



Center for
Molecular Modeling

Faster Potential Energy Surfaces: The *art* of Making Force Fields

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IPAM Workshop IV: Physical Frameworks for Sampling Chemical Compound Space

May 2011, Los Angeles



Overview

Introduction

Exploring Chemical Compounds
Force Field 101
Making Force Fields

Atoms in Molecules

Partitioning of Electron Density
Iterative Hirshfeld
ESP Fitted charges
Benchmarks

The Hydrogen Bond

AIM Application to FF development
Charge Transfer Matters!

Split Charge Equilibration

Partitioning of Polarization
Split Charges
Benchmarks

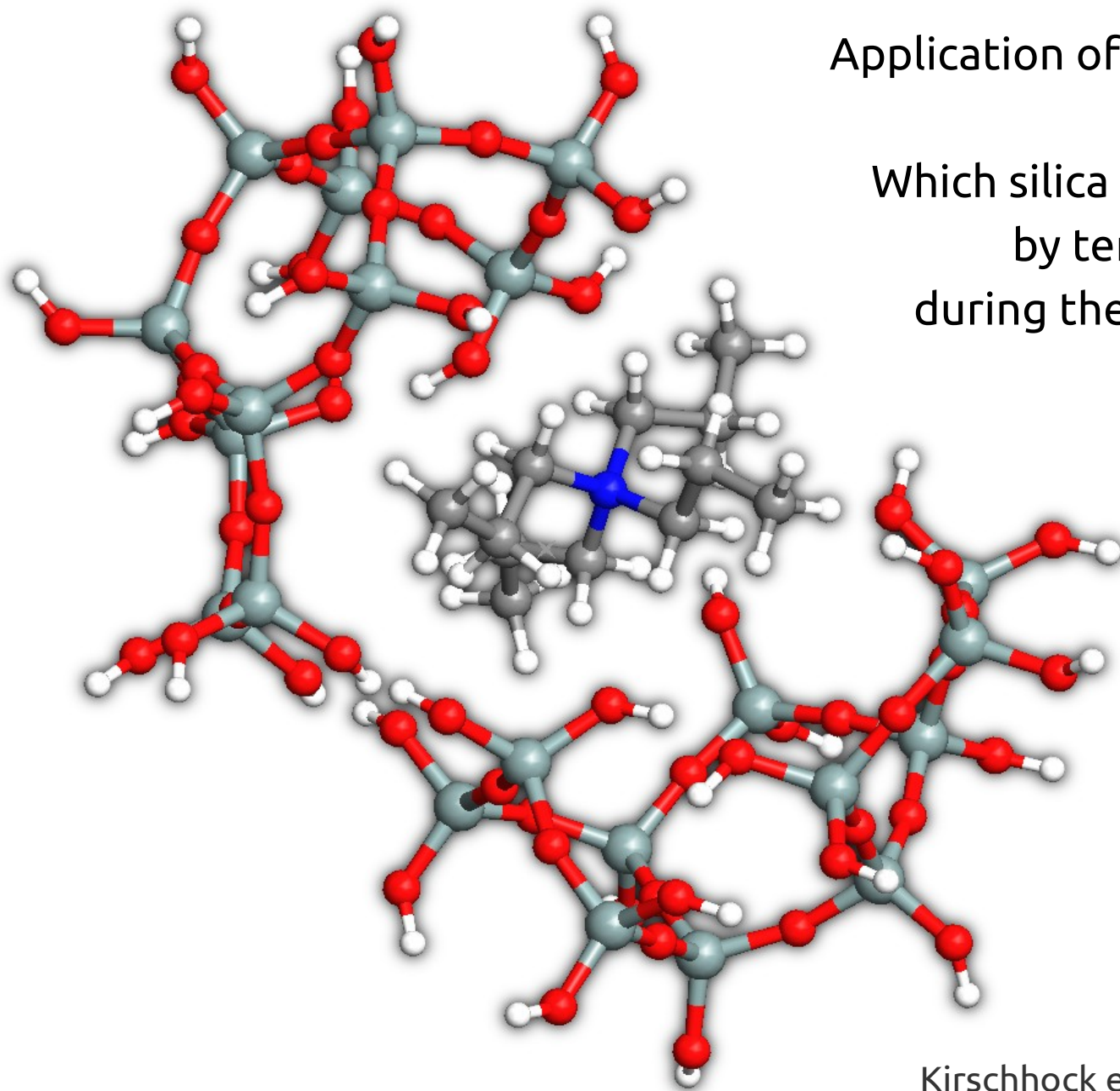
Harmonic Models

Force Constants & Rest Values

Conclusion

Application of interest: Zeolite synthesis

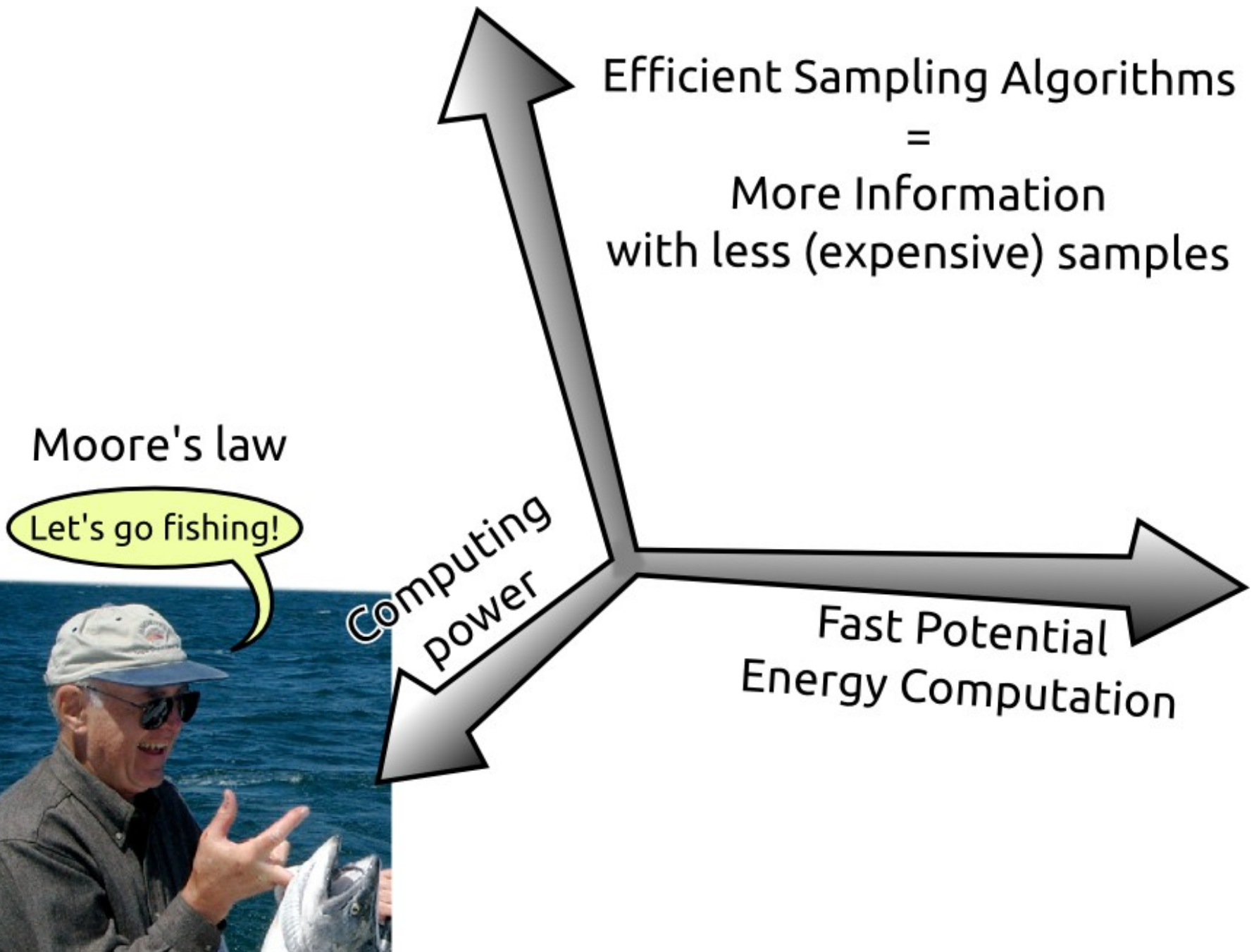
Which silica oligomers are promoted
by template molecules
during the prenucleation phase?



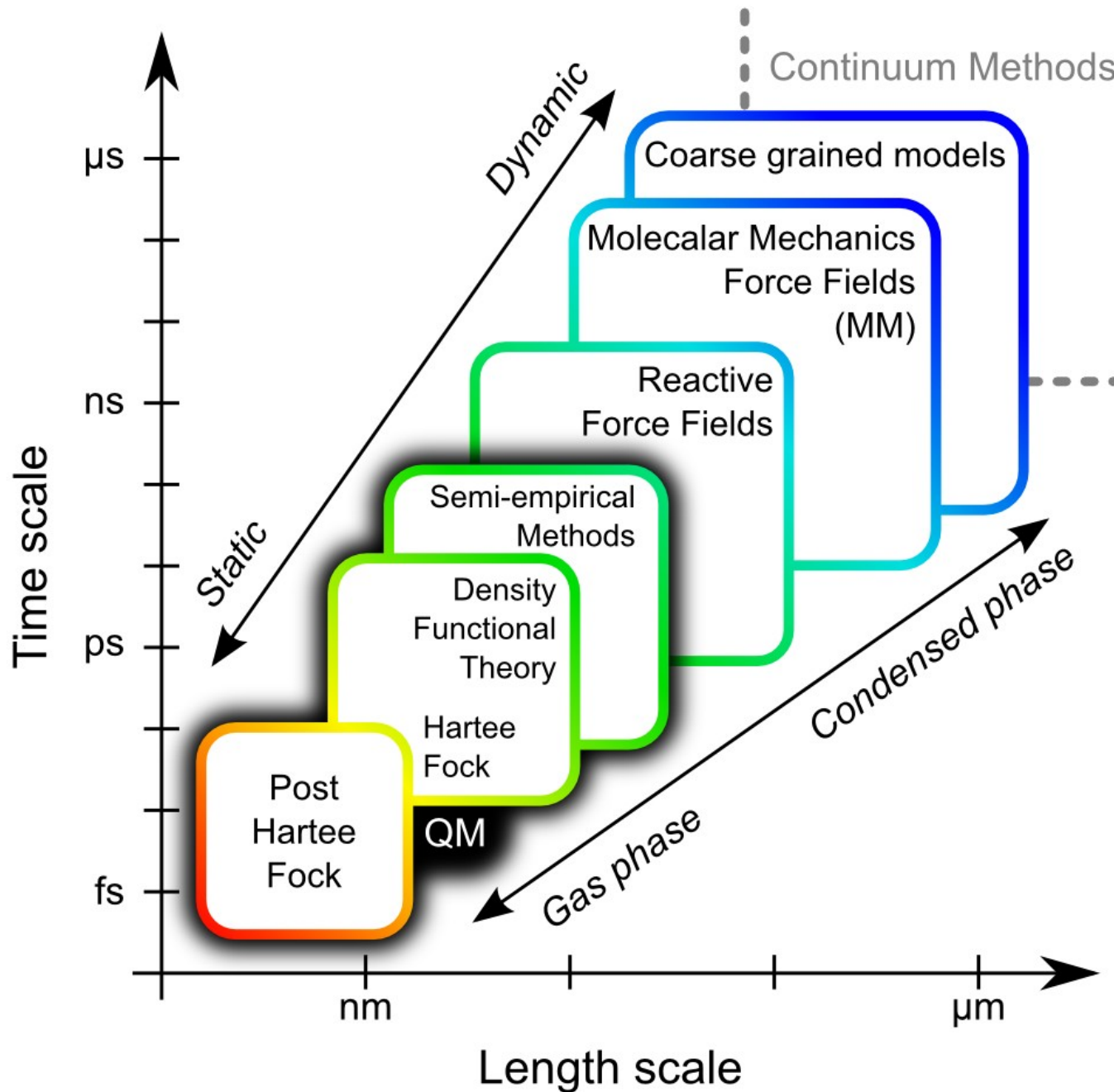
Work of Catlow & Lewis

Kirschhock et al. *Chem. Eur. J.* **2005**, 11, pp4306

Exploring Chemical Compounds



Exploring Chemical Compounds



(1,2,3,4) Valence Force Fields

$$\sum_k \frac{1}{2} K \left(q_k - q_k^{(0)} \right)^2$$

(5) Electrostatics

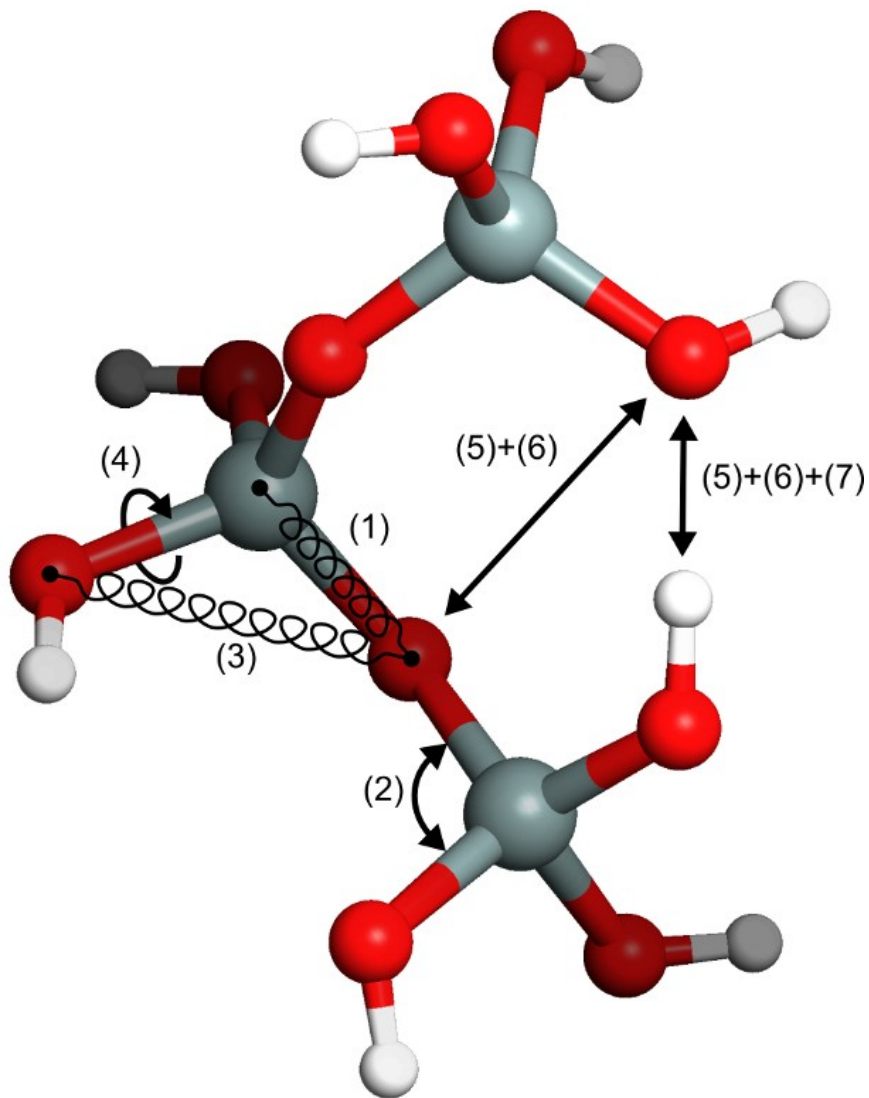
$$\sum_{i>j} \frac{q_i q_j}{r_{ij}}$$

(6) London Dispersion & Pauli Repulsion

$$\sum_{i>j} A_{ij}^{(p)} \exp\left(-\frac{r_{ij}}{B_{ij}^{(p)}}\right) - C_{ij}^{(6)} r_{ij}^{-6}$$

(7) Hydrogen Bonding

$$\sum_{i>j} -A_{ij}^{(hb)} \exp\left(-\frac{r_{ij}}{B_{ij}^{(hb)}}\right)$$



Chemical bonds are fixed during an FF simulation!

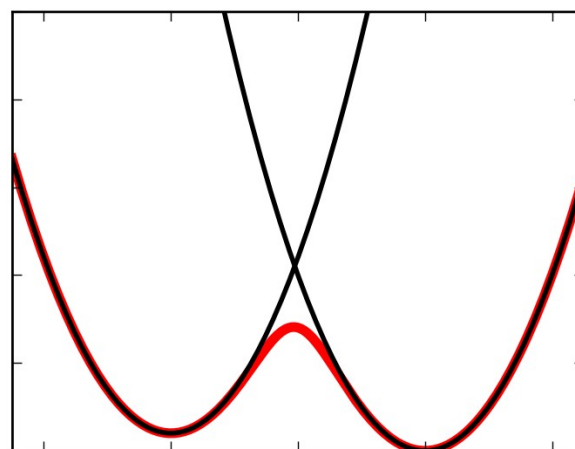
- * Terms in the valence FF
- * Exclusion rules for non-bonding interactions
- ▀ Is chemical sampling possible?

Reactive FF's

- Use empirical model for the bond order
- Computationally more expensive than traditional FF
- e.g. Tersoff, Brenner, ReaxFF, ...

