} - F_{ij} \delta_{ab} \quad (2)$$

functional	$E_{es}$	$E_{exch=rep}$	$E_{FC}$	$E_{OI+BSE}$	total interaction
HF (5D)	-8.26	6.88	-1.38	-2.16	-3.55
HF (6D)	-8.27	6.91	-1.36 (-1.37)	-2.18 (-2.19)	-3.55
B-LYP	-8.00	7.42	-0.58	-3.28	-3.86
B-LYP (6D)	-8.03	7.48	-0.55 (-0.61)	-3.30 (-3.26)	-3.86
PW91	-7.91	6.22	-1.69	-3.30	-4.99
PW91 (6D)	-7.93	6.27	-1.66 (-1.50)	-3.33 (-3.48)	-4.99
B3LYP	-8.05	6.67	-1.38	-2.97	-4.35
B3LYP (6D)	-8.07	6.71	-1.36 (-1.45)	-2.99 (-2.95)	-4.35 (4.40)



# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed  
(SIBFA)

Full QM based **Bottom up approach**

- Energetic scheme based on simplified QM.
- **Efficient reproduction of EDA schemes.**
- Anisotropy of the potential through **non atomic centers.**

Unified treatment of inter- and intra-molecular interactions.

# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)

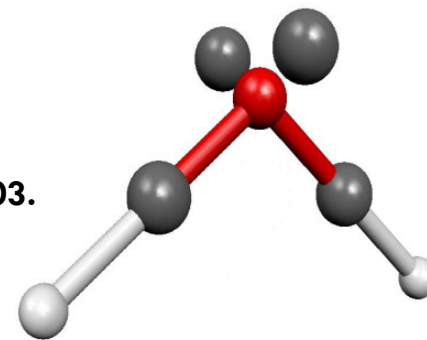
$$\Delta E = E_c + E_{\text{exch-rep}} + E_{\text{pol}} + E_{\text{CT}} + E_{\text{disp}}$$

Fully separable/anisotropic energy function.  
QM references: CSOV (or RVS), SAPT

- $E_c$ : Multipoles up to quadrupoles (atoms+bond midpoints)  
+ short-range electrostatics (penetration correction) (\*)
- Simplified LMO scheme for Exchange-Repulsion (Murrell, 1964)  
 $S^2/R + S^2/R^2$  (\*\*) (bond-bond+bond-LP+LP-LP)

(\*) Piquemal, Gresh, Giessner-Prettre, J. Phys. Chem A, Comput. Chem. 107, 10353, 2003.

(\*\*) Piquemal, Chevreau, Gresh, J. Chem. Theo. Comput., 3, 824, 2007.



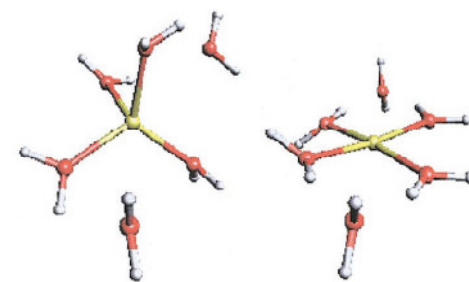


# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)

$$\Delta E = E_c + E_{\text{exch-rep}} + E_{\text{pol}} + E_{\text{CT}} + E_{\text{disp}}$$

- Polarization using ab initio polarizabilities on each LMO (Garner-Stevens)+ short range correction for exchange.
- **Quadrupolar polarizabilities on metals** (Cu(I), Cu(II)...)
- SIBFA includes **charge transfer** (Murrell et al., 1966)
- **Dispersion** ( $C_6, C_8, C_{10}$ ) + **exchange-dispersion**
- Effective Hamiltonian for open shell complexes: inclusion of **ligand field effects**



A  
SIBFA

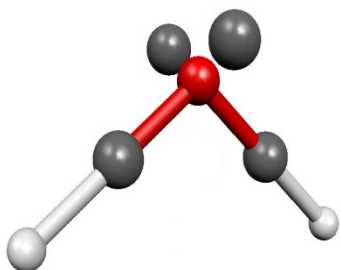
B  
SIBFA-LF

(\*) Gresh, Cisneros, Darden, Piquemal, J. Chem. Theo. Comput. 3, 1960, **2007**.

(\*\*) Piquemal, Gresh, Deeth, Giessner-Prettre, J. Comput. Chem. 24, 1963, **2003**.

# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)

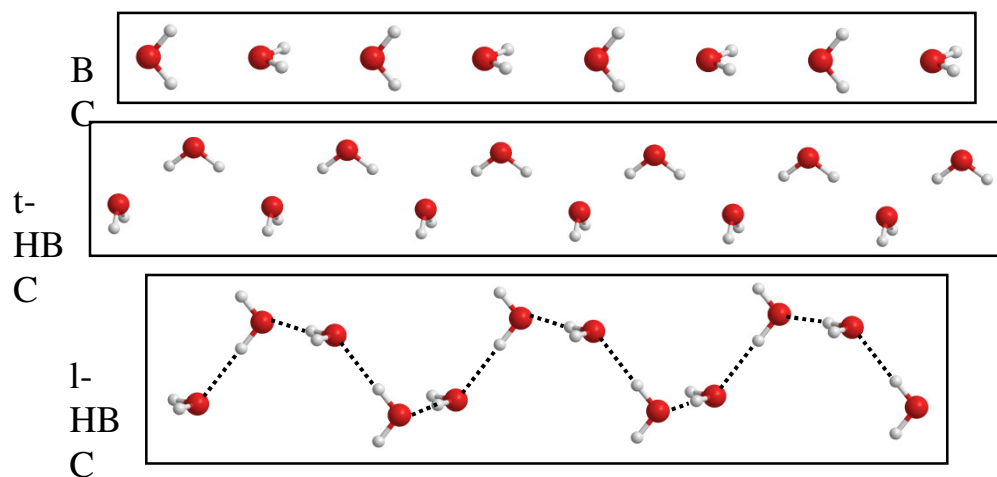


SIBFA vs. CPE (Chemical Potential Equalization) vs. RVS

Can SIBFA be a quantitative approach?

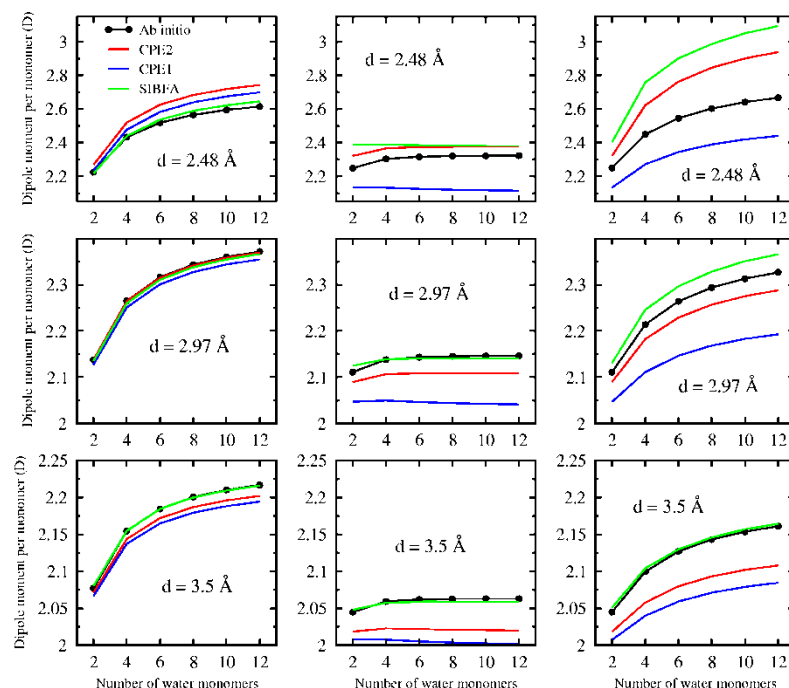
(To do that, it would need to reproduce both QM energies and QM average dipole moment/per molecule.)

→ It **does** thanks to its inclusion of polarizable lone pairs:  
importance of **electron localization**.



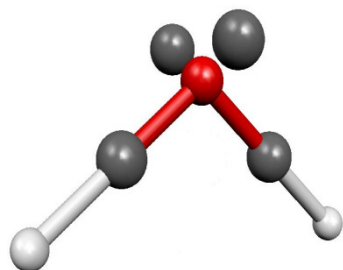
Piquemal et al., J. Phys. Chem. A 111, 8170, 2007.

Piquemal et al., J. Chem. Theo. Comput. 3, 824, 2007.



# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)



SIBFA vs. CPE (Chemical Potential Equalization)

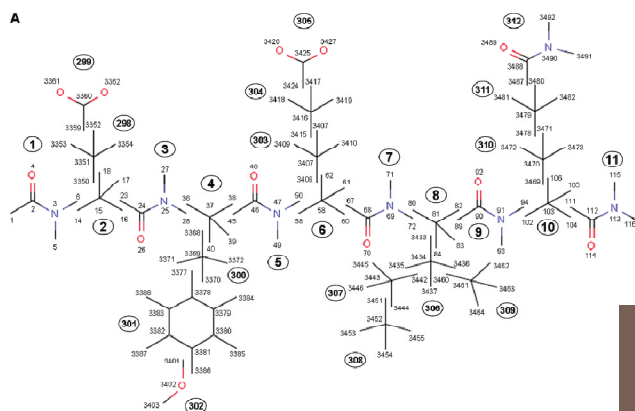
energies (kcal/mol)	$E_c$	$E_{\text{exch}}$	$E_1$	$E_{\text{pol}}$	$E_{\text{ct}}$	$E_2$	$\Delta E$
HBC SIBFA	-81.6	54.0	-27.6	-18.2 (-14.0)	-9.5	-27.7	-55.4
HBC RVS	-81.2	54.3	-26.8	-17.3 (-14.5)	-9.8	-24.3	-53.1
t-HBC SIBFA	-58.8	29.9	-28.8	-9.0 (-7.2)	-3.6	-12.6	-41.4
t-HBC RVS	-53.5	27.3	-26.3	-9.8 (-8.2)	-3.5		-39.2
I-HBC SIBFA	-60.9	54.0	-6.9	-3.9 (-3.6)	-7.5	-11.4	-18.3
I-HBC RVS	-60.5	55.1	-5.4	-5.5 (-4.7)	-7.7		-17.8

Piquemal et al., J. Phys. Chem. A 111, 8170, 2007.

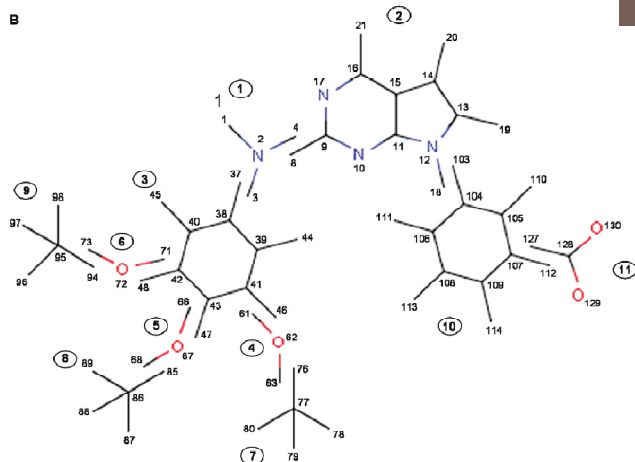
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# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)



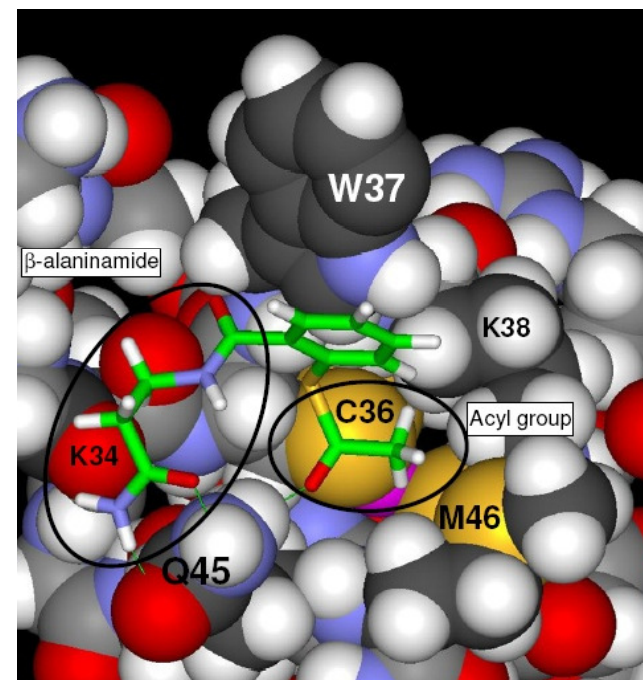
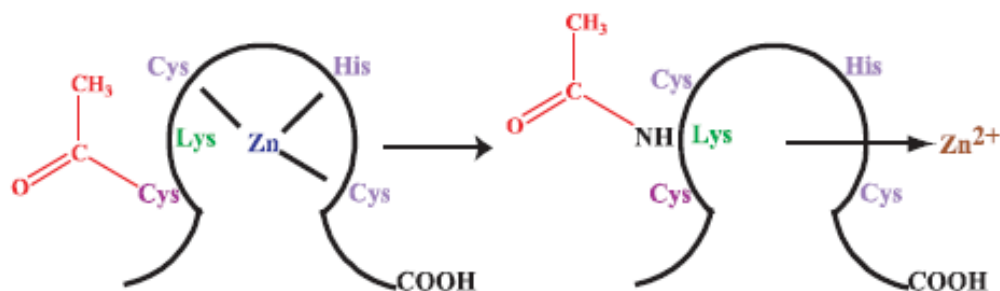
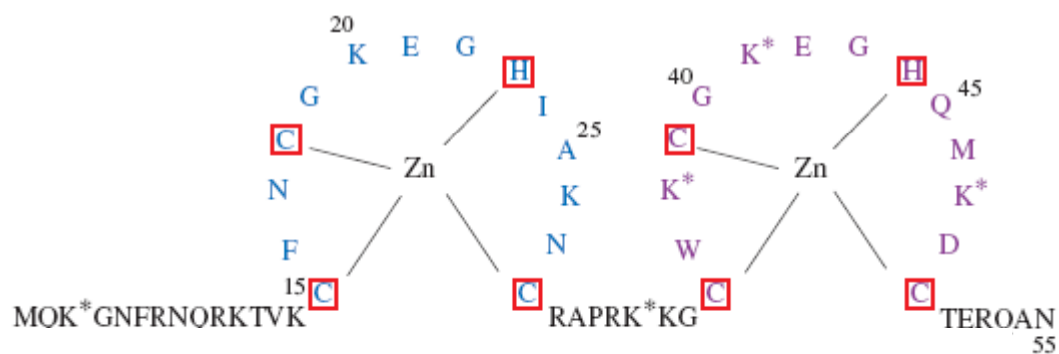
Macromolecules built upon using fragment libraries.  
Multipoles & polarizabilities redistribution.



# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)

Experimental collaboration: Ettore Appella  
NIH Immunology group (Bethesda, MA, USA)  
Phase II molecule (Thioester)

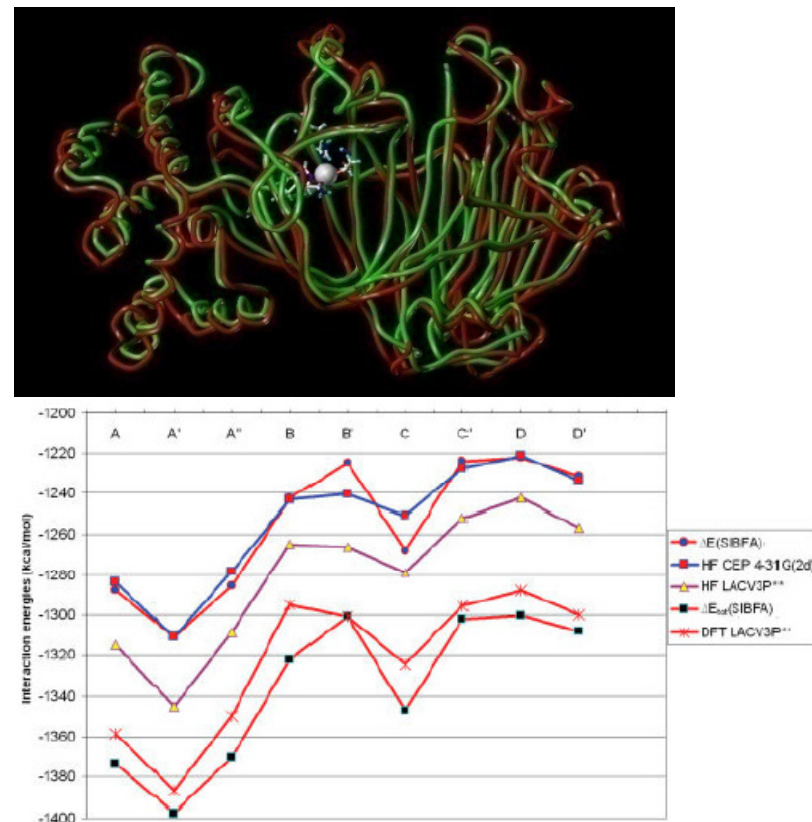
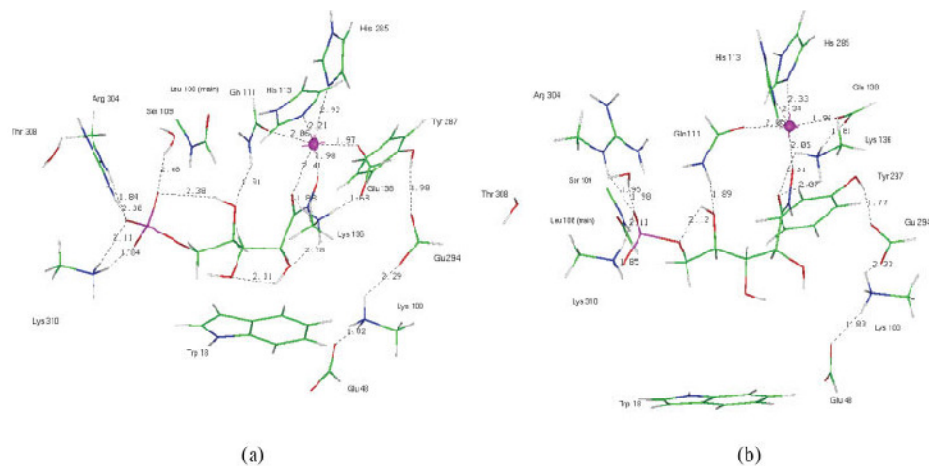