Enveloping Distribution Sampling

Christ et al, *JCP* 2007, 126, p184110



$$\frac{\log(\exp(-\beta V_1) + \exp(-\beta V_2))}{\beta}$$

FF Interpolation

$$(1 - \lambda)V_1(\mathbf{r}^N) + \lambda V_2(\mathbf{r}^N)$$

(e.g. alchemical change)

Reference energy may be included in FF.

Old school approach:

- * Start from known parameters or make a guess
- * Try and fix parameters repetitively until some target is reached

Drawbacks: labor intensive & boring

Better approach:

- * Define an objective function that measures the FF quality
- * Minimize the objective numerically

Drawbacks: no physical insight & rank deficiency

Even better:

- * Associate FF parameters with QM 'observables'
(where possible, avoid parameter fitting)
- * Carefully designed cost functions

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FF = Potential energy for atoms in molecules

⇒ Obvious question: **"What are atoms in molecules?"**

Strict **definition is not possible**, but several *partitioning schemes* exist.

1955: Mulliken Population Analysis

1970: Löwdin Population Analysis

1977: Hirshfeld Partitioning

1985: Natural Population Analysis

1994: Bader Partitioning

2007: Iterative Hirshfeld Partitioning

Properties that can be derived:

- * Atomic charges (& multipole expansion)
- * Bond orders
- * Pairwise electrostatic interactions
- * Atomic spin density
- * Overlap of atomic densities
- * Condensed reactivity descriptors
- * ...

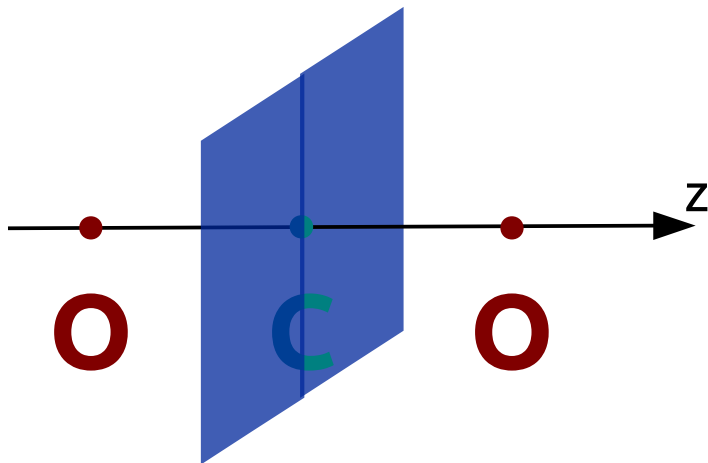
Hirshfeld partitioning

Density partitioning in general:

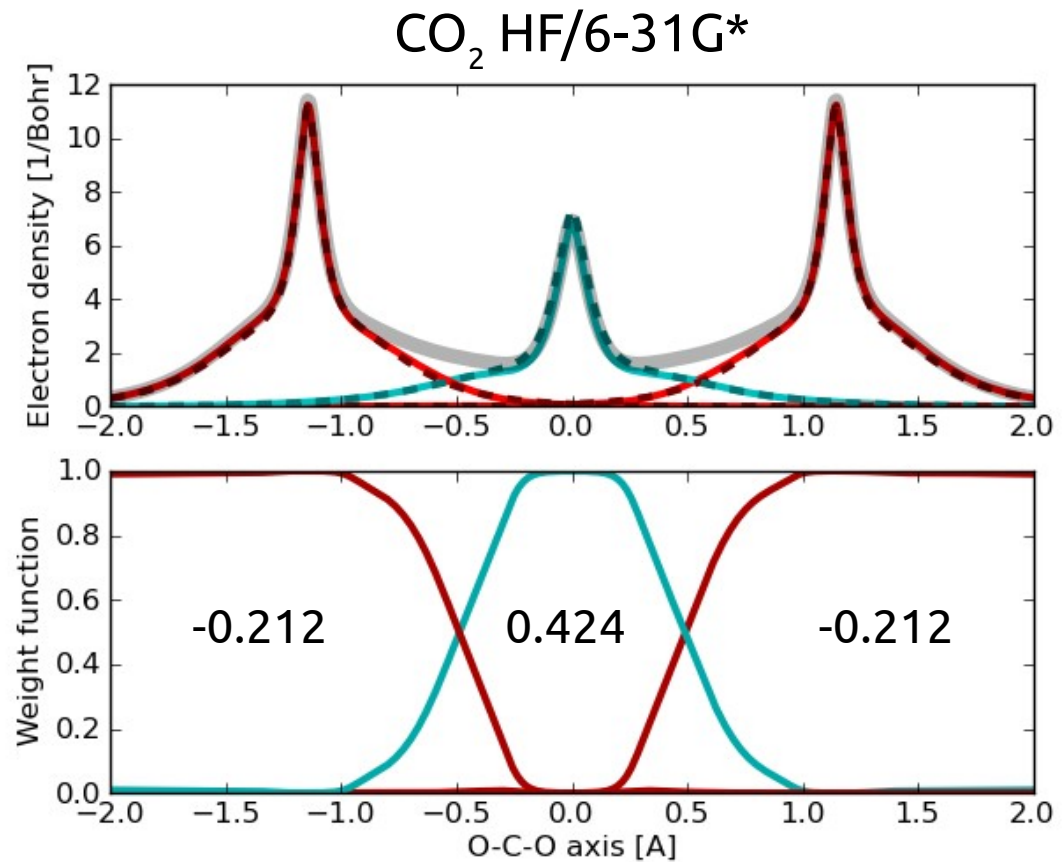
$$\rho_A(\mathbf{r}) = w_A(\mathbf{r})\rho_{\text{mol}}(\mathbf{r}) \quad \sum_A w_A(\mathbf{r}) = 1 \quad N_A = \int \rho_A(\mathbf{r}) d\mathbf{r}$$

Hirshfeld partitioning:

$$w_A(\mathbf{r}) = \frac{\rho_{0,A}(|\mathbf{r} - \mathbf{R}_A|)}{\sum_B \rho_{0,B}(|\mathbf{r} - \mathbf{R}_B|)}$$



Hirshfeld, *Theoret. Chim. Act.* **1977**, 44, p129



Iterative Hirshfeld Partitioning

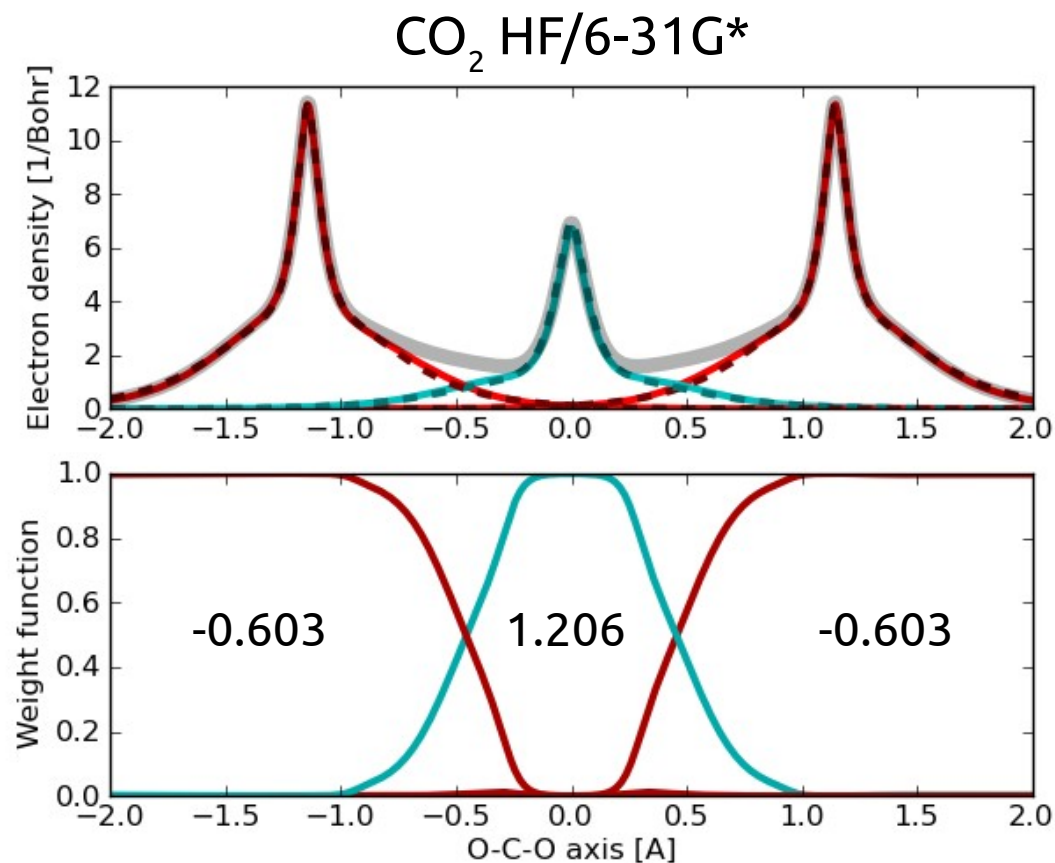
Density partitioning in general:

$$\rho_A(\mathbf{r}) = w_A(\mathbf{r})\rho_{\text{mol}}(\mathbf{r}) \quad \sum_A w_A(\mathbf{r}) = 1 \quad N_A = \int \rho_A(\mathbf{r})d\mathbf{r}$$

Hirshfeld-I partitioning:

$$w_A(\mathbf{r}) = \frac{\rho'_{0,A}(|\mathbf{r} - \mathbf{R}_A|, N_A)}{\sum_B \rho'_{0,B}(|\mathbf{r} - \mathbf{R}_B|, N_B)}$$

$$\rho'_{0,A}(r, N_A) = (N_A - \lfloor N_A \rfloor)\rho'_{0,A}(r, \lfloor N_A \rfloor) + (\lceil N_A \rceil - N_A)\rho'_{0,A}(r, \lceil N_A \rceil)$$



Bultinck et al., *JCP* **2007**, 126, p144111

Other Properties

Hirshfeld

Hirshfeld-I

Atomic multipole expansion [a.u.]

	c	dz	qzz	ozzz
O1	-0.212	-0.114	0.036	0.506
C2	0.424	0.0	-1.160	0.0
O3	-0.212	0.114	0.036	

	c	dz	qzz	ozzz
O1	-0.603	0.217	-0.001	0.219
C2	1.206	0.0	-0.292	0.00
O3	-0.603	-0.217	-0.001	0.0

Pairwise atomic Coulomb interaction [E_h]

	O1	C2
C2	-0.771	
O3	0.023	-0.771

	O1	C2
C2	-0.783	
O3	0.093	-0.783

$$\iint \frac{\rho_A(\mathbf{r})\rho_B(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d\mathbf{r}d\mathbf{r}'$$

Overlap of atomic densities [a.u.]

	O1	C2
C2	0.253	
O3	0.002	0.253

	O1	C2
C2	0.214	
O3	0.004	0.214

$$\int \rho_A(\mathbf{r})\rho_B(\mathbf{r})d\mathbf{r}$$

Bond orders

	O1	C2
C2	2.46	
O3	0.30	2.46

	O1	C2
C2	2.08	
O3	0.39	2.08

$$\iint [\rho^{(2)}(\mathbf{r}, \mathbf{r}') - \rho(\mathbf{r})\rho(\mathbf{r}')]w_A(\mathbf{r})w_B(\mathbf{r}')d\mathbf{r}d\mathbf{r}'$$

Why Hirshfeld-I?

Pragmatic reasons (benchmarks on large sets of molecules)

- * Charges reproduce the ESP
 - some small molecules (like CO₂) are exceptions
 - 10% to 20% error on ESP with charges
 - 5% to 10% error on ESP with charges and dipoles
- * Not sensitive to basis set (e.g. diffuse functions)
- * Robust with respect to conformational changes
- * Does not suffer from buried atom problem (like ESP charges)
- * 'Easy' to compute

Conceptual reason

- * Hirshfeld-I atoms are very similar to atoms in vacuum

**Don't get too enthusiastic!
Hirshfeld-I is still just a scheme.**

ESP Fitted charges minimize:

$$\chi^2 = \sum_{g=1}^{N_g} w_g \left(V_{\text{AI}}(\mathbf{r}_g) - \sum_{a=1}^{N_a} \frac{q_a}{\|\mathbf{r}_a - \mathbf{r}_g\|} \right)^2 + \text{some penalty}$$

Often used for FF development (e.g. RESP for AMBER)

but ...

- 1) Buried charges are ill-defined, sensitive to choice of grid
- 2) Sensitive to conformational changes

because ...

- 1) Ill-conditioned least-squares system
- 2) Atomic multipole expansions are truncated after the monopole.

This error is compensated by overfitting the charges.

**This is also just a scheme,
i.e. one to partition the ESP.**

Computational details

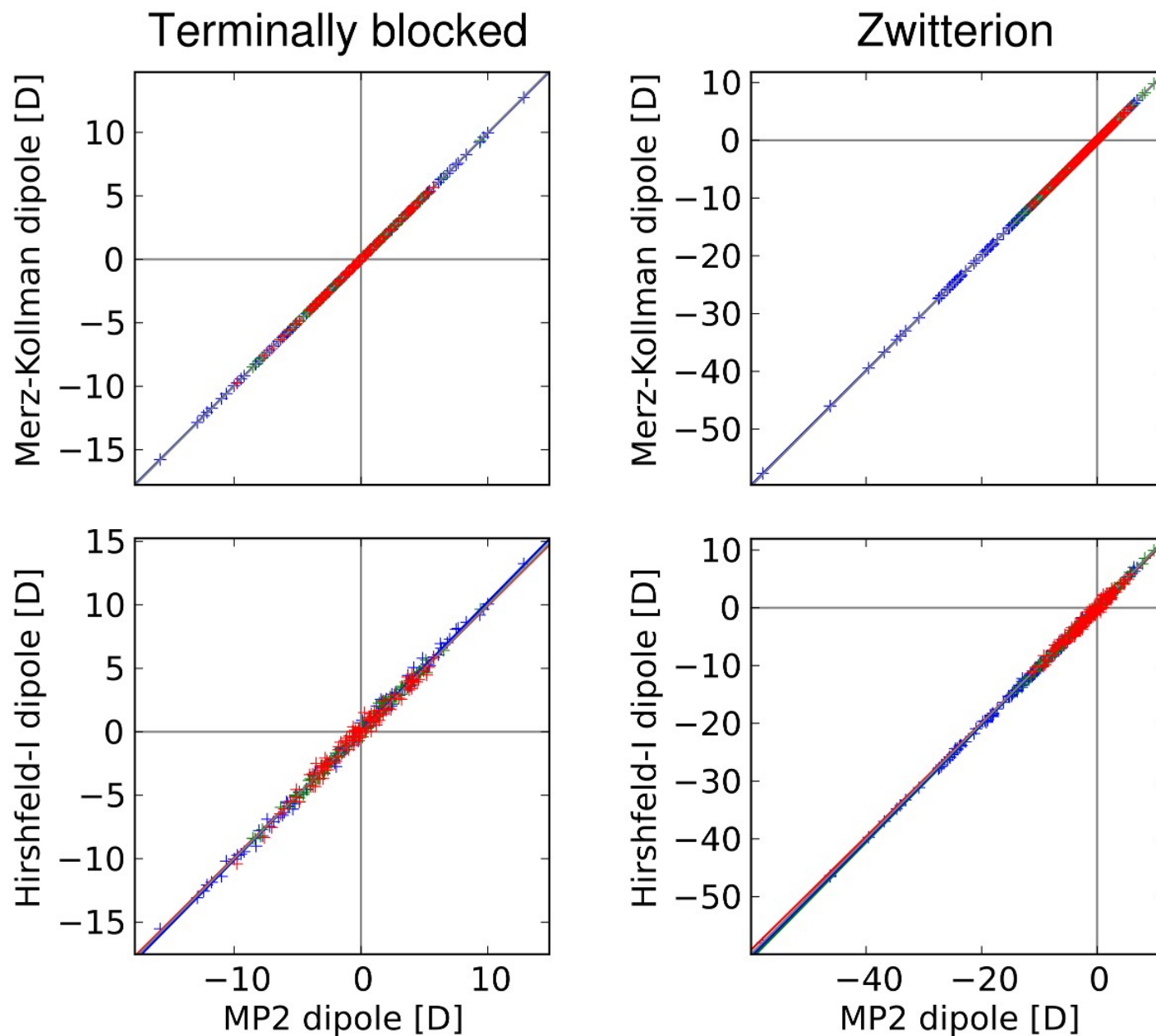
- 1) Random penta alanine conformers
 - * 103 Terminally blocked
 - * 134 Zwitterionic
- 2) B3LYP/6-31Gd geometry optimizations
- 3) MP2/Aug-cc-pVTZ single point computations
- 4) Charges derived with a selection of schemes:
 - * Merz-Kollman
 - * Hirshfeld-I

Benchmarks

- 1) Comparison of MP2 dipole with charge-derived dipole
- 2) Sensitivity of charges to conformational changes

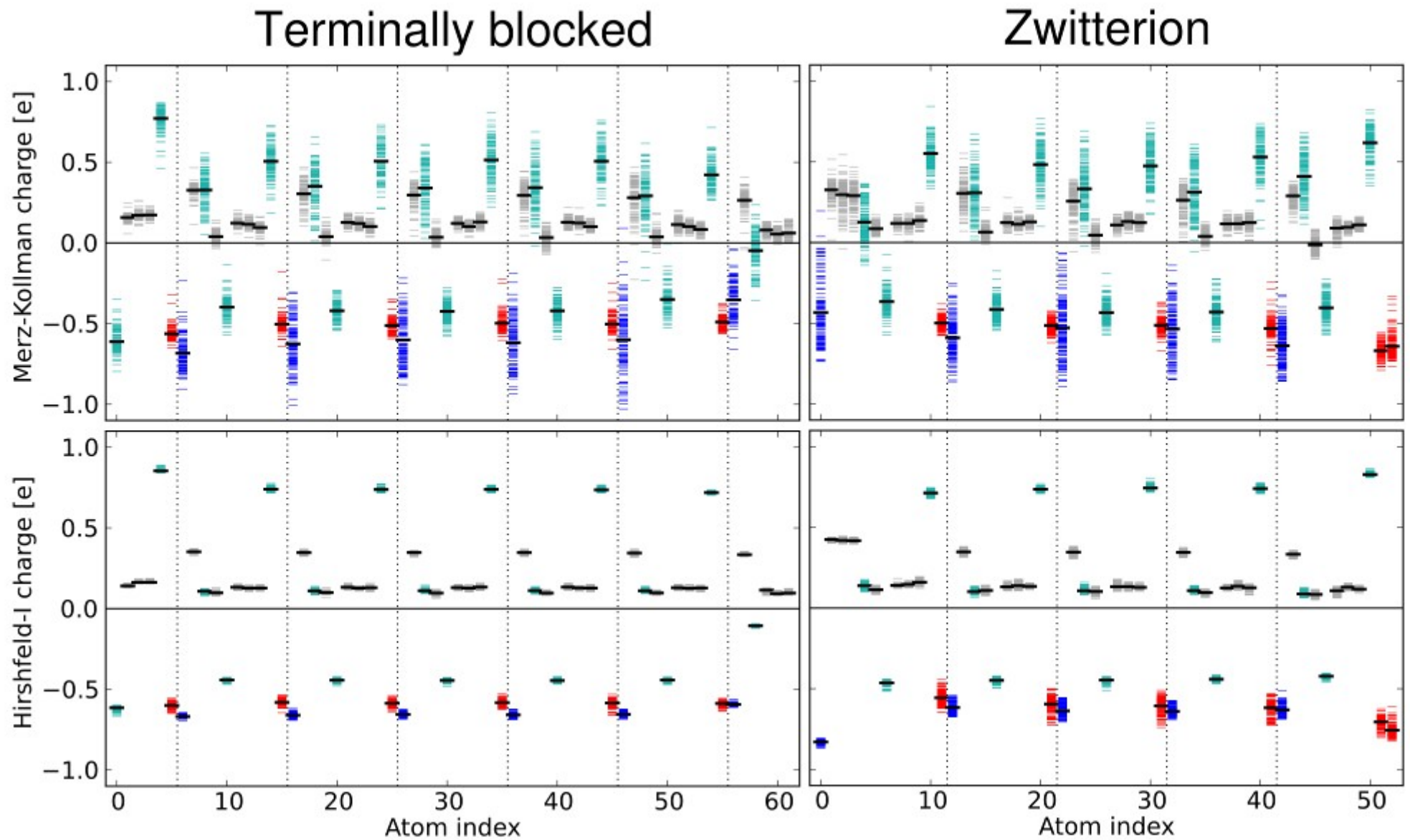
(paper in preparation)

Penta Alanine Benchmark



(paper in preparation)

Penta Alanine Benchmark



(paper in preparation)

One limitation observed so far ...

Poor point charge ESP's in very polar systems:

- Zeolites
- Oxide part in metal-organic frameworks

Hirshfeld-I is still young, and needs more testing

(paper in preparation)

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Atomic densities (or weight functions) provide...

1) Breakdown of the electrostatic interactions

- *MM charges*
- *Polarizable FF's (charge equilibration, inducible dipoles)*
- *Intermolecular charge-transfer*

$$q_A, d_{A,x}, d_{A,y}, d_{A,z}, \dots$$
$$\iint \frac{\rho_A(\mathbf{r})\rho_B(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d\mathbf{r}d\mathbf{r}'$$

2) Pauli repulsion ~ overlap of (singlet state) densities

- *Overlap of partitioned atoms -> pairwise potentials*

$$\int \rho_A(\mathbf{r})\rho_B(\mathbf{r})d\mathbf{r}$$

3) Bond order in dimers

- *Pairwise breakdown of weak covalent bonding*

$$\iint [\rho^{(2)}(\mathbf{r}, \mathbf{r}') - \rho(\mathbf{r})\rho(\mathbf{r}')]w_A(\mathbf{r})w_B(\mathbf{r}')d\mathbf{r}d\mathbf{r}'$$

4) Partitioning of linear response

- *Changes in charges/dipoles due to external field*
=> *Second order parameters in polarizable FF*

cfr. WOFF: Rotenberg, B. et al *PRL* **2010**, 104, 138301

Water dimer

Two scans of the water dimer

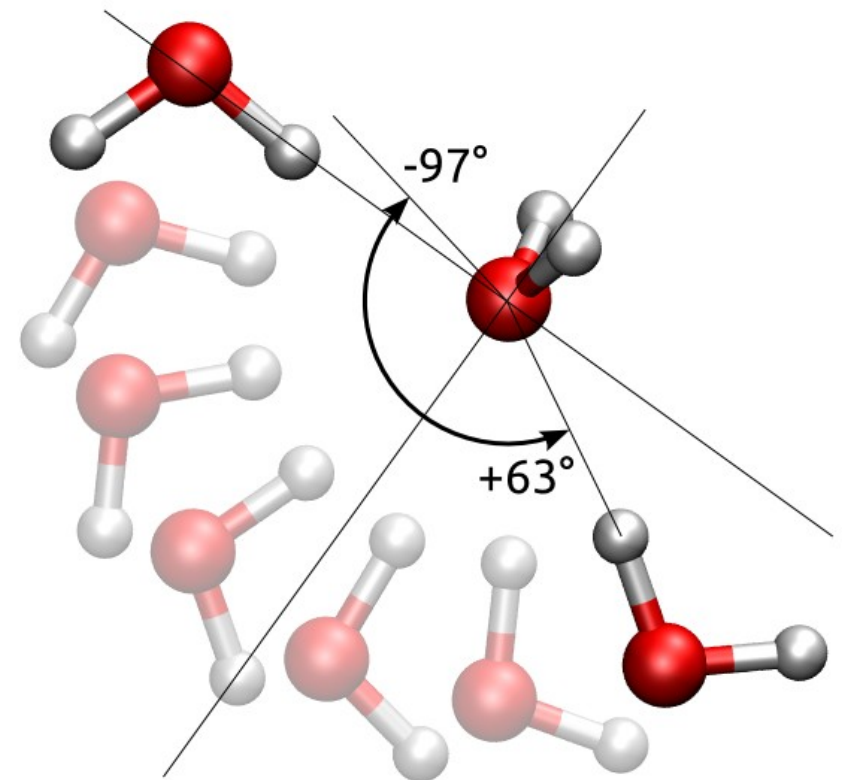
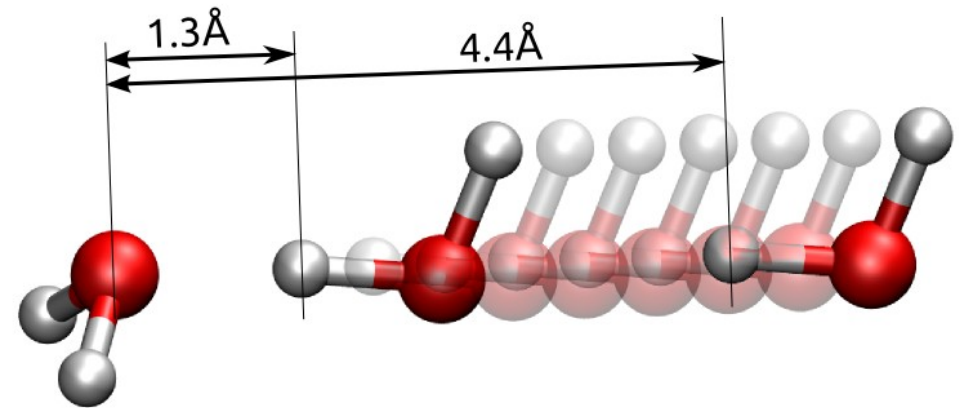
- * Both go through optimal geometry
- * PBE/DGTZVP
- * cp. cor. Interaction energies

AIM properties computed at each point

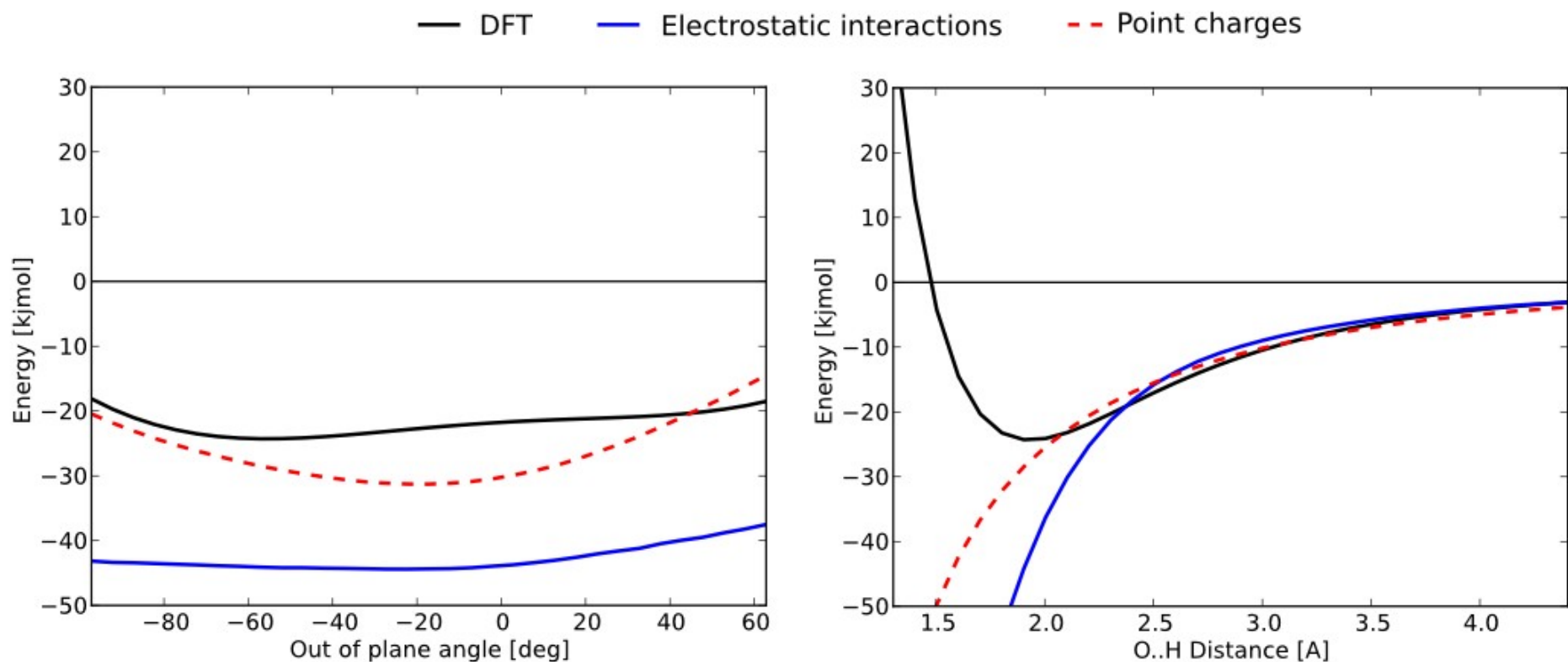
- * Charges

AIM pairwise properties

- * Electrostatic interaction
- * Bond orders
- * Pauli Overlap

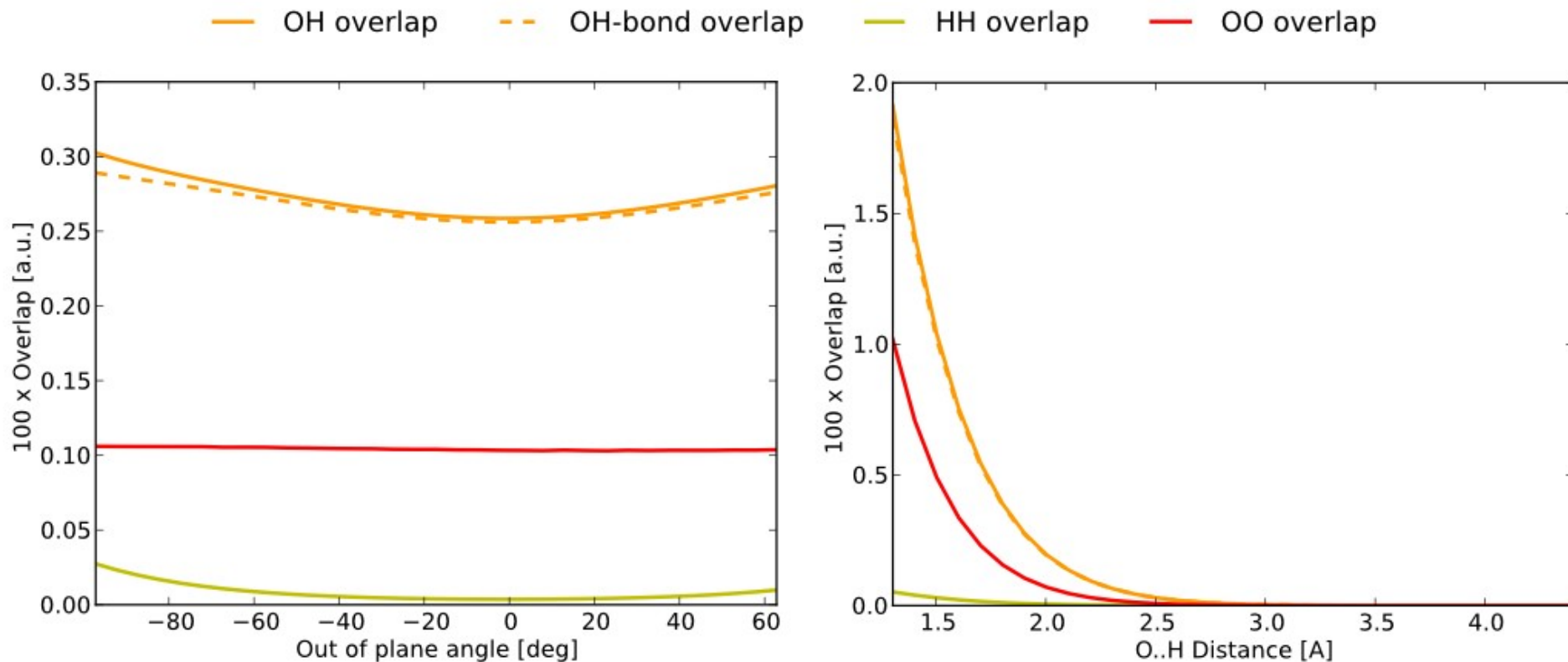


Electrostatic Interactions



- * Electrostatic interactions OK at long distances
- * Hirshfeld-I dipoles (& multipoles) are needed at short distance.
- * Subtle angular variations are missing