Miller Jenkins, Hara, Durell, Hayashi, Inman, Piquemal, Gresh, Appella, *J. Am. Chem. Soc.* 129, 11067, 2007.

# New generation force fields.

Sum of Interactions Between Fragments Ab initio computed (SIBFA)

Experimental collaboration:  
L. Salmon, ICCMO, Orsay (France).  
Phosphomannose Isomerase (PMI)  
164 residues.



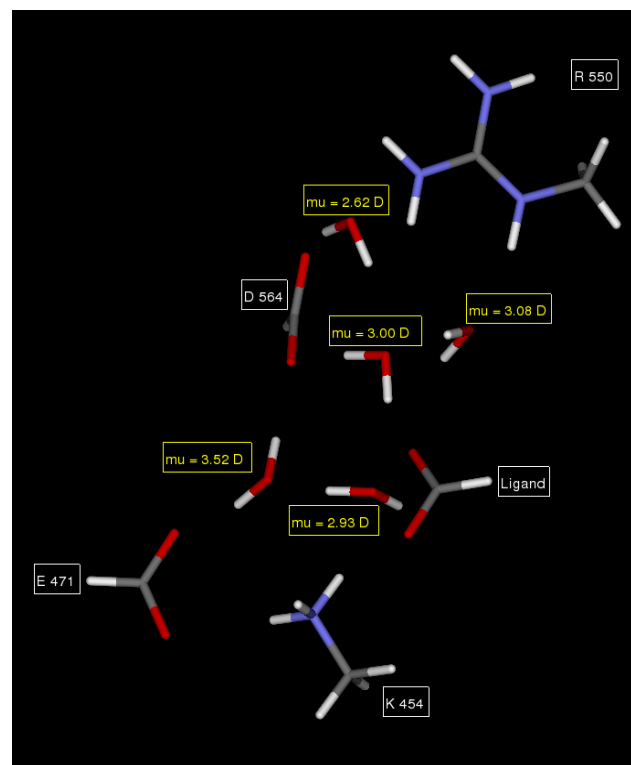
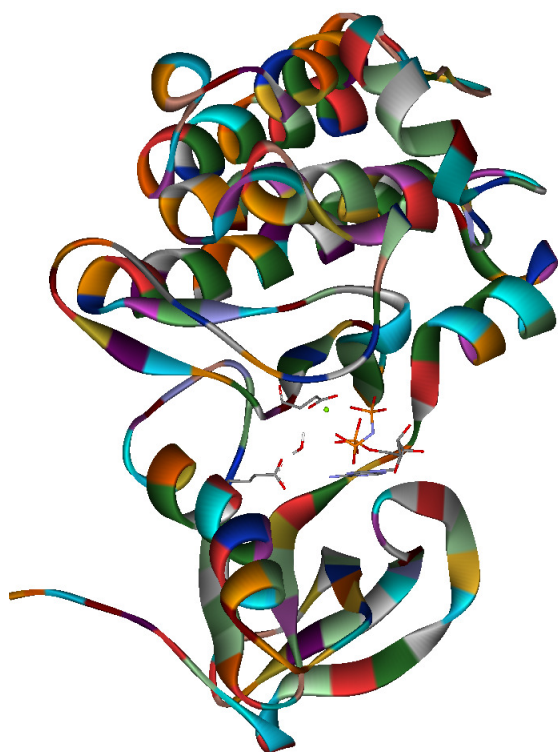
C. Roux, N. Gresh, L. Perera, J-P Piquemal and L. Salmon, *J. Comput. Chem.* 28, 938, **2007**.

J. Foret, B. de Courcy, N. Gresh, J-P. Piquemal, L. Salmon, **2009**, *Bioorg. Med. Chem.*, online.

# New generation force fields.

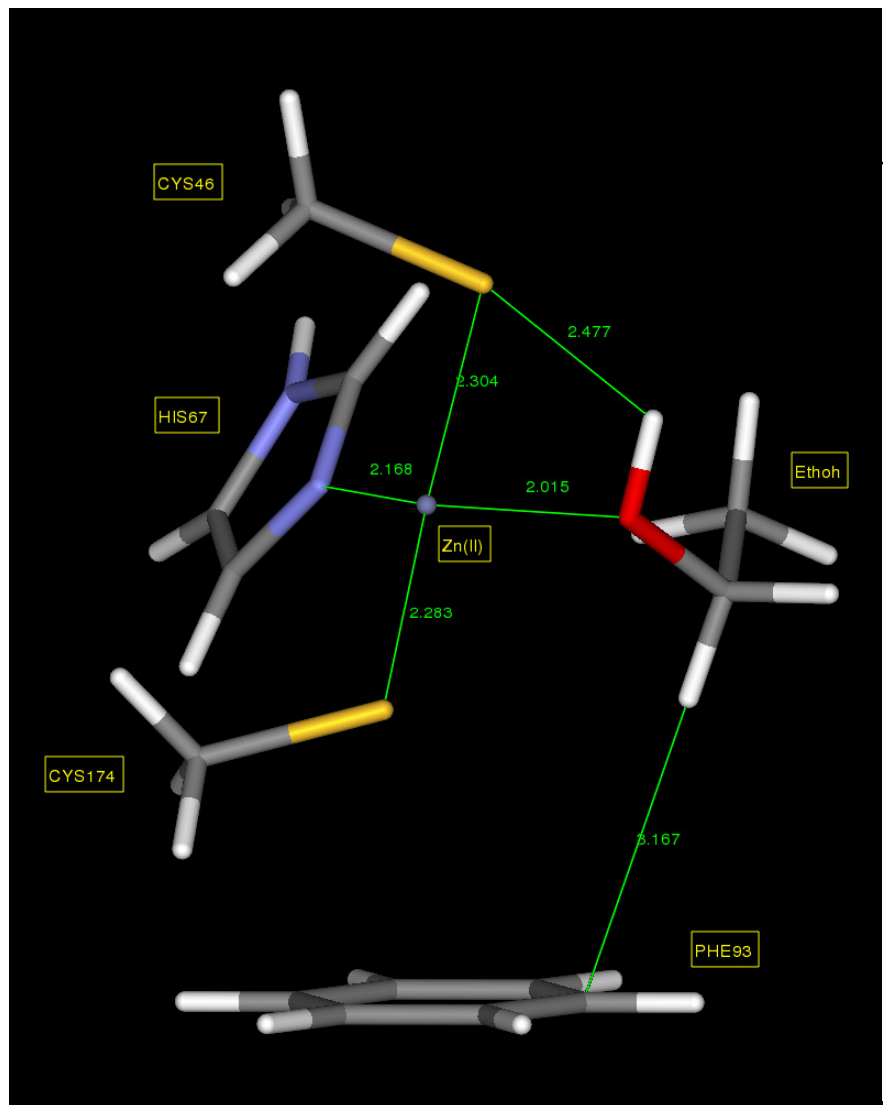
Sum of Interactions Between Fragments Ab initio computed (SIBFA)

Focal Adhesion Kinase (FAK)