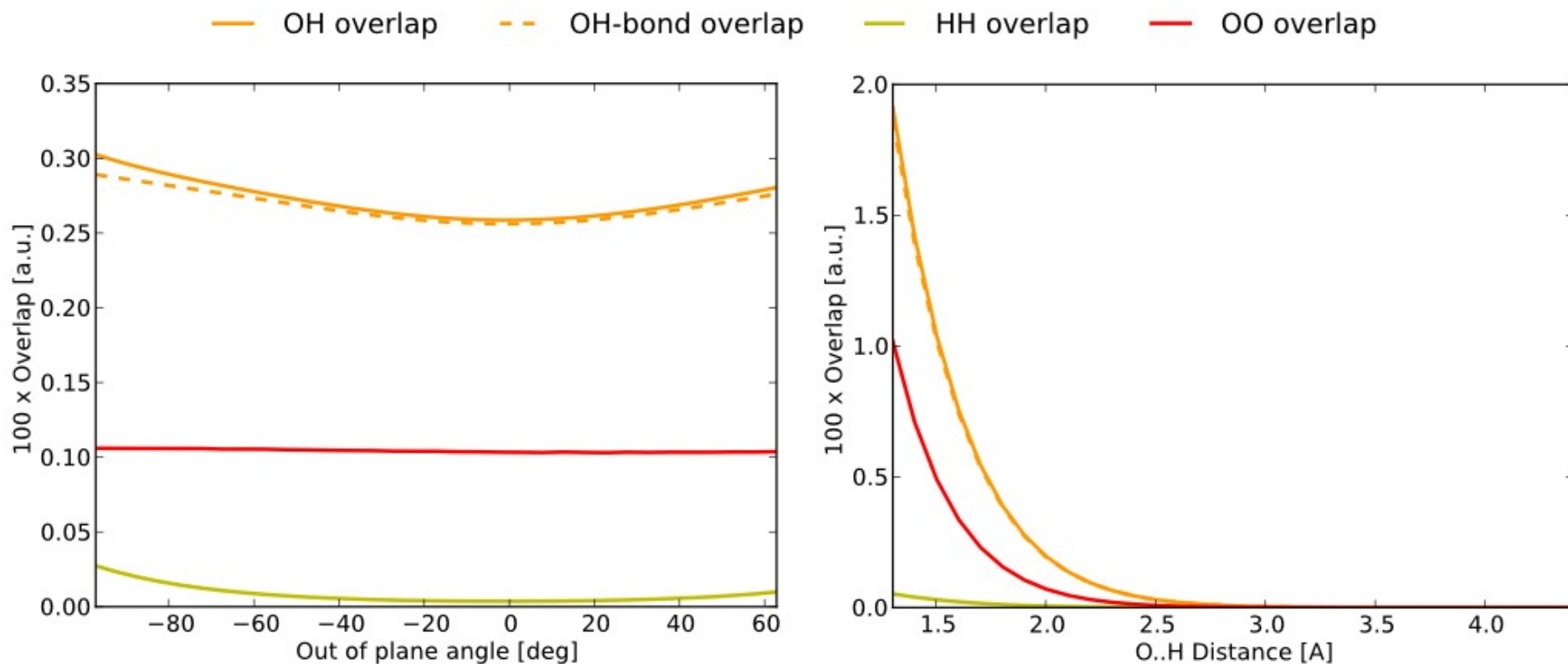
Pauli Repulsion



$$O_{ij} = \int \rho_i(\mathbf{r})\rho_j(\mathbf{r})d\mathbf{r}$$

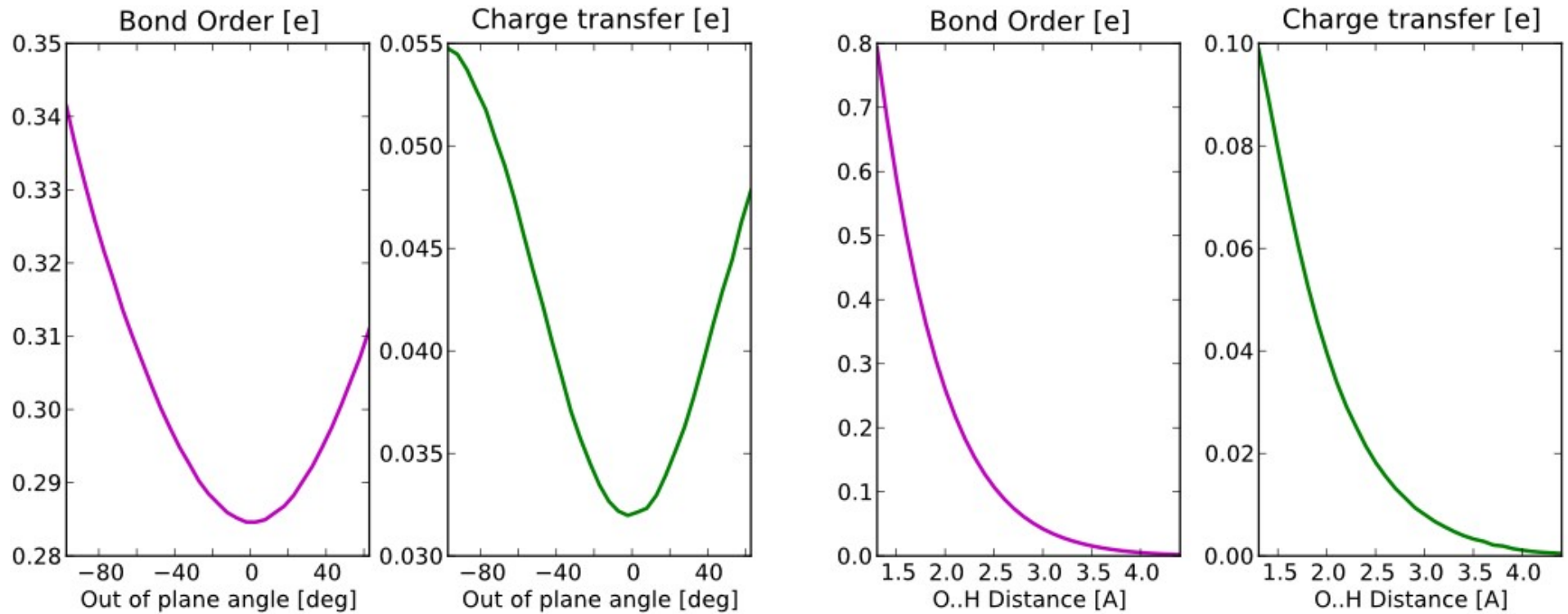
XY overlap = sum of overlap integrals over all pairs X and Y
OH-bond overlap = overlap integral for O and H in the HBond only

Pauli Repulsion



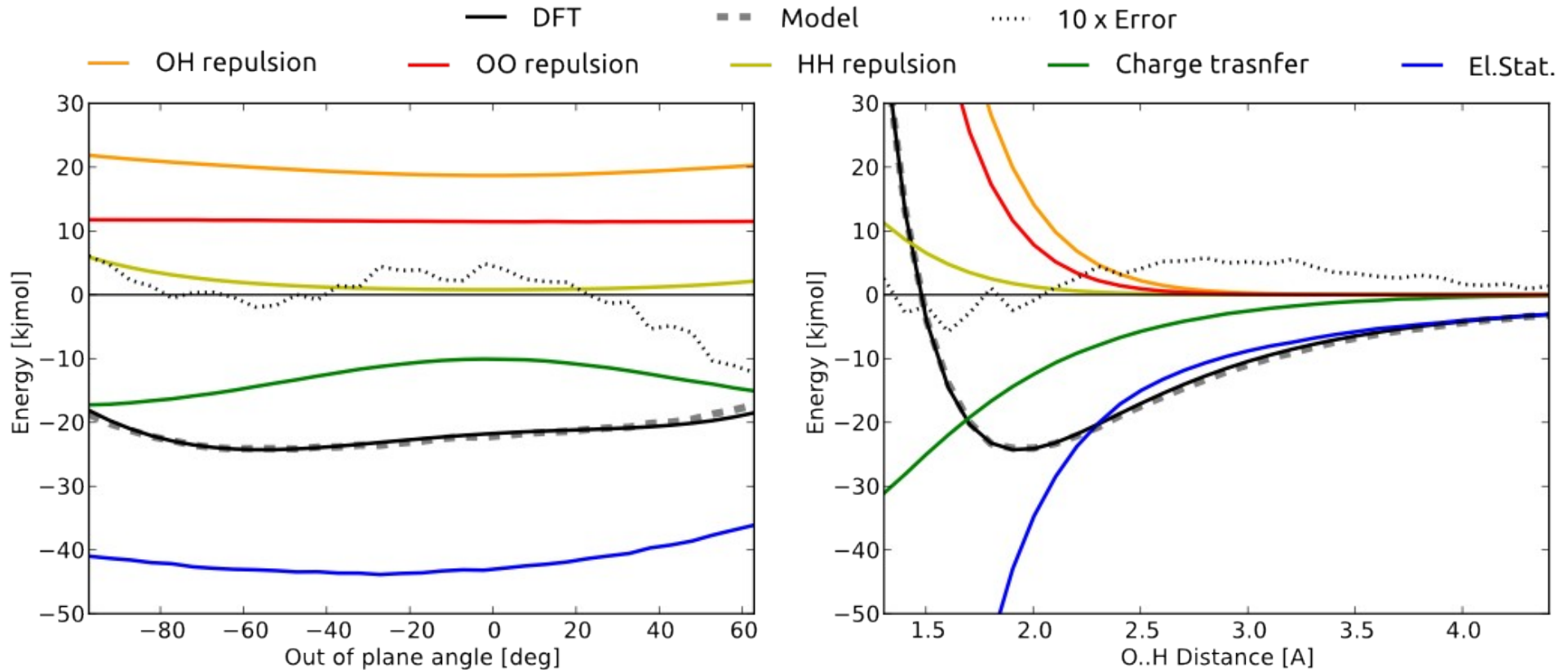
- * No effect of lone pairs
- * $A \cdot \exp(-B \cdot R)$ is a good approximation
- * OH pair in hydrogen bond has angular overlap dependence

Bond order & charge transfer



- * Electrons go from acceptor H_2O to donor H_2O during HBond formation
- * Charge transfer and Bond order are similar but not identical
- * OH overlap $\sim (\text{Bond order})^2$
- * Large relative fluctuations in charge transfer

Model for the interaction energy



$$E_{\text{DFT}} \approx A_1 O_{\text{OH}} + A_2 O_{\text{HH}} + A_3 O_{\text{OO}} - A_4 Q_{\text{trans}} + E_{\text{el.stat.}}$$

in atomic units: $A_1 = 2.75$ $A_2 = 8.22$ $A_3 = 4.22$ $A_4 = 0.12$

- * Insightful, but still using AIM data from DFT computations.
- * Pairwise repulsion parameters are trivial to extract.
- * Explicit exchange term (\sim bond order) is not helpful.
- * Should be combined with EDA (Morokuma) or SAPT



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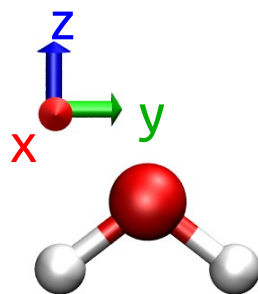
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Dipole Polarizability

Definition:

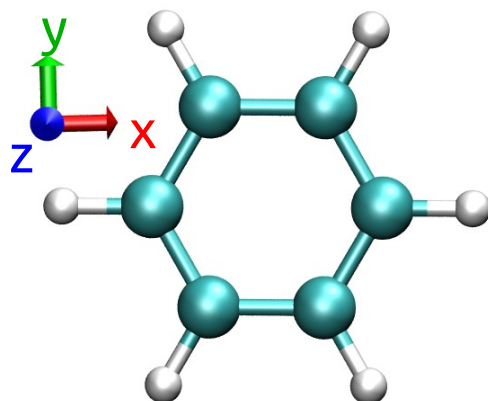
$$P_{\alpha\beta} = \frac{\partial d_{\alpha}^{(mol)}}{\partial E_{\beta}}$$



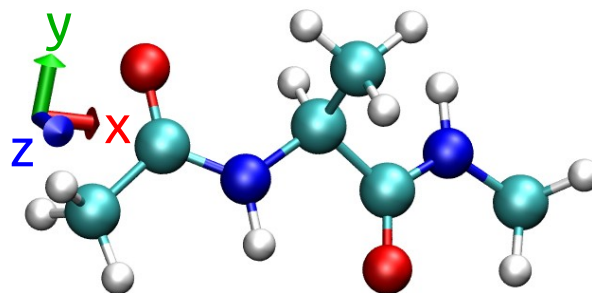
[4.28	0	0]
[0	8.19	0]
[0	0	6.78]

A few examples
computed with
PBE/DGTZVP:

- water
 - benzene
 - alaninedipeptide
- (values in a.u.)



[77.38	0	0]
[0	77.38	0]
[0	0	34.15]



[122.8	4.19	-1.41]
[4.19	89.57	-5.27]
[-1.41	-5.27	73.80]

Definition

Empirical model for the linear response
(of the e^- density to a change in the external field)

Basic mathematical form is quadratic

$$E = \frac{1}{2} q^T J^{(qq)} q + q^T J^{(qd)} d + \frac{1}{2} d^T J^{(dd)} d$$

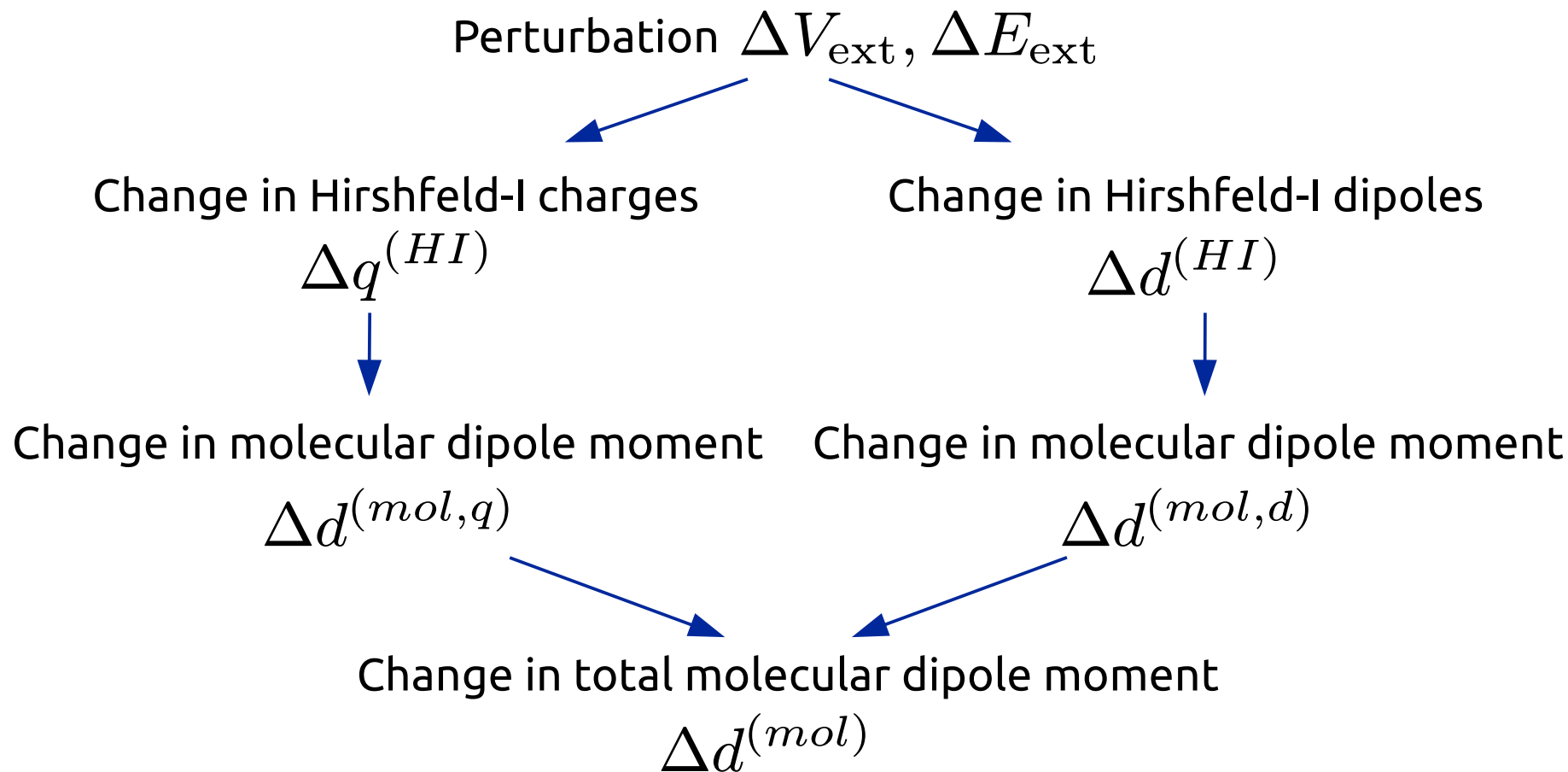
$q =$ vector with N atomic charges

$d =$ vector with $3N$ atomic dipole components

$$+ q^T x^{(q)} + d^T x^{(d)}$$
$$+ q^T \Delta V_{\text{ext}} + d^T \Delta E_{\text{ext}}$$

e.g. $J_{ij}^{(qq)} = \frac{1}{r_{ij}}$

Partitioning of Induced Dipole Moment



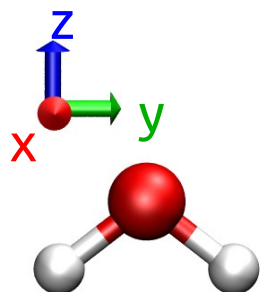
Do this with uniform X, Y and Z fields:

=> atomic charge contribution to $P_{\alpha\beta}$

=> atomic dipole contribution to $P_{\alpha\beta}$

Krishtal et al, *JCP* **2006**, 125, p034312.

Partitioning of Induced Dipole Moment

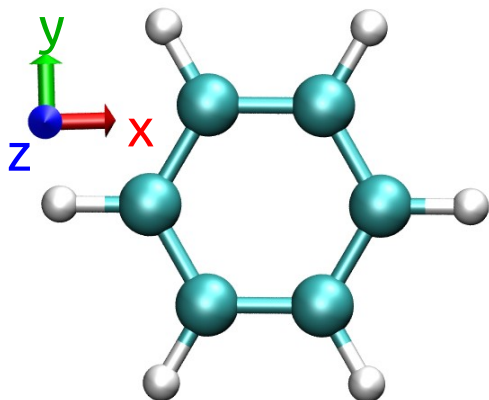


Charge contribution to $P_{\alpha\beta}$

$$\begin{bmatrix} 0 & 0 & 0 \\ 0 & 6.8 & 0 \\ 0 & 0 & 4.92 \end{bmatrix}$$

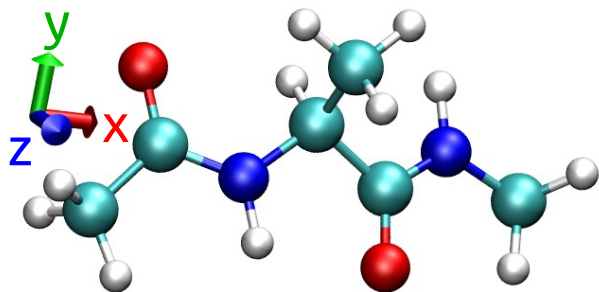
Dipole contribution to $P_{\alpha\beta}$

$$\begin{bmatrix} 4.28 & 0 & 0 \\ 0 & 1.39 & 0 \\ 0 & 0 & 1.86 \end{bmatrix}$$



$$\begin{bmatrix} 70.13 & -0.03 & 0 \\ 0.34 & 70.38 & 0 \\ 0.05 & 0.51 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 7.14 & 0.01 & 0.03 \\ -0.45 & 7.1 & -0.05 \\ -0.03 & -0.54 & 34.16 \end{bmatrix}$$



$$\begin{bmatrix} 101.7 & 2.63 & -0.14 \\ 2.63 & 69.05 & -12.97 \\ -0.17 & -12.9 & 45.07 \end{bmatrix}$$

$$\begin{bmatrix} 21.68 & 1.61 & -1.22 \\ 0.97 & 20.58 & 7.73 \\ -1.04 & 7.62 & 28.78 \end{bmatrix}$$

Mathematical form for the 'Electronic energy'

$$E_{\text{EEM}} = \left(\sum_i \chi_i q_i + \frac{1}{2} \eta_i q_i^2 \right) + \left(\frac{1}{2} \sum_{i,j \neq i} \frac{q_i q_j}{|\mathbf{r}_i - \mathbf{r}_j|} \right) + \left(\sum_i V_{\text{ext}}(\mathbf{r}_i) q_i \right)$$

Variables: q_i

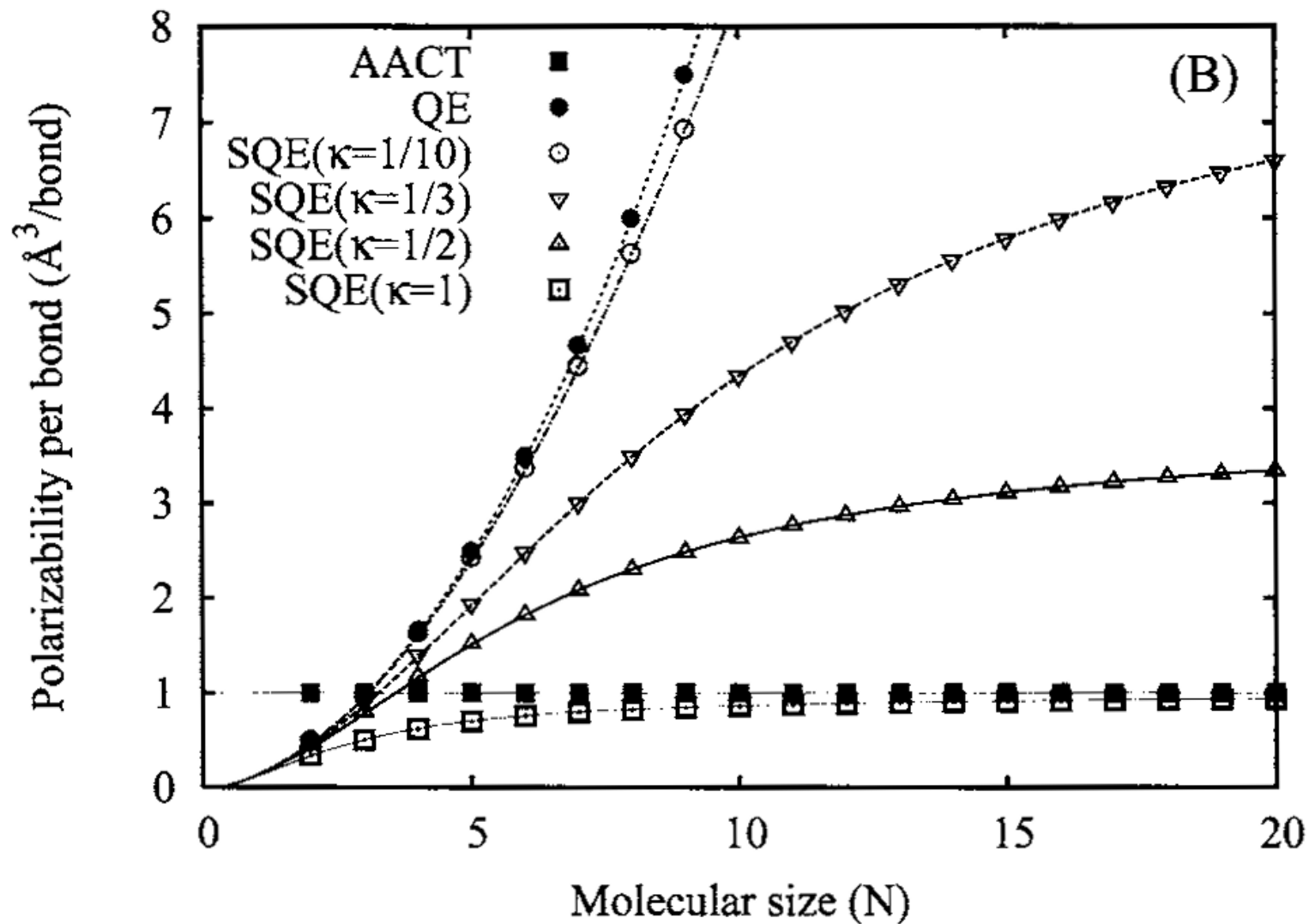
Parameters: $\chi_i, \eta_i, \bar{r}_i$

Charges minimize E_{EEM} (with total charge constraint)

$$\frac{\partial E_{\text{EEM}}}{\partial q_i} = \sum_i \left(\chi_i + V_{\text{ext}}(\mathbf{r}) + \eta_i q_i + \sum_{j \neq i} \frac{q_j}{|\mathbf{r}_i - \mathbf{r}_j|} \right) = \chi_{\text{mol}}$$

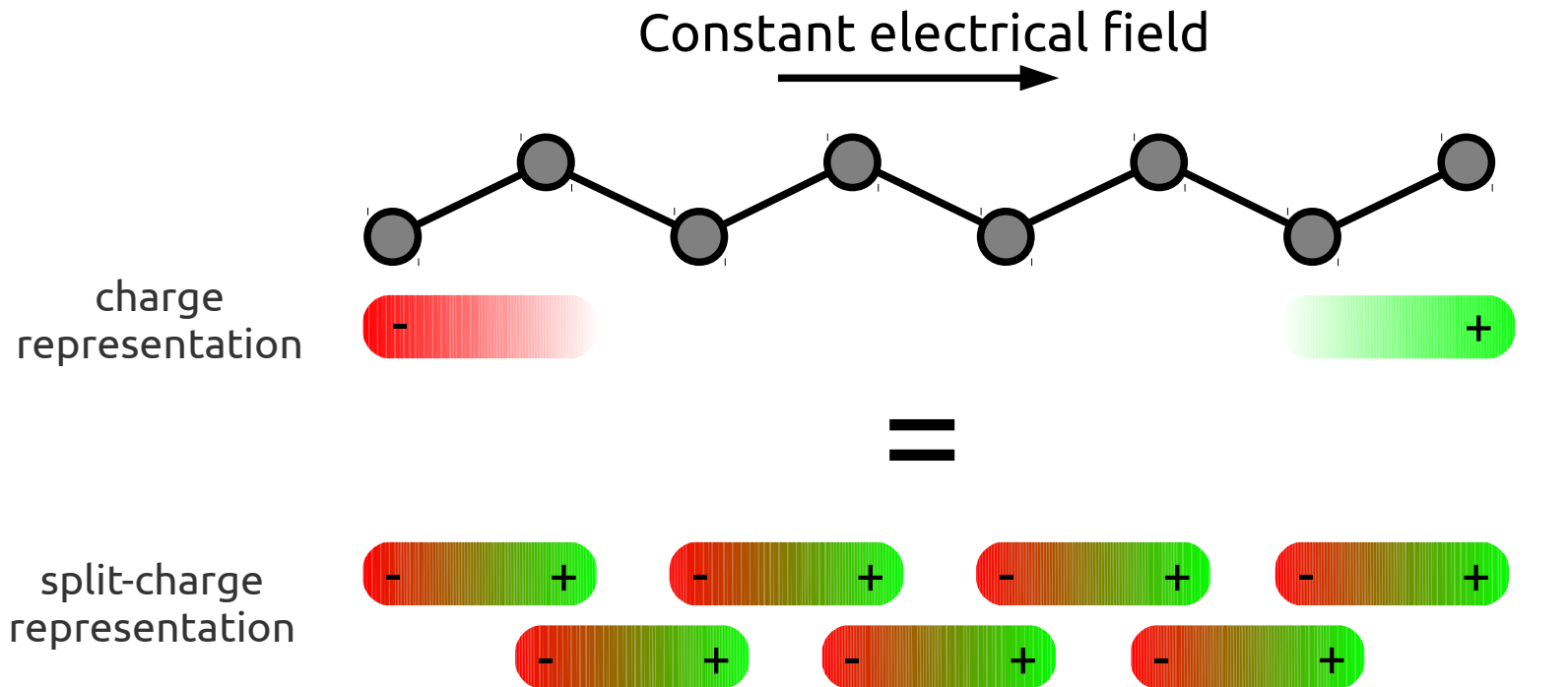
Mortier, W.; Ghosh, S.; Shankar, S. *JACS* **1986**, 108, 4315-4320.

Superlinear Scaling of Polarizability



Warren et al, *JCP* 2008, 128, p144110

Split Charge Equilibration



$$\begin{bmatrix} q_1 \\ \vdots \\ q_{N^a} \end{bmatrix} = \begin{bmatrix} +1 & 0 & \dots \\ -1 & +1 & \\ 0 & -1 & \\ \vdots & & \ddots \end{bmatrix} \begin{bmatrix} p_1 \\ \vdots \\ p_{N^b} \end{bmatrix}$$

$$q = Tp$$

$$(q = Tp + q_0)$$

Electronic energy (EEM)

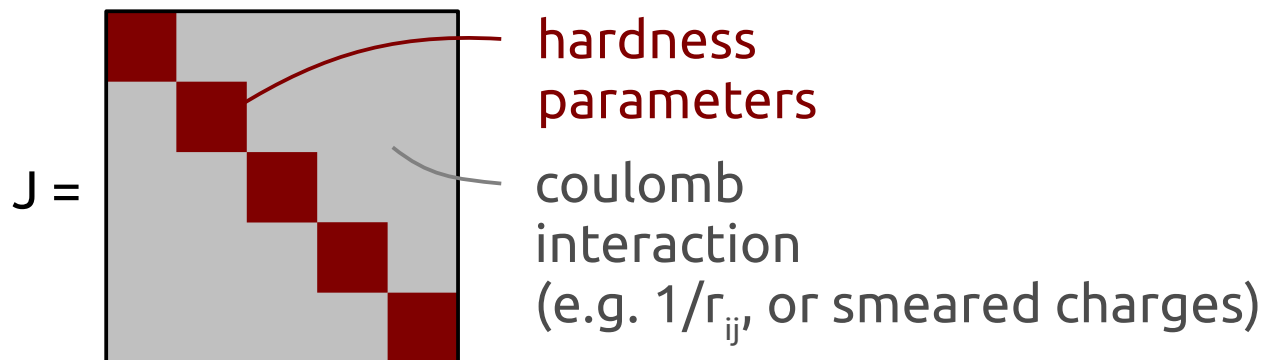
$$E_{\text{EEM}} = \frac{1}{2} q^T J q + x^T q \quad \text{with} \quad \sum_i q_i = 0$$

$$J_{ij} = \frac{1}{|\mathbf{r}_i - \mathbf{r}_j|} \quad \text{when} \quad i \neq j$$

$$J_{ii} = \eta_i$$

$$x_i = \chi_i + V_{\text{ext}}(\mathbf{r}_i)$$

Second order contributions



Electronic energy (EEM)

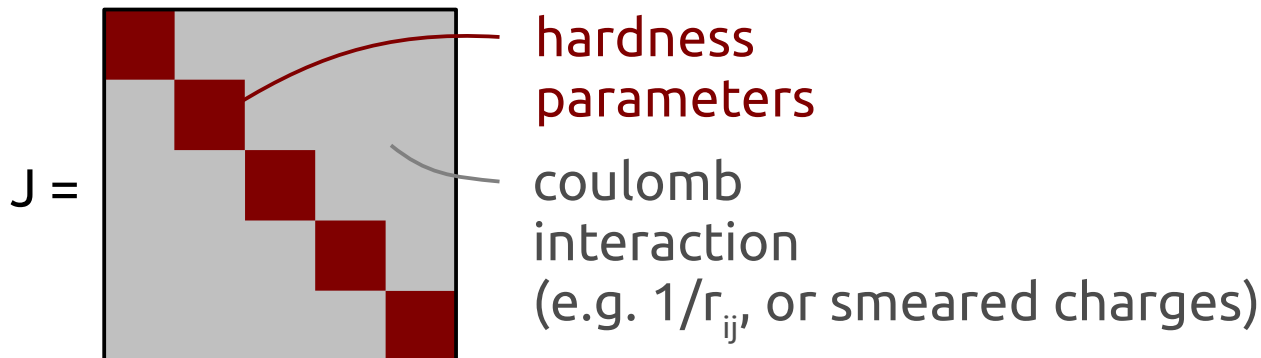
$$E_{\text{EEM}} = \frac{1}{2} p^T T^T J T p + x^T T p \quad \text{with} \quad q = T p$$

$$J_{ij} = \frac{1}{|\mathbf{r}_i - \mathbf{r}_j|} \quad \text{when} \quad i \neq j$$

$$J_{ii} = \eta_i$$

$$x_i = \chi_i + V_{\text{ext}}(\mathbf{r}_i)$$

Second order contributions



Electronic energy (SQE)

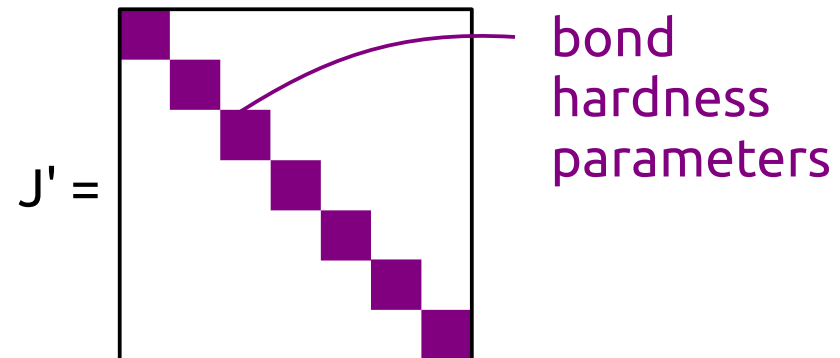
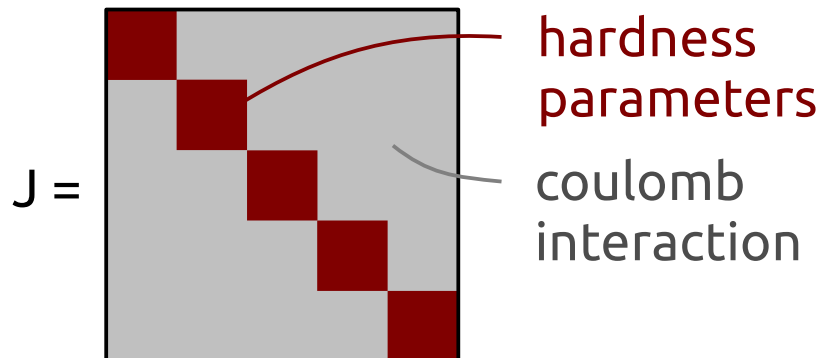
$$E_{\text{SQE}} = \frac{1}{2} p^T T^T J T p + \frac{1}{2} p^T J' p + x^T T p \quad \text{with} \quad q = T p$$

$$J_{ij} = \frac{1}{|\mathbf{r}_i - \mathbf{r}_j|} \quad \text{when} \quad i \neq j \quad J'_{kl} = \delta_{kl} \kappa_k \quad \text{bond hardness}$$