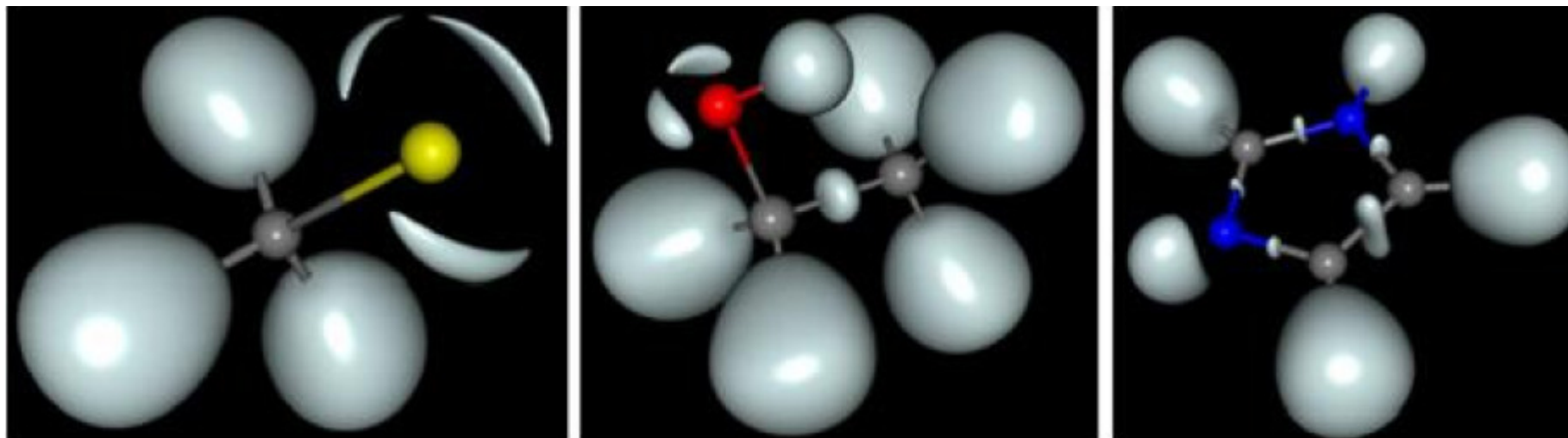
**EXPLICIT SOLVATION REQUIRED!!!**

De Courcy, Piquemal\*, Garbay, Gresh\*, to appear in *J. Am. Chem. Soc.*, **2009**



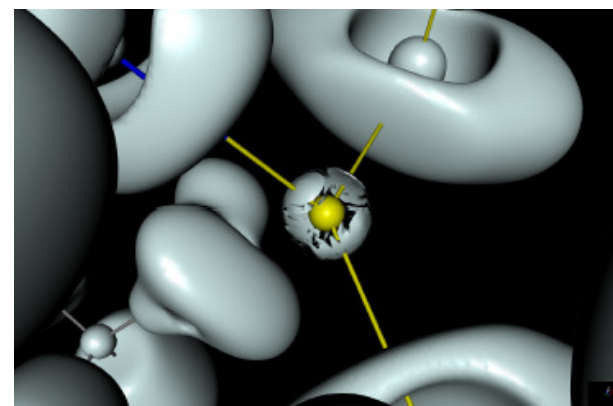
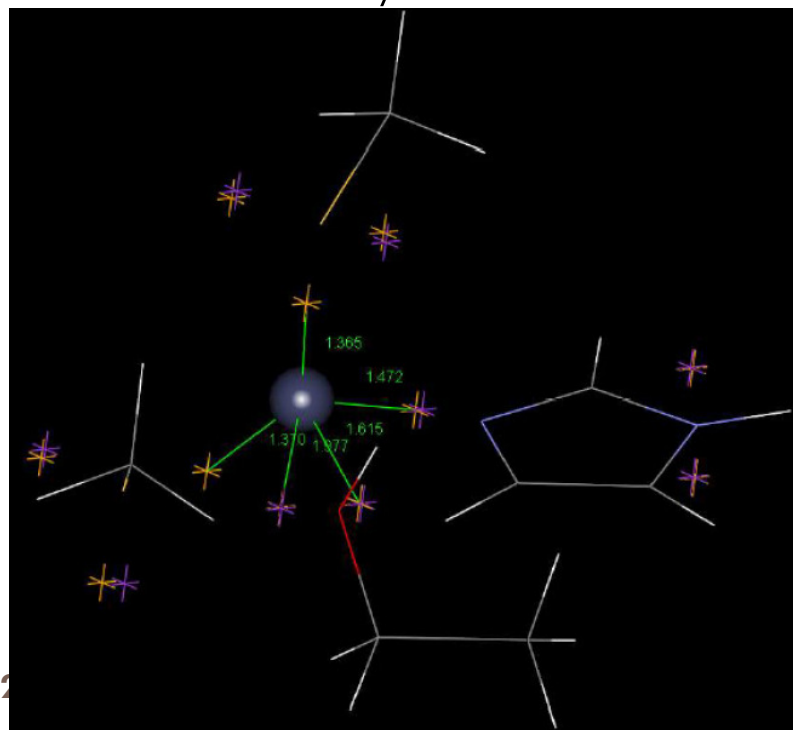
	Complex <i>a</i>		Complex <i>b</i>	
	Ab initio	SIBFA	Ab initio	SIBFA
$E_{\text{Coul.}} / E_{\text{MTP}}$	-661,6	-657,9	-664,4	-662,4
$E_{\text{exch.}} / E_{\text{rep}}$	177,4	168,9	180,2	173,5
$E_1$	-484,2	-489,0	-484,2	-488,9
$E_{\text{pol}}(\text{HF}) / E_{\text{pol}}$	-85,5	-94,4	-84,5	-93,0
$E_{\text{pol}}(\text{RVS}) / E_{\text{pol}^*}$	-113,2	-121,6	-111,8	-120,0
$E_{\text{ct}}(\text{RVS})$	-48,5		-48,6	
BSSE	-5,1		-5,6	
$E_{\text{ct}^*} / E_{\text{ct}}$	-43,4	-41,3	-43,0	-41,3
$E_2$	-156,6	-162,9	-154,8	-161,3
$\Delta E$	<b>-618,2</b>	<b>-624,7</b>	<b>-617,3</b>	<b>-623,2</b>
$\delta E(\text{MP2}) / E_{\text{disp}}$	-48,4	-65,9	-57,8	-70,5
$\Delta E_{\text{tot}}$	<b>-666,6</b>	<b>-690,6</b>	<b>-675,1</b>	<b>-693,7</b>



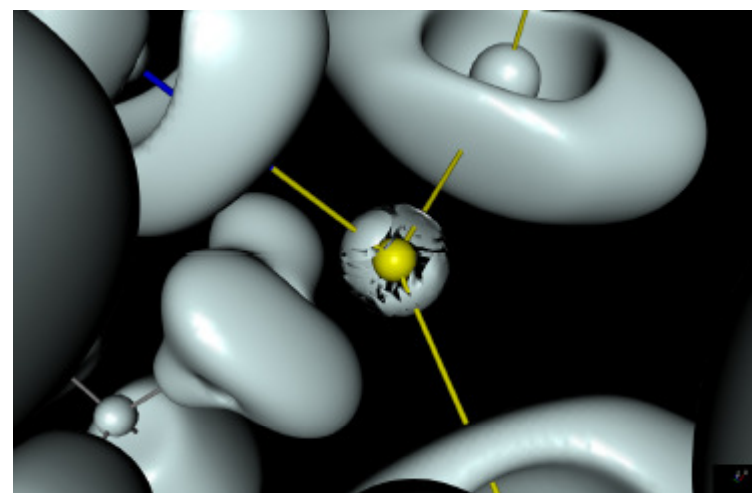
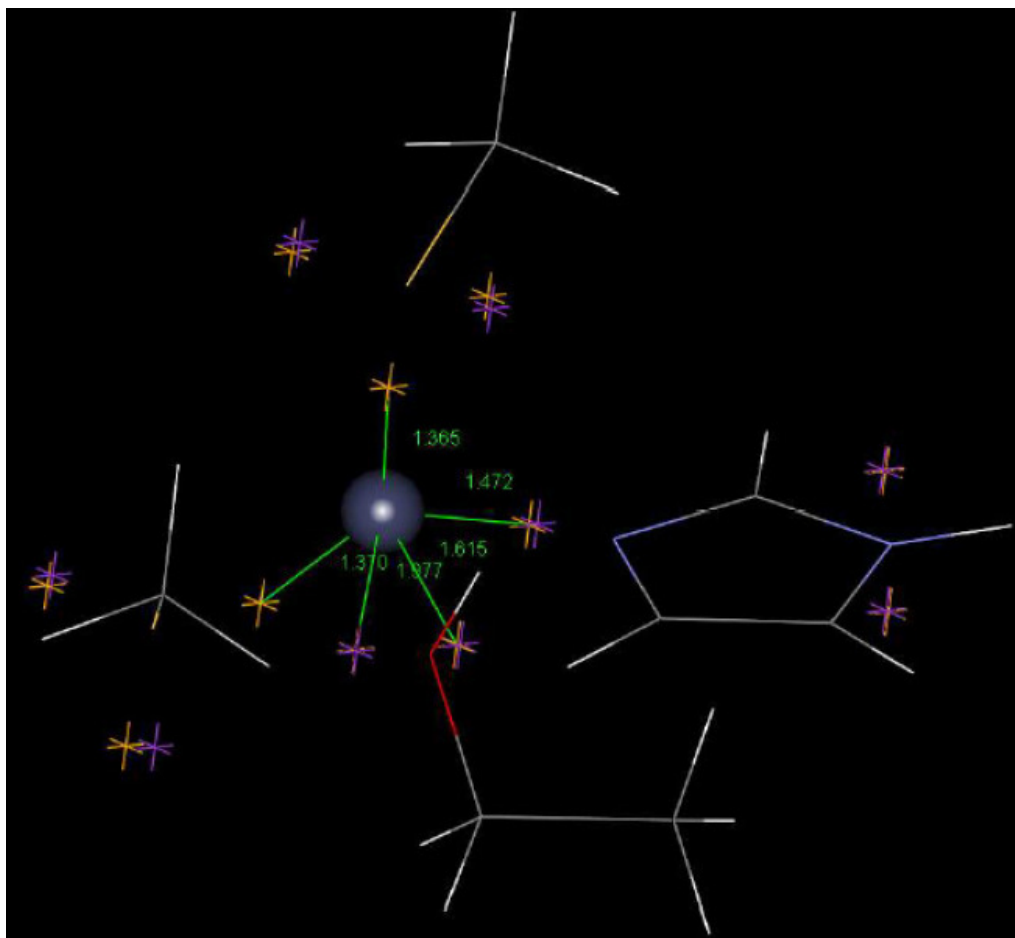