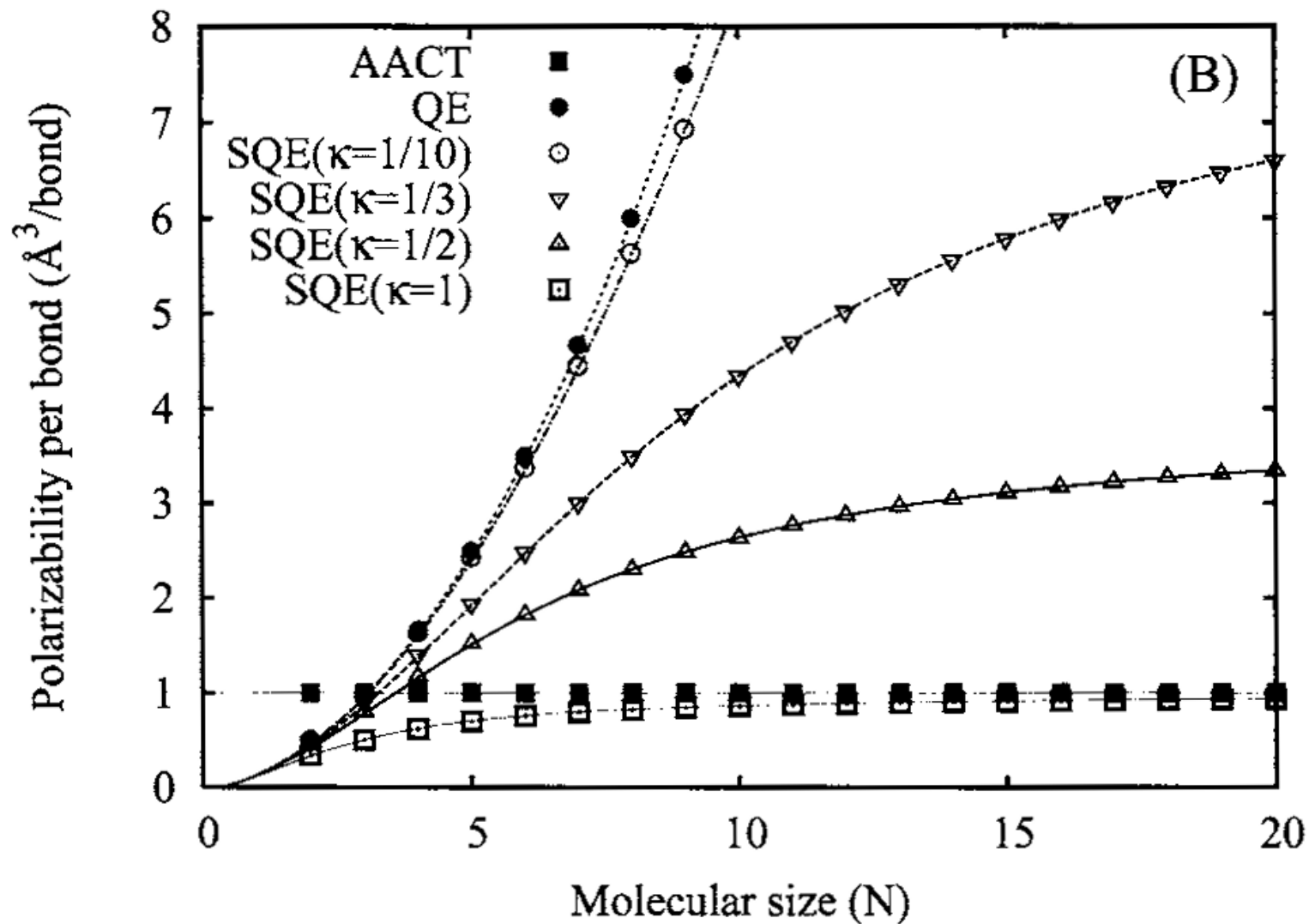
$$J_{ii} = \eta_i$$

$$x_i = \chi_i + V_{\text{ext}}(\mathbf{r}_i)$$

Second order contributions



Superlinear Scaling of Polarizability



Warren et al, *JCP* 2008, 128, p144110

Training set

- * 500 small organic molecules
- * Algorithmic selection from Pubchem
 - maximized diversity
 - constraints on atomtypes and number of atoms

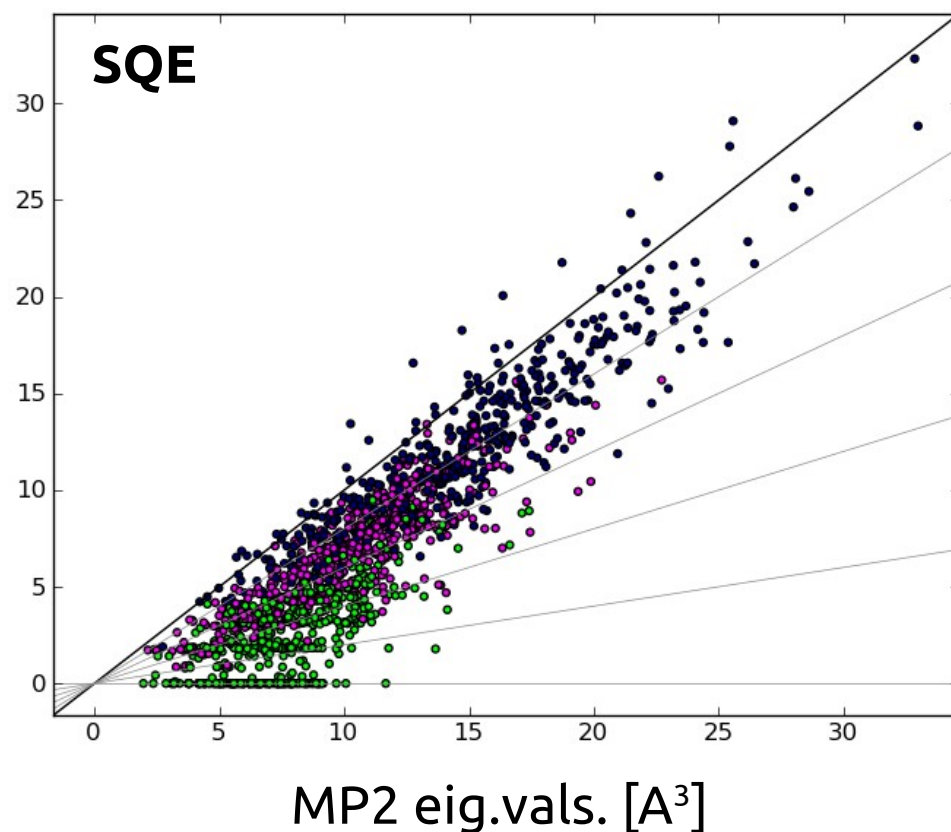
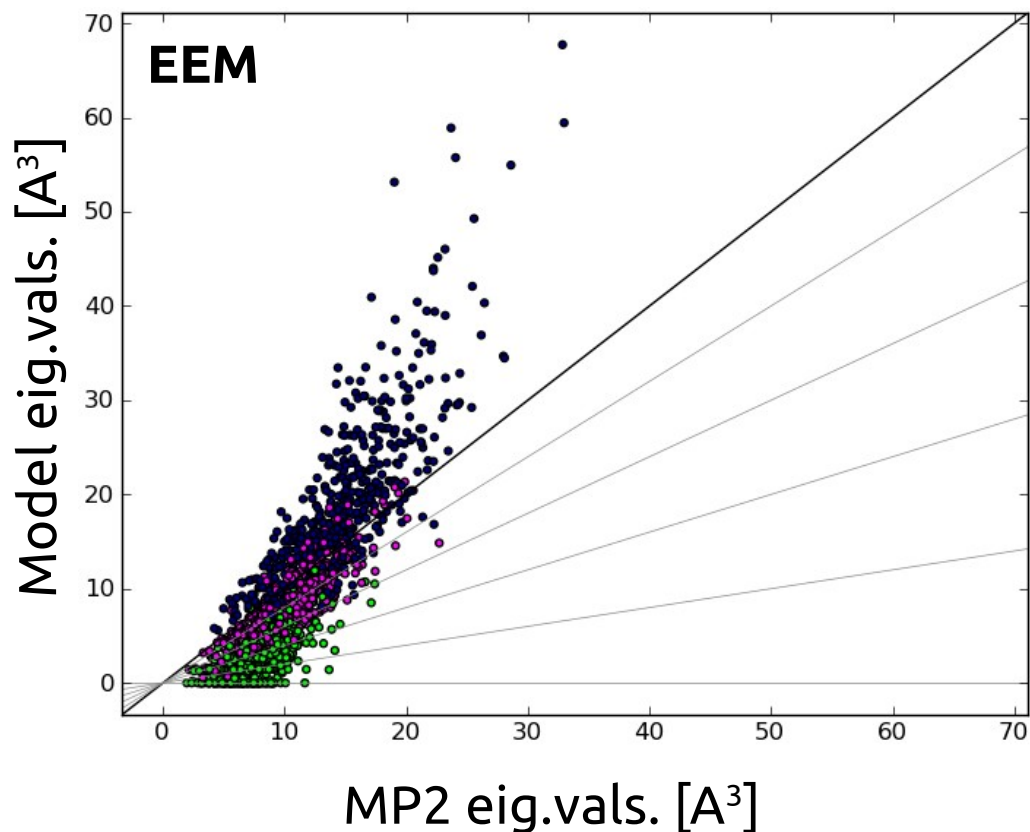
Calibration of EEM/SQE in 12 different ways

- * 3 charge schemes: Mulliken, Natural, Hirshfeld-I
- * 2 atom types: elements & elements+number of bonds
- * 2 cost functions: with and without linear response data

Conclusions (limited to small molecules)

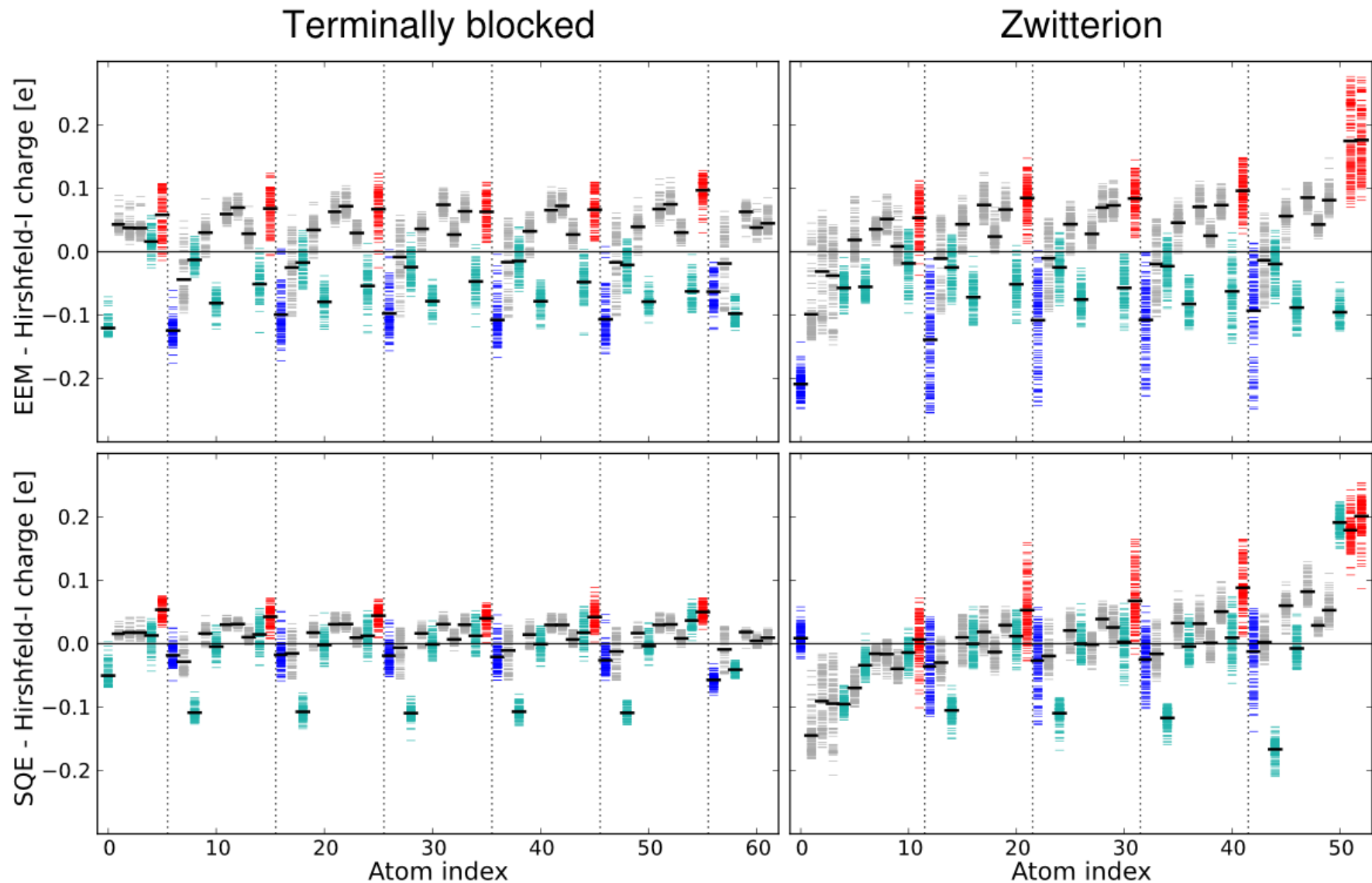
- * SQE outperforms EEM in all tests
- * Cross validation confirms transferability of parameters
- * Hirshfeld-I based calibrations are most useful
- * SQE model gives good dipole polarizabilities

Comparison of dipole polarizability eigenvalues (500 small organic molecules)



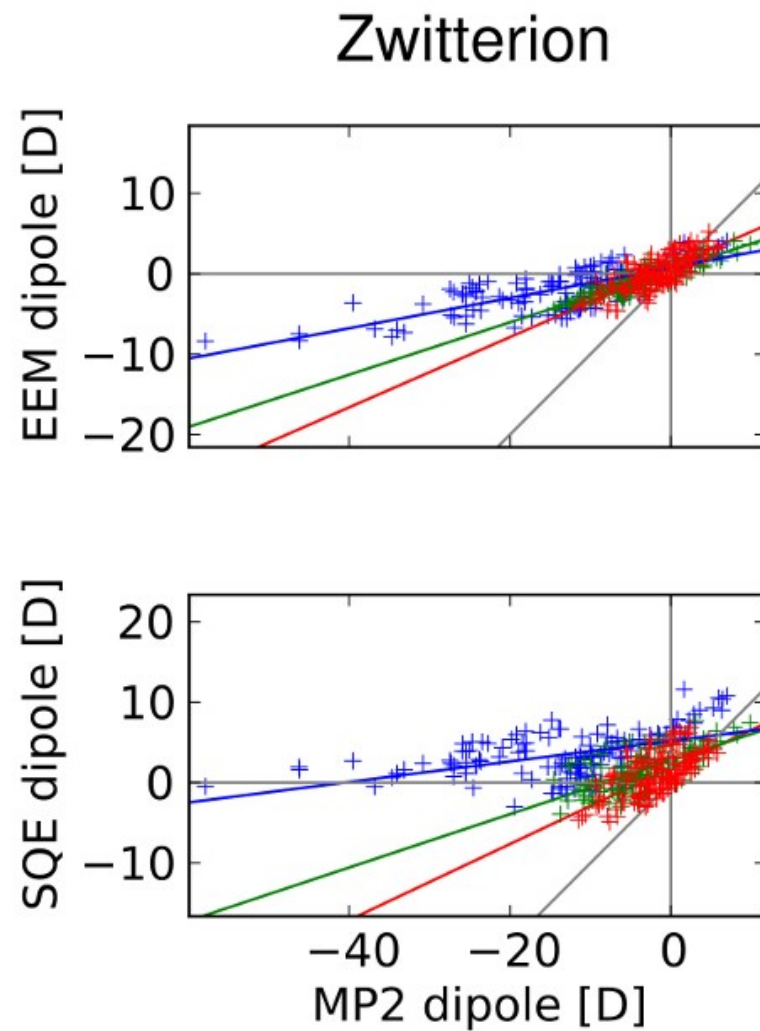
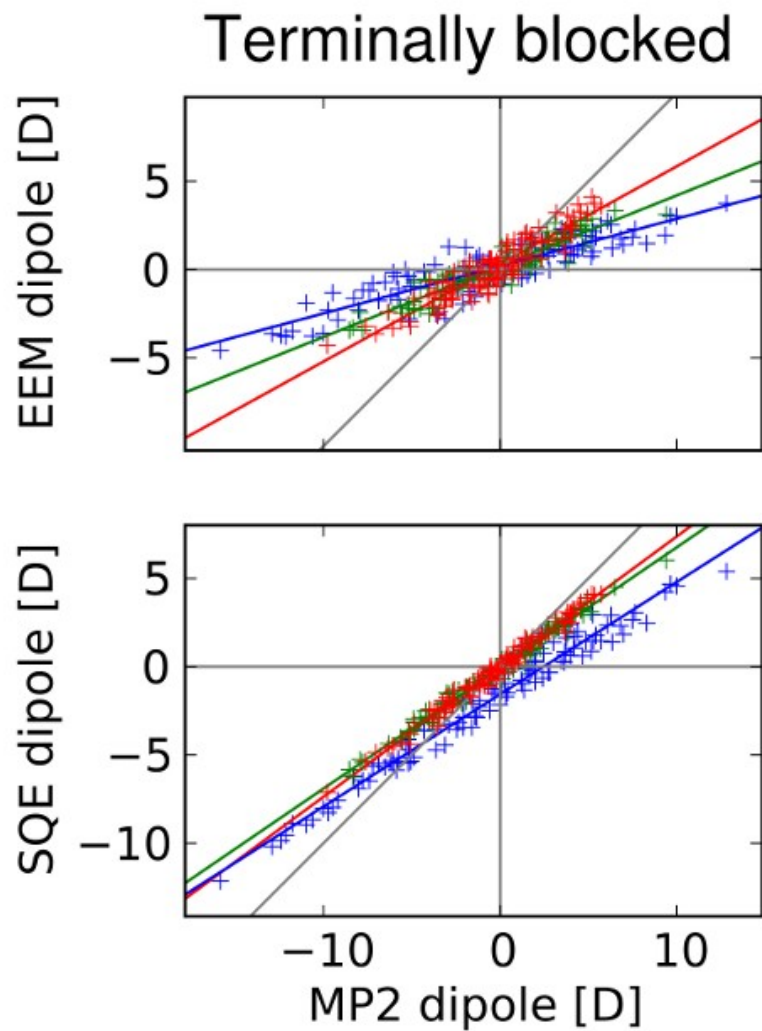
Verstraelen, T et al *JCP* **2009**, 131, p044127

Penta Alanine Benchmark



(paper in preparation)

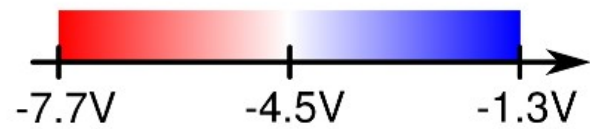
Penta Alanine Benchmark



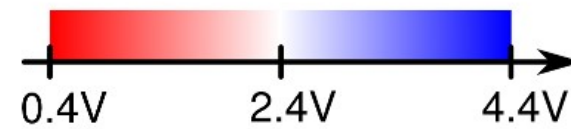
(paper in preparation)

Penta Alanine Benchmark

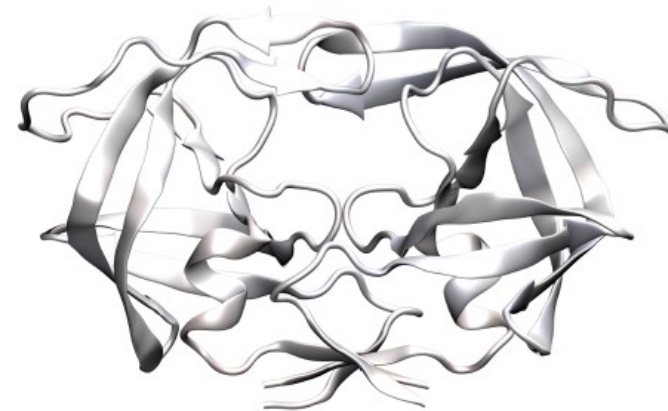
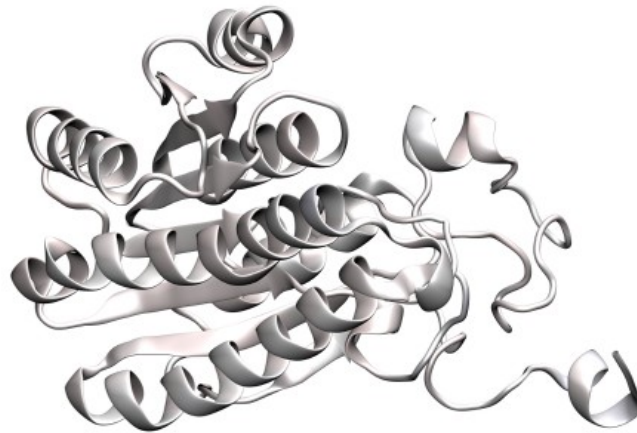
1HDC, $Q_{\text{tot}}=-7e$



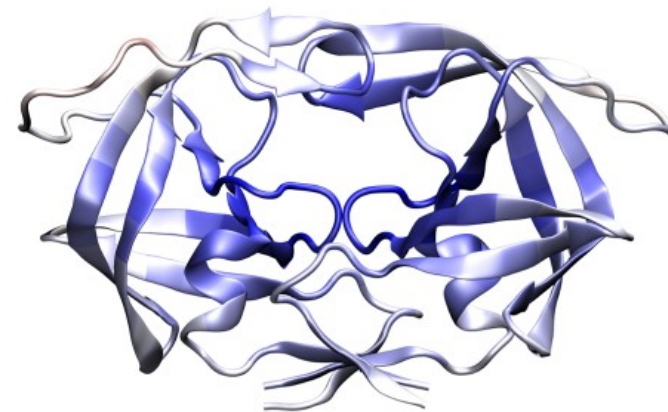
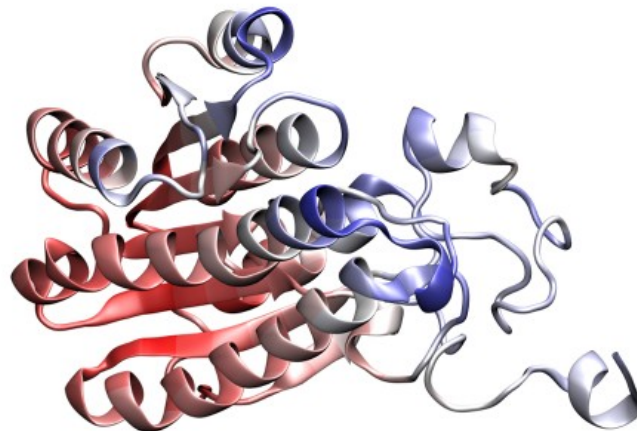
1HSG, $Q_{\text{tot}}=+2e$



EEM



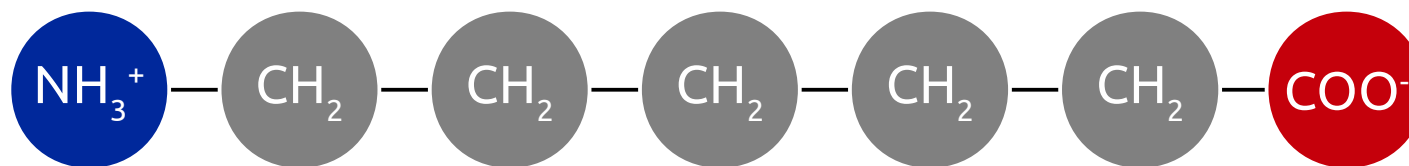
SQE



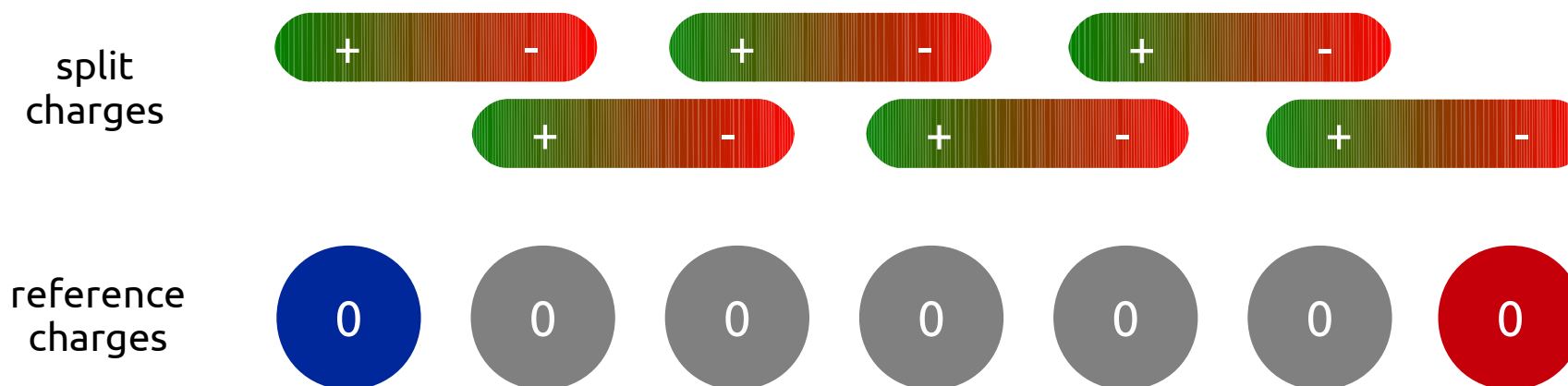
(paper in preparation)

Penta Alanine Benchmark

Charged molecules (and zwitter ions) are problematic.



SQE model: $q = Tp$

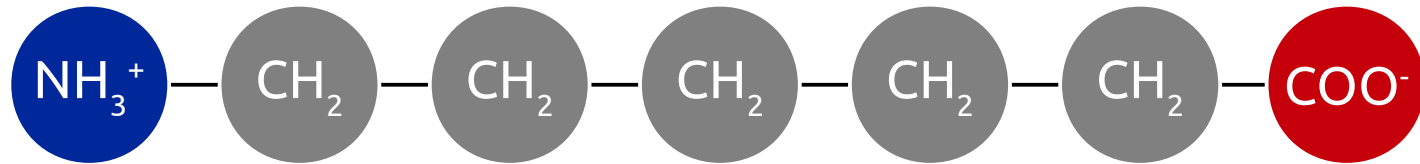


→ electronegativities of endpoints depend on chain length

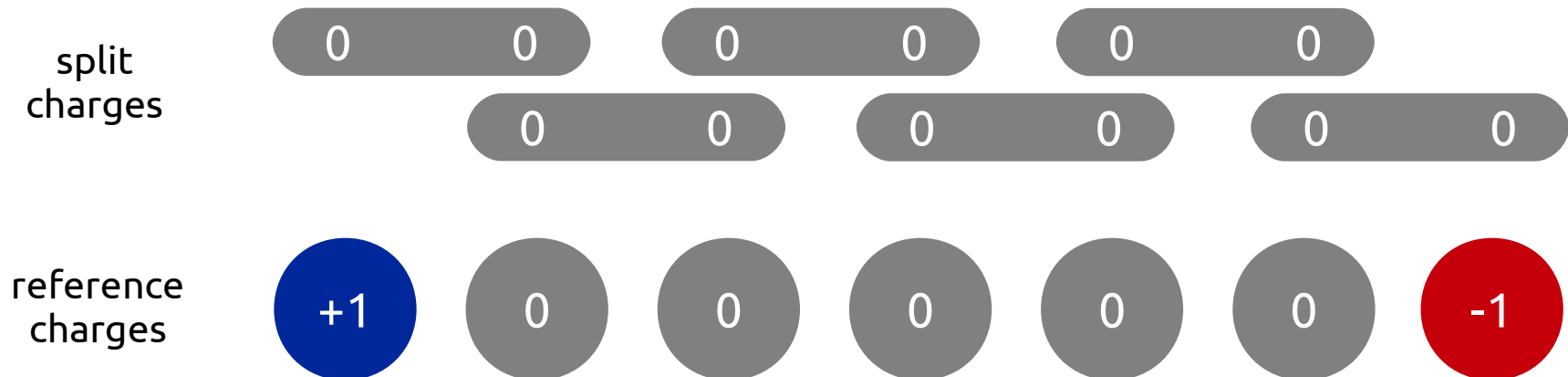
(paper in preparation)

Penta Alanine Benchmark

Charged molecules (and zwitter ions) are problematic.



Better choice (for transferable parameters): $q = Tp + q_0$



→ More parameters. How to model the reference charges?

(paper in preparation)

Overview

Introduction

Exploring Chemical Compounds
Force Field 101
Making Force Fields

Atoms in Molecules

Partitioning of Electron Density
Iterative Hirshfeld
ESP Fitted charges
Benchmarks

The Hydrogen Bond

AIM Application to FF development
Charge Transfer Matters!

Split Charge Equilibration

Partitioning of Polarization
Split Charges
Benchmarks

Harmonic Models

Force Constants & Rest Values

Conclusion

The Harmonic Model (1)

Mathematical form

$$E_{\text{harm}} = \frac{1}{2} \sum_{i=1}^{N_q} \sum_{j=1}^{N_q} K_{ij} (q_i - q_i^{(0)}) (q_j - q_j^{(0)})$$

where q_i are a series of internal coordinates (distances, angles, ...)

Purpose of the harmonic model

- Reproduce geometries and vibrational frequencies
- To be included in FF model, next to electrostatics and others
- Should represent the covalent interactions

Traditional approach to get parameters

- set $q_i^{(0)}$ to the internal coordinates of a QM/XRD optimized structure
- *Invert* the QM Hessian, i.e. find **a** solution for

$$\frac{\partial^2 E^{(\text{QM})}}{\partial x_k \partial x_l} = H_{kl} = \sum_{i=1}^{N_q} \sum_{j=1}^{N_q} K_{ij} \frac{\partial q_i}{\partial x_k} \frac{\partial q_j}{\partial x_l} \quad \text{or} \quad H = J^T K J$$

Ermoshin et al, *Chem. Phys.* **1996**, 202, p53

Problems with the traditional approach

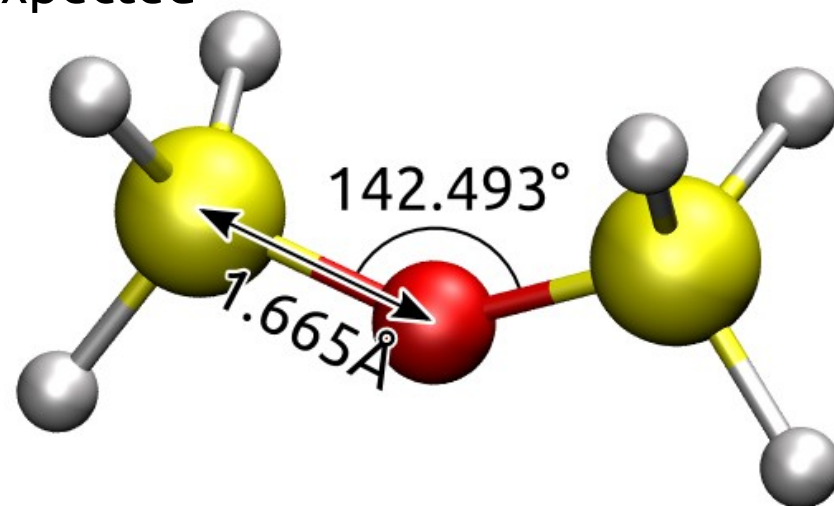
- * **Not optimal:** which Jacobian inverse is best?
- * **Not optimal:** it is common to neglect many cross terms
- * One assumes that other terms (electrostatics, ...) do not affect geometry
⇒ requires 1-2, 1-3 or 1-4 exclusion rules

However ...

- * 1-2, 1-3 and 1-4 electrostatics are real
- * For EEM or SQE, all electrostatic terms must be present

Calibration of valence parameters must be reinvented!

- * Turns out to be more complicated than expected
- * Demo with a 2T silica cluster
- * QM training data
 - PBE/DGTZVP
 - geometry optimization
 - frequency computation



Valence Part (without cross terms)

Harmonic bond stretch terms: **Si-H; Si-O**

$$\frac{1}{2}K_b \left(d - d_b^{(0)} \right)^2$$

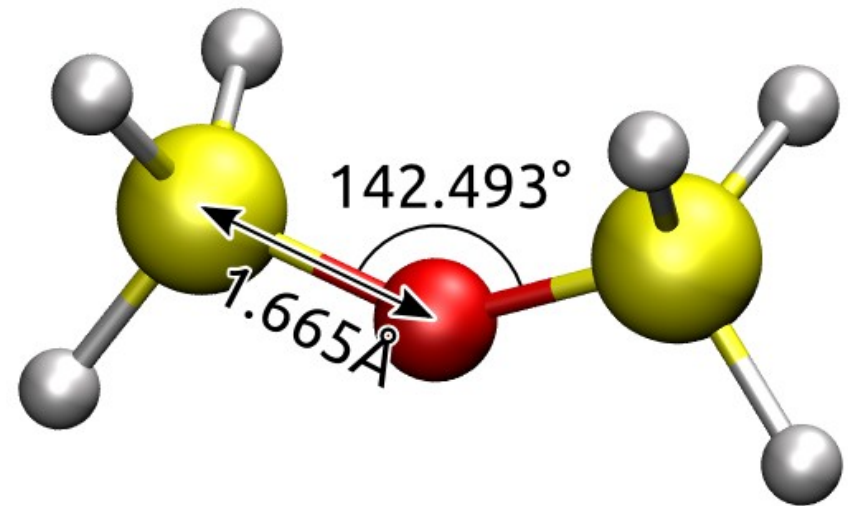
Harmonic Cosine terms: **H-Si-H, H-Si-O, Si-O-Si**

$$\frac{1}{2}K_a \left(\cos(\alpha) - \cos \left(\alpha_a^{(0)} \right) \right)^2$$

Electrostatics

Fixed charge transfer: **Si-H, Si-O**

$$\frac{1}{2} \sum_{i \neq j} \frac{q_i q_j}{r_{ij}}$$



(with or without exclusion rules, both cases will be considered)

Charge transfers are ESP-fitted: $q_{\text{Si} \rightarrow \text{H}} = -0.118e$ $q_{\text{Si} \rightarrow \text{O}} = -0.229e$

Traditional approach

- * Force constants
 - Use Moore-Penrose inverse of Jacobian.
 - Neglect cross terms.
- * Rest values
 - Optimal internal coordinates
- * 1-3 exclusion lists (bonds and bends are excluded)

Fitted parameters

- * Numbers in **red** are compromises
- * Many significant off-diagonal K_{ij} (mainly H-Si-H)

	Force constant	Rest value
Si-H	1636.7 kJ mol ⁻¹ Å ⁻²	1.50 Å
Si-O	2876.1 kJ mol ⁻¹ Å ⁻²	1.67 Å
H-Si-H	344.0 kJ mol ⁻¹	109.38 °
H-Si-O	265.2 kJ mol ⁻¹	109.56 °
Si-O-Si	159.0 kJ mol ⁻¹	142.49 °