De Courcy, Gresh, Piquemal\*, Interdiscip Sci Comput Life Sci (2009) 1, 55

DOI: 10.1007/s12539-008-0027-0



	$E_1 / \Delta E$	$E_2 / \Delta E$
Cy <sup>-</sup> 46/ Zn(II)	66,8%	33,2%
Imh / Zn(II)	44,1%	55,9%
Cy <sup>-</sup> 174/ Zn(II)	67,1%	32,9%
EtOH / Zn(II)	39,7%	60,3%



	$E_1 / \Delta E$	$E_2 / \Delta E$
Cy <sup>-</sup> 46/ Zn(II)	66,8%	33,2%
Imh / Zn(II)	44,1%	55,9%
Cy <sup>-</sup> 174/ Zn(II)	67,1%	32,9%
EtOH / Zn(II)	39,7%	60,3%

# Next generation force field.

- Beyond multipoles: Gaussian Electrostatics Model.(\*).



## Density Fitting

A fundamental calculation of the energy surface for the system of three hydrogen atoms  
**S F Boys and I Shavitt**, University of Wisconsin, Report WIS-AF-13 (1959) NTIS AD212985

“In this method, each two-center charge distribution is expanded by least-squares as a sum of single-centre charge distributions centered on various points on the line passing through the two centres. . .”

$$\tilde{\rho} = \sum_{k=1}^N x_k k(r) \approx \rho = \sum_{\mu\nu} P_{\mu\nu} \phi_{\mu}(r) \phi_{\nu}^*(r)$$

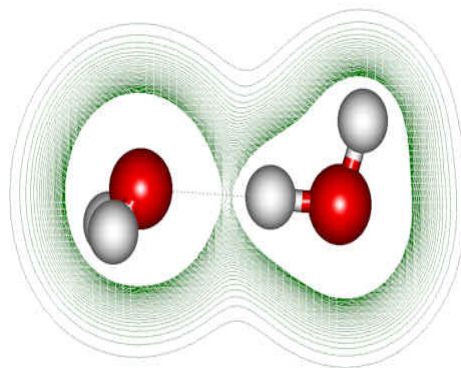
$$E_{\text{coulomb}} = \frac{Z_A Z_B}{r_{AB}} - \int \frac{Z_A \tilde{\rho}^B(\mathbf{r}_B)}{r_{AB}} dr - \int \frac{Z_B \tilde{\rho}^A(\mathbf{r}_A)}{r_{AB}} dr + \int \frac{\tilde{\rho}^A(\mathbf{r}_A) \tilde{\rho}^B(\mathbf{r}_B)}{r_{AB}} dr$$

(\*). Cisneros, Piquemal, Darden, J. Chem. Phys. 123, 055109, 2005

Piquemal, Marquez, Parisel, Giessner-Prettre, J. Comput. Chem., 26, 1052, 2005..

# Next generation force field.

GEM :what if MM=QM??



$$\Delta E = E_c + E_{\text{éch-rép}} + E_{\text{pol}} + E_{\text{TC}} + E_{\text{disp}}$$

Frozen density of a rigid fragment  
Hermite gaussians  
(recursions using PME –fourier space  $n \log(n)$ )

Piquemal, Cisneros, Reinhardt, Gresh, Darden, J. Chem. Phys. 124, 104101, **2006**

Cisneros, Piquemal, Darden, J. Chem. Phys. 125, 184101, **2006**

Cisneros, Piquemal, Darden, J. Phys. Chem. B. 110, 13682, **2006**

Cisneros, Elking, Piquemal, Darden, J. Phys. Chem. A 111, 12049, **2007**

Gresh, Cisneros, Darden, Piquemal, J. Chem. Theo. Comp. 3, 1960, **2007**

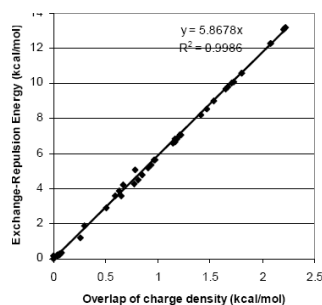
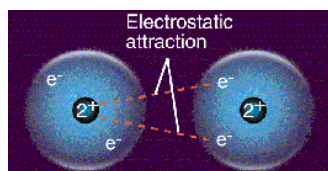
# Next generation force field.

GEM :what if MM=QM??

Auxiliary Coefficients

Integrales

$E_{\text{coul}} + E_{\text{exch-rep}}$



$$E_{\text{coulomb}} = \frac{Z_A Z_B}{r_{AB}} - \int \frac{Z_A \tilde{\rho}^B(r_B)}{r_{AB}} dr - \int \frac{Z_B \tilde{\rho}^A(r_A)}{r_{AB}} dr + \int \frac{\tilde{\rho}^A(r_A) \tilde{\rho}^B(r_B)}{r_{AB}} dr$$

$$S_{\rho} = \int \rho_a(r) \rho_b(r) dr \approx \int \tilde{\rho}_a(r) \tilde{\rho}_b(r) dr$$

$$E_{\text{exch/rep}} \approx KSP \quad (\text{Wheatley and Price})$$

Electrostatic Potentials & Field from frozen densities

Polarization

+

Charge transfer

$$-0.5 * \sum_j^{xyz} \Delta\mu(i) * (\gamma E_0(j))$$

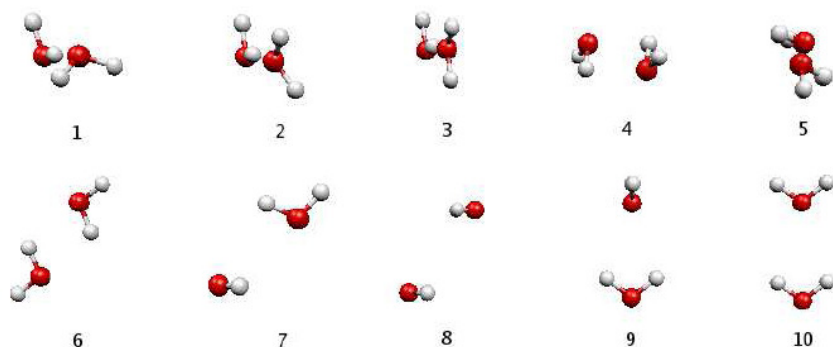
(Garmer et Stevens)

$$E_{ct} = -2 \sum_{\alpha} \sum_{\beta^*} \frac{(I_{\alpha\beta^*})^2}{\Delta E_{\alpha\beta^*}}$$

(Gresh, Claverie et Pullman)

# Next generation force field.

GEM :what if MM=QM?? Water clusters: aug-cc-pVTZ

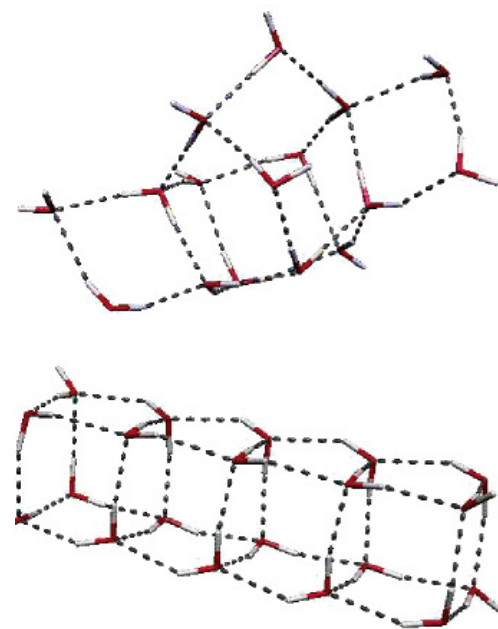


Chemical precision?

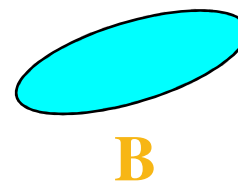
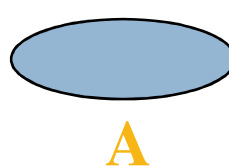
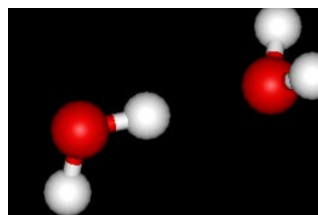
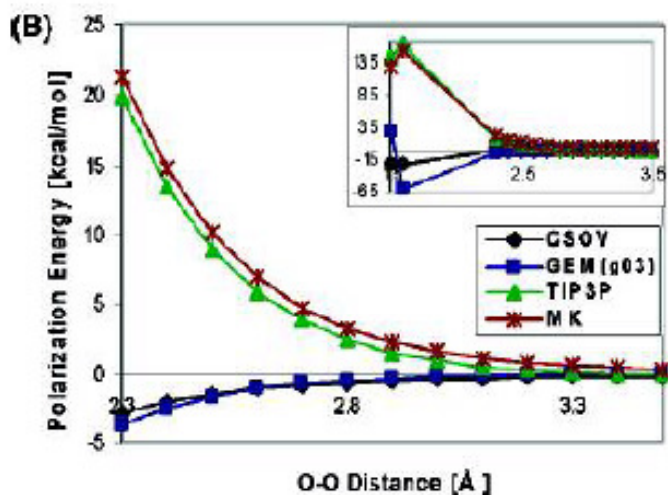
Average errors on total interaction energies:  
< 0.1 kcal/mol .

Errors < 0.1 kcal on **each** of the components.

n	$E_{\text{coulomb}}$ GEM-0	$E_{\text{coulomb}}$ CSOV	$E_{\text{coulomb}}$ GEM-0 (BD)	$E_{\text{exch-rep}}$ GEM-0	$E_{\text{exch-rep}}$ CSOV
16	-186.842	-186.379 (-186.842)	-184.798	164.946	166.539 (166.549)
20	-309.380	-307.199 (-309.204)	-305.838	292.25	292.164 (292.190)
64	-449.519	-446.122 (-454.077)	-443.537	336.48	-



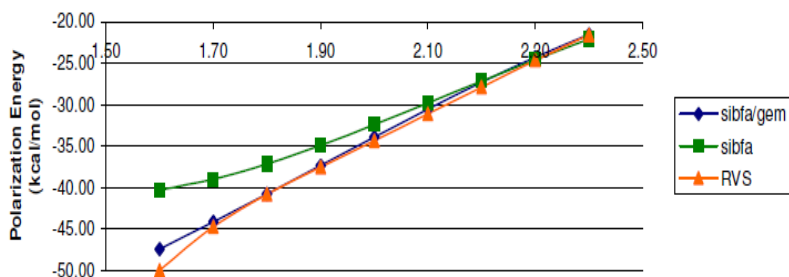
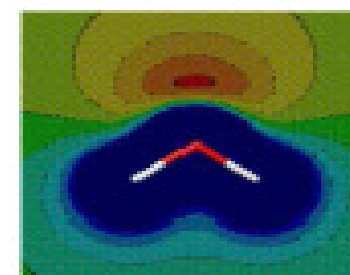
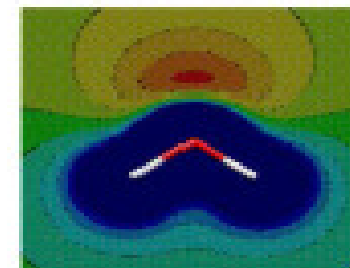
# The Gaussian Electrostatic Model (GEM): QM/MM and MM'/MM



Molecule A (QM)

Molecule B (GEM)

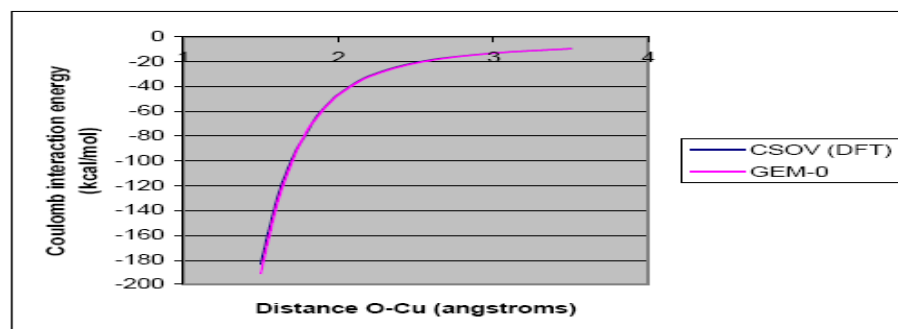
$$H_A = H_A^\circ + V_{AB}$$



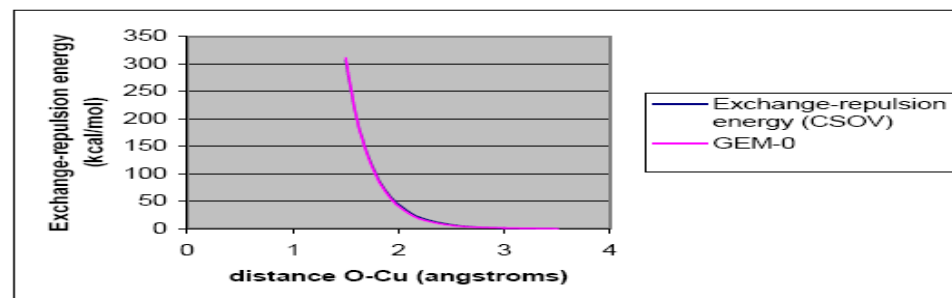
**Inclusion of scalar relativity via pseudopotentials: Hg(II) potential including quadrupolar polarisation & back-donation.**

# Cu(I)-H<sub>2</sub>O Complex

32



**Figure 2 :** Intermolecular coulomb energy for the water-Cu(I) complex at various distances for GEM-0 and CSOV at the B3LYP/aug-cc-PVTZ level for the water and B3LYP/6-31G\* for the cation. The water is immobile at the position of the molecule A of the dimer 1.



**Figure 3 :** Intermolecular exchange-repulsion energy for the water-Cu(I) complex at various distances for GEM-0 and CSOV at the B3LYP/aug-cc-PVTZ level for the water and B3LYP/6-31G\* for the cation. The water is immobile at the position of the molecule A of the dimer 1. The K parameter is 2.8942 ( $R^2=0.9991$ ).



# Continuous Electrostatic Model

$$\tilde{\rho} = \sum_{k=1}^N x_k k(r) \approx \rho = \sum_{\mu\nu} P_{\mu\nu} \phi_{\mu}(r) \phi_{\nu}^*(r)$$

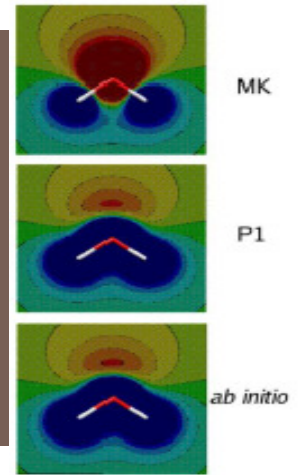
Efficient recursion for cartesian multipoles :  
Cipriani et Silvi (1982) (L<sup>4</sup>)

**Hermites (gaussian derivative) : link to cartesian multipole.**  
Challacombe, Schwegler, Amlof (1995) (L<sup>2</sup>)

$$\Lambda_{tuv}(\mathbf{r}, \alpha, \mathbf{P}) = \left( \frac{\partial}{\partial P_x} \right)^t \left( \frac{\partial}{\partial P_y} \right)^u \left( \frac{\partial}{\partial P_z} \right)^v \exp(-\alpha r_{\mathbf{P}}^2)$$

**Fast convergence of the multipole serie:**  
**Quadrupole is enough!**

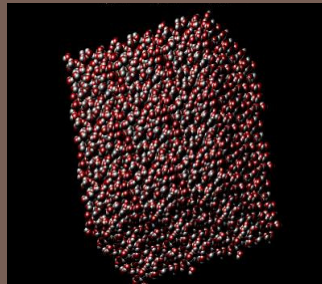
+ efficient generalized McMurchie-Davidson recursion



# How fast it is?

34

- It depends on the basis sets and can be improved by a dual density/ESP grid fitting neglecting the core.



4096 water molecules, single CPU 3.6 Ghz

A2 basis sets fitted on B3LYP 6-31G\*

2.29 s

To be compared to 0.2 s for AMBER (TIP<sub>3</sub>P)

It could be less using spherical harmonics/Slater-type expressions  
(JCTC, 2009, in press)



## Rethinking Amoeba: metals and PBC

(Atomic Multipole Optimized Energetics for Biomolecular Applications)  
Ren et Ponder, 2003

A « reasonable » first step towards precision (protein ready!)

Piquemal, Perera, Cisneros, Ren, Pedersen, Darden, J. Chem. Phys., **2006**, 125, 054511

# AMOEBA: no universal Thole damping!!!

Atomic Multipoles (Stone)

Polarization: Applequist/Tholé

van der Waals: Halgren's function

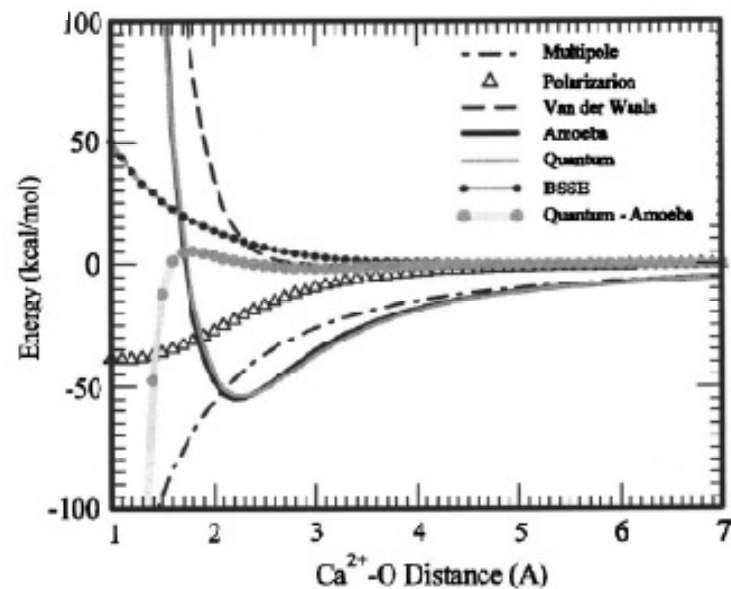
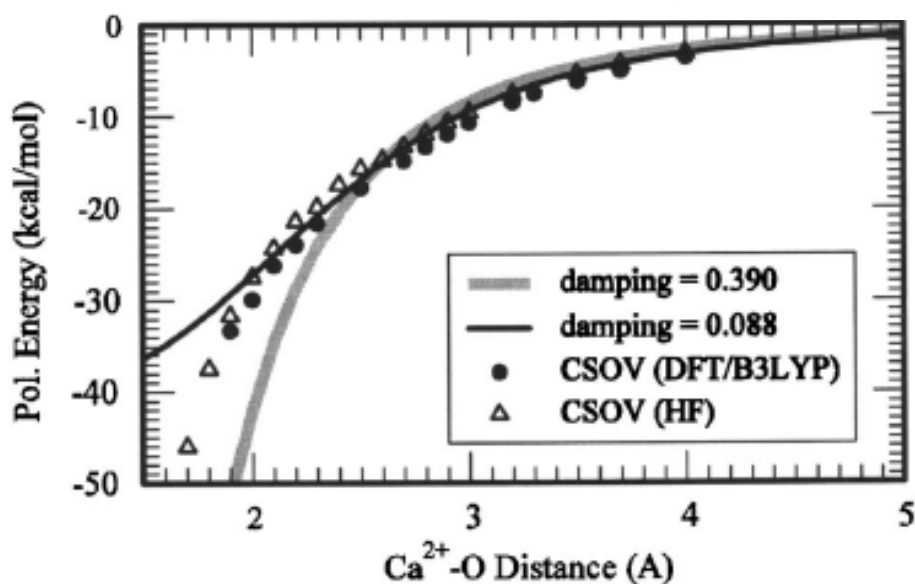
$$E_{\text{Halgren}}(14-7) = E_{\text{exc-rep}} + E_{\text{disp}} + E_{\text{TC}}$$

$$= E_{\text{Halgren}}(\text{direct}) + E_{\text{Halgren}}(\text{recip.})$$

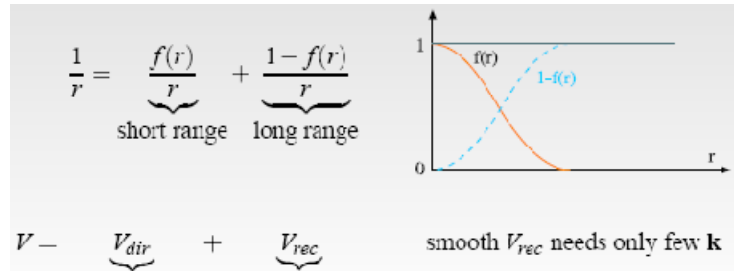
$$E_{\text{Halgren}} = \Delta E_{\text{tot}}(\text{MP2BSSE corrected})$$

$$- E_{\text{Coulomb}}(\text{amoeba multipoles})$$

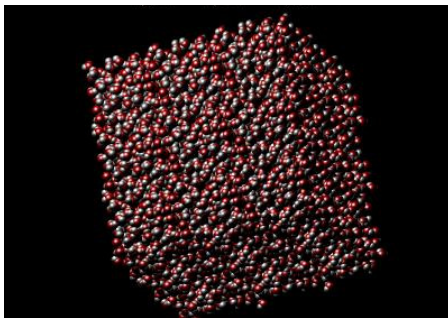
$$- E_{\text{polarization}}(\text{amoeba}).$$



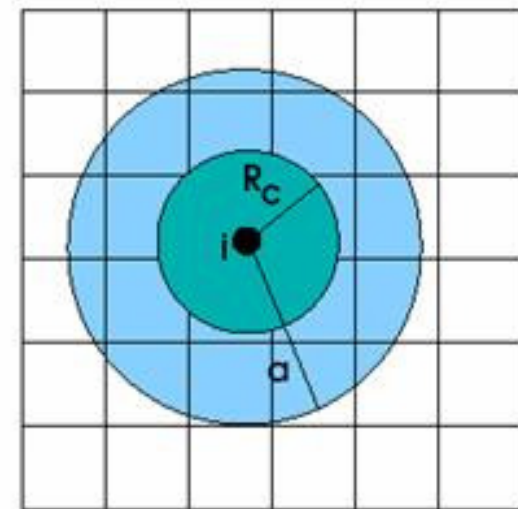
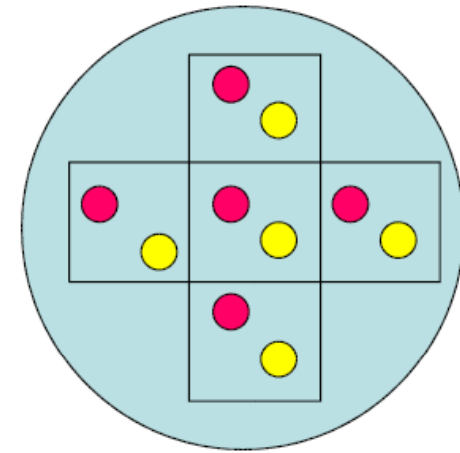
# PBC for Amoeba

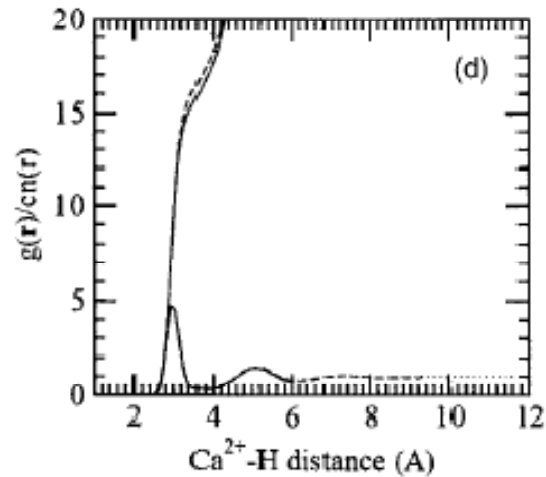
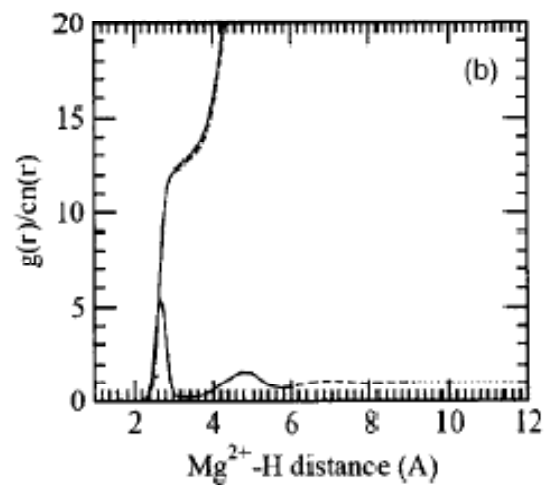
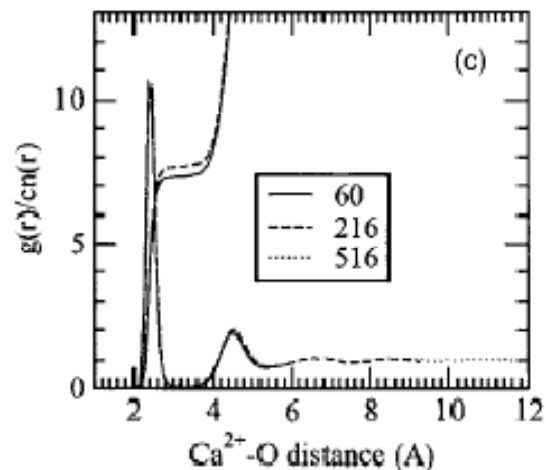
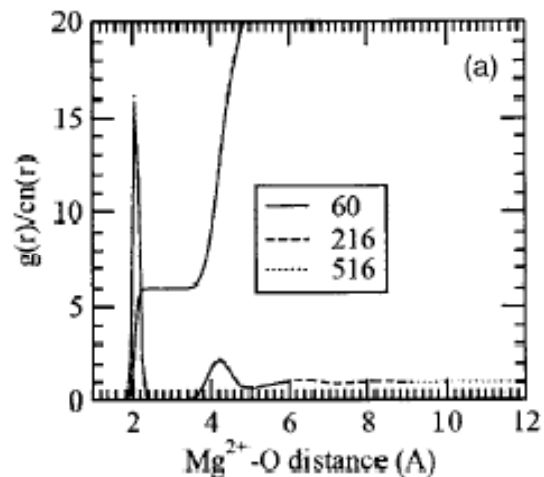
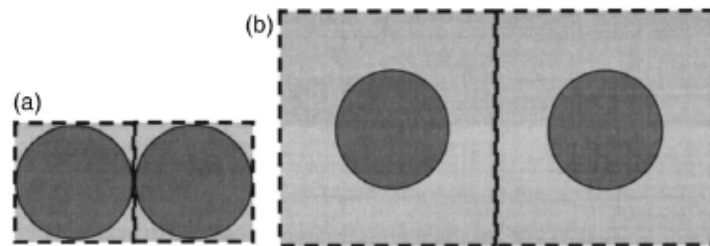


Particle Mesh Ewald  
 $N^2 \rightarrow N \log(N)$



Amber:  
 van der Waals are in reciprocal sum too!!!





## MD

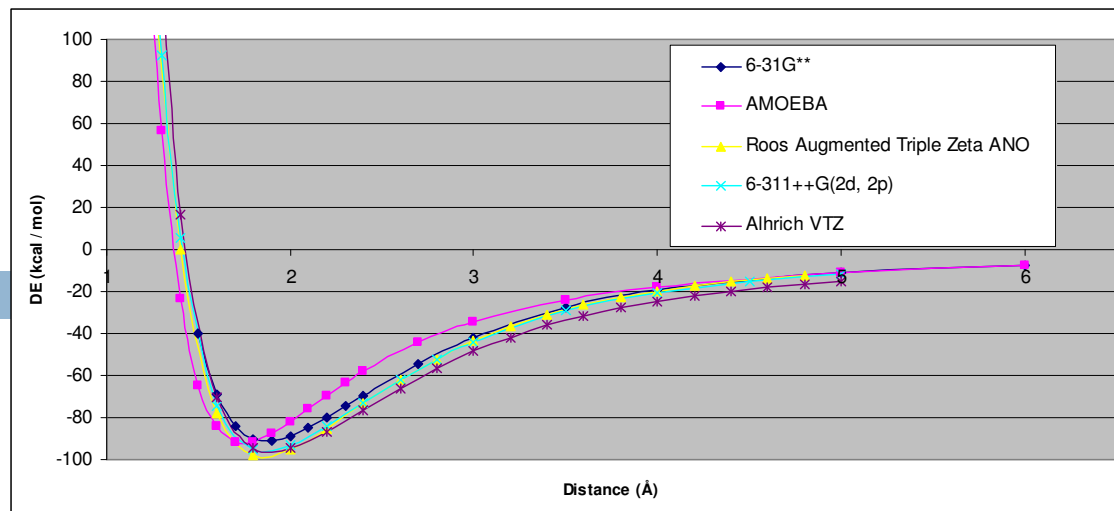
- Intégration step: 1 fs
- Successive over relaxation (SOR)
- Direct sum: 6,7 Å cutoff (8 Å vdw)

CCI :  
10-12 time Amber  
but robust MD, Proteines ready

1 ns = 34 H (216 H<sub>2</sub>O)  
= 94 H (512 H<sub>2</sub>O)

1.8 Ghz Opteron AMD

	$M = \text{Ca}^{2+}$			$M = \text{Mg}^{2+}$
Water coordination				
Experimental <sup>a,b</sup>	7.0–8.0			6.0
Car-Parrinello <sup>c</sup>	6.2–7, 7.2, 8–9			6.0
This work	7.7 (216 and 516 waters), 7.2 (60 waters)			6.0
First $M(\text{II})-\text{O}$ peak (Å)				
Experimental <sup>d</sup>	2.41–2.44; 2.437; 2.46			2.09
Car-Parrinello <sup>e</sup>	2.43–2.44			2.13
This work	2.42–2.46			2.08
First $M(\text{II})-\text{H}$ peak (Å)				
Experimental <sup>a</sup>	2.97–3.07			...
Car-Parrinello <sup>f</sup>	3.03–3.07			...
This work	3.01–3.05			2.63–2.73
Average first shell water tilt angle				
Experimental <sup>g</sup>	34.0–38.0; 39.7			...
Car-Parrinello <sup>f</sup>	40.1			...
This work	40.2–40.7			39.7–40.6
First shell water residence time (ps) for $n$ water molecules	$N$	$t^* = 0$ ps	$t^* = 2$ ps	No exchange of water during the simulation time
	60	131	154	
	216	158	182	
	516	160	203	
Percentage coordination with $n$ water molecules	$n=60$	$n=216$	$n=516$	
6	0.84	0	0.2	100.0
7	46.16	33.16	30.28	0.0
8	52.72	66.32	69.16	0.0
9	0.28	0.52	0.36	0.0



Zn (II):

Free energy of solvation

Zn <sup>2+</sup>	-457.01	-467.26 (Marcus)
Mg <sup>2+</sup>	-431.1 (2.9)	-435.4 (Schmid)
Ca <sup>2+</sup>	-354.9 (1.7)	-357.2 (Schmid)

Piquemal, Wu, Reinhardt, Chaudret, Ren, 2009, to appear, JPCB.

Incoming developments: Charge transfer/IR spectroscopy.

(Chaudret, Gresh, Darden, Piquemal)



Design of next generation polarizable force fields from ab initio computations: beyond point charges.

Cisneros, Darden, Gresh, Reinhardt, Parisel, Pilmé and Piquemal  
in *Multi-scale Quantum Models for Biocatalysis: Modern Techniques and Applications*, for the Book Series: *Challenges and Advances in Computational Chemistry and Physics* , ed. D. M. York and T.-S. Lee, **2009**, 137-172, Springer Verlag

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Thomas Darden (Open Eyes Inc.)

Pengyu Ren (University of Texas, Austin)

Dennis Salahub (Calgary)

Michel Mons, Jean-Pierre Dognon (French Atomic Commission, CEA)