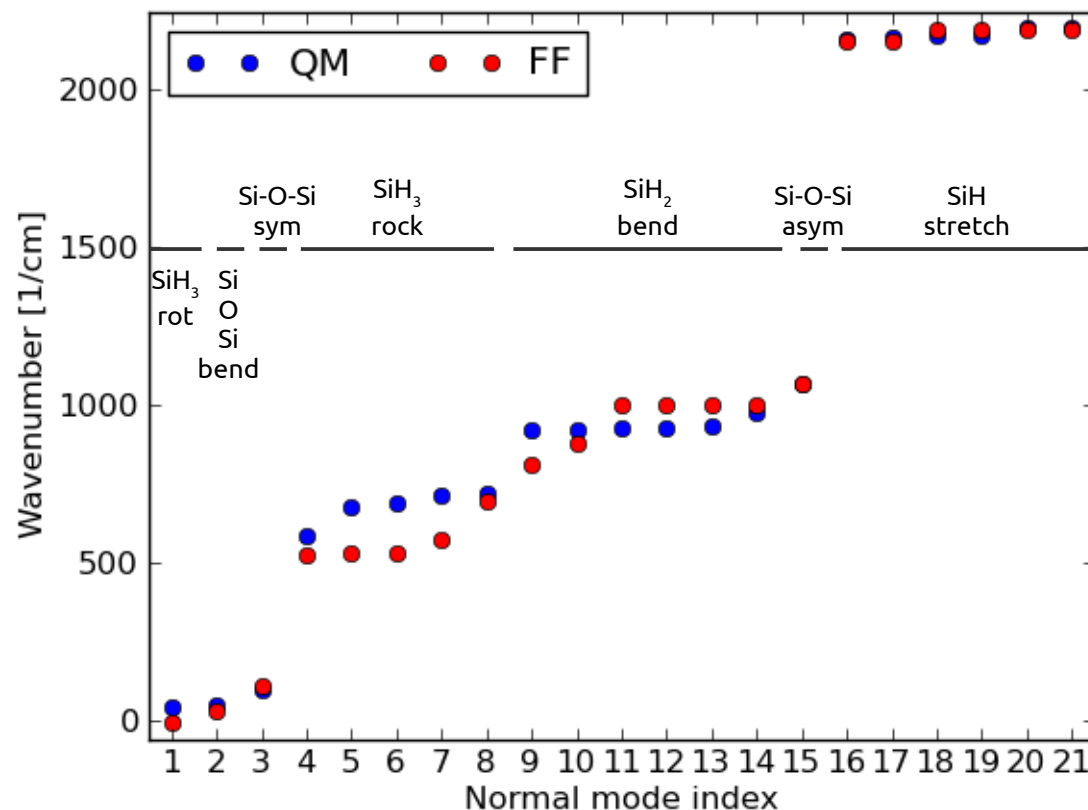
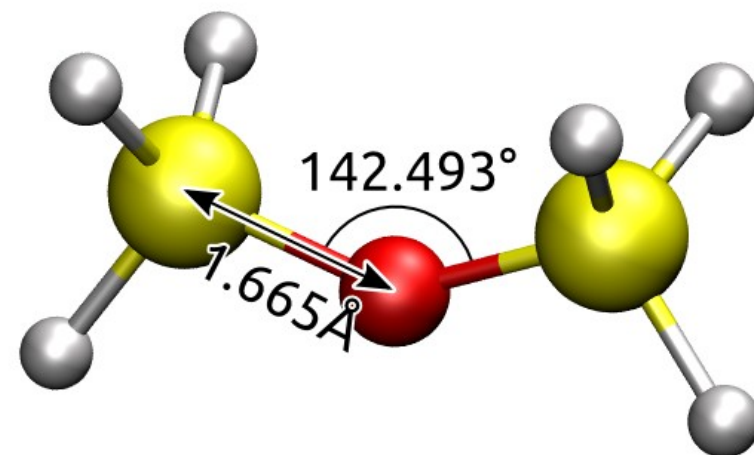
Results for Method A

Optimized internal coordinates

Si-O-Si angle 125.055 °
Si-O bond 1.655 Å

Comments

- Si-O-Si angle is wrong due to 1-4 and 1-5 electrostatics
- FF Spectrum is reasonable
- Only way to improve is "trial and error"



Least-squares approach

No Exclusion lists

1) Guess force constants: 'approximate' least-squares fit

$$\frac{\partial^2 E^{(\text{QM})}}{\partial x_k \partial x_l} = \frac{\partial^2 E^{(\text{ES})}}{\partial x_k \partial x_l} + \sum_{i=1}^{N_q} K_{ii} \frac{\partial q_i}{\partial x_k} \frac{\partial q_i}{\partial x_l}$$

2) Rest values: least squares fit of (zero gradient)

$$0 = \frac{\partial E^{(\text{QM})}}{\partial x_k} = \frac{\partial E^{(\text{ES})}}{\partial x_k} + \sum_{i=1}^{N_q} K_{ii} (q_i - q_i^{(o)}) \frac{\partial q_i}{\partial x_k}$$

3) Redo force constants: 'correct' least-squares fit

$$\frac{\partial^2 E^{(\text{QM})}}{\partial x_k \partial x_l} = \frac{\partial^2 E^{(\text{ES})}}{\partial x_k \partial x_l} + \sum_{i=1}^{N_q} K_{ii} \frac{\partial q_i}{\partial x_k} \frac{\partial q_i}{\partial x_l} + \sum_{i=1}^{N_q} K_{ii} (q_i - q_i^{(o)}) \frac{\partial^2 q_i}{\partial x_k \partial x_l}$$

Least-squares approach

After step 2:

	Force constant	Rest value
Si-H	1664.3 kJ mol ⁻¹ Å ⁻²	1.51 Å
Si-O	2713.8 kJ mol ⁻¹ Å ⁻²	1.70 Å
H-Si-H	287.9 kJ mol ⁻¹	140.69 °
H-Si-O	374.7 kJ mol ⁻¹	131.95 °
Si-O-Si	24.6 kJ mol ⁻¹	23.12 °

	Traditional f.c.'s
Si-O	1636.7
Si-O	2876.1
H-Si-H	344.0
H-Si-O	265.2
Si-O-Si	159.0

- * Poor rest values for angles
- * Bending f.c.'s change a lot
- * Bond rest values increase to compensate electrostatics

Least-squares approach

After step 3:

	Force constant	Rest value
Si-H	1659.2 kJ mol ⁻¹ Å ⁻²	1.51 Å
Si-O	2709.9 kJ mol ⁻¹ Å ⁻²	1.70 Å
H-Si-H	172.6 kJ mol ⁻¹	140.69 °
H-Si-O	250.3 kJ mol ⁻¹	131.95 °
Si-O-Si	0.0 kJ mol ⁻¹	23.12 °

	Traditional f.c.'s
Si-O	1636.7
Si-O	2876.1
H-Si-H	344.0
H-Si-O	265.2
Si-O-Si	159.0

- * This step is supposed to fine-tune the f.c.'s, but...
- * Bending f.c.'s values do not become more realistic.

In general: parameters are not encouraging

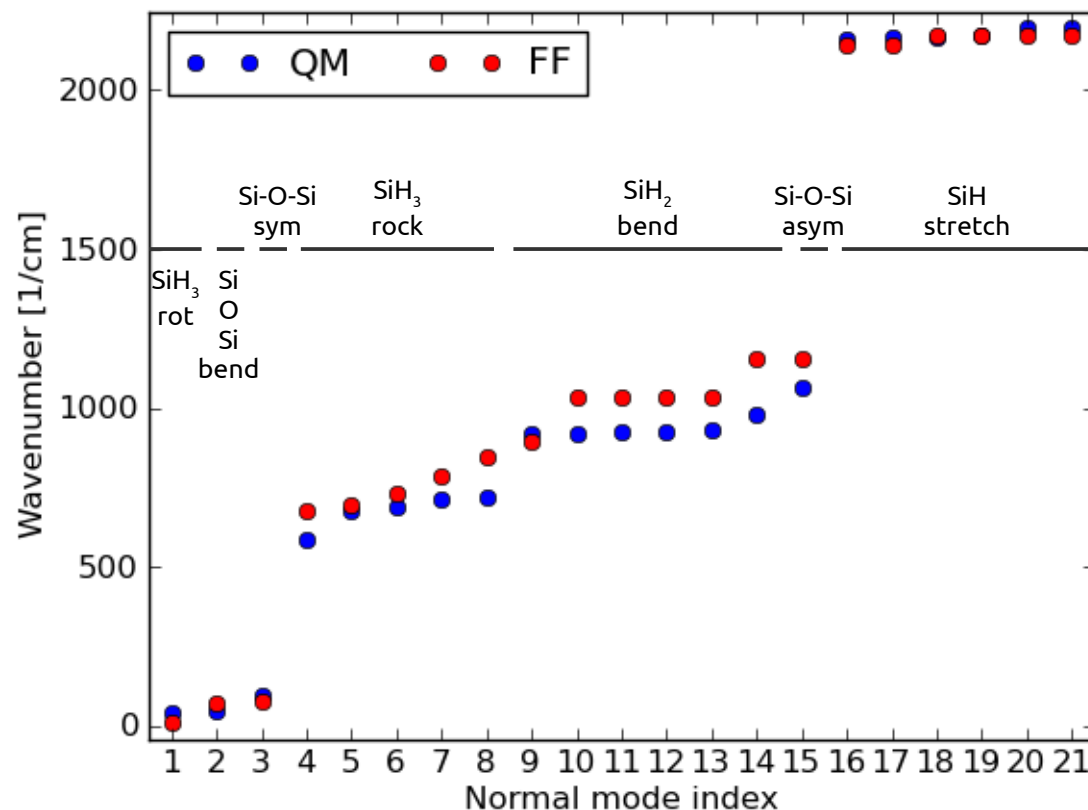
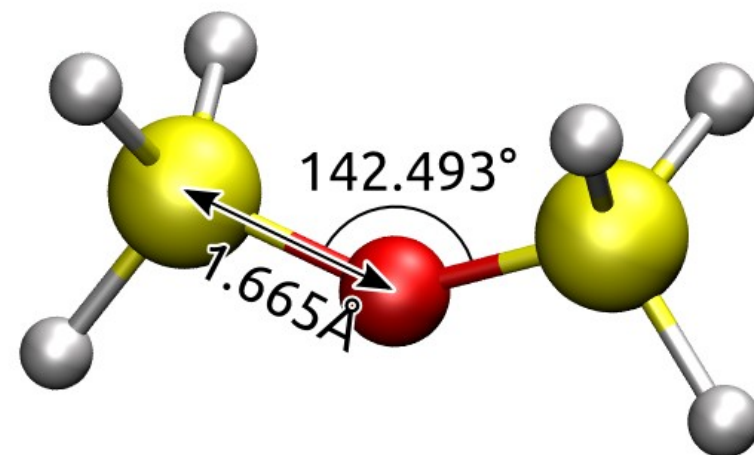
Results for Method B(2)

Optimized internal coordinates

Si-O-Si angle 108.204 °
Si-O bond 1.668 Å

Comments

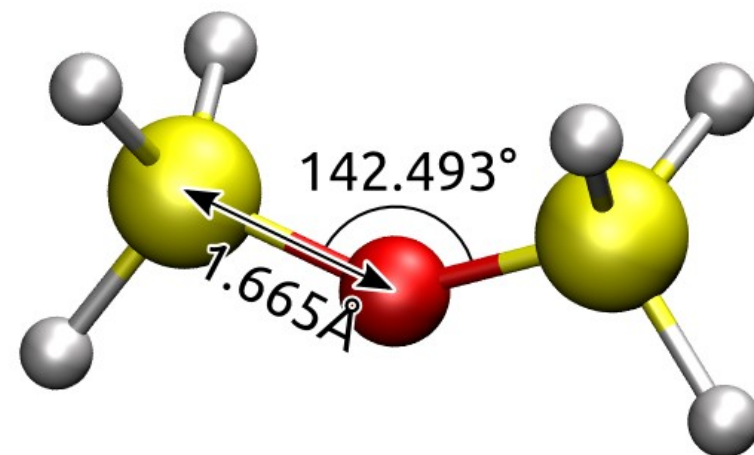
- Si-O-Si angle is wrong because a small error in the gradient along a low modes causes a large error in geometry
- Bond length is better
- FF Spectrum is worse



Results for Method B(3)

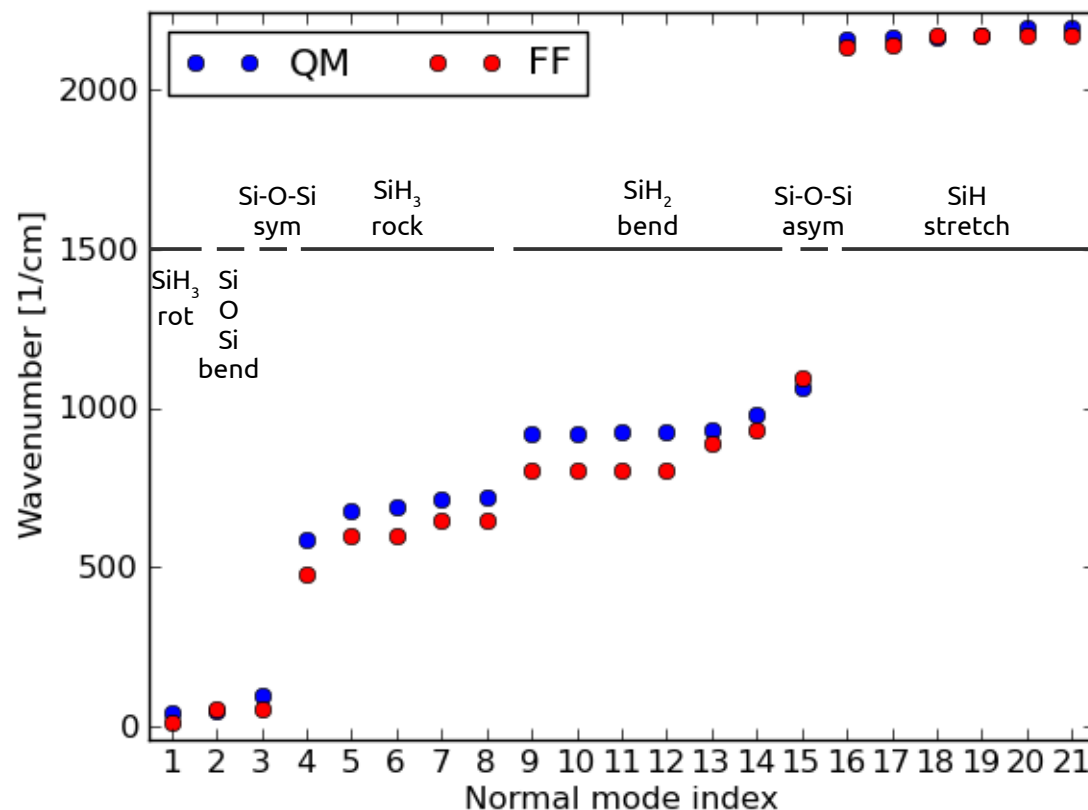
Optimized internal coordinates

Si-O-Si angle 180.00 °
Si-O bond 1.666 Å



Comments

- Pfff...



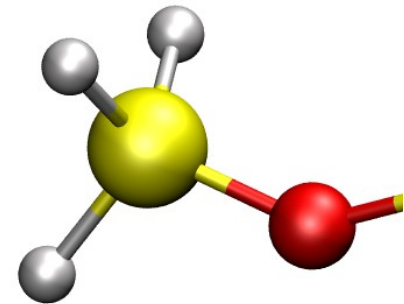
Improved least-squares approach

- * Still no exclusion lists
- * Try to fix all the issues encountered in B

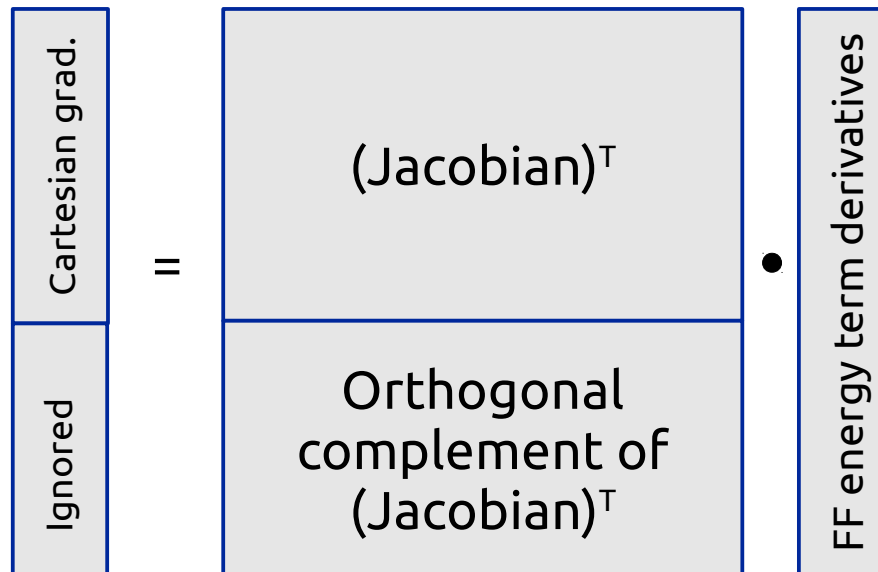
Problem 1: the computation of the Cartesian gradient is a projection

N_q derivatives of FF energy terms \rightarrow $3N-6$ independent

\Rightarrow rest values for **H-Si-H** and **H-Si-O** are ill-defined



6 angles
5 independent



(paper in preparation)

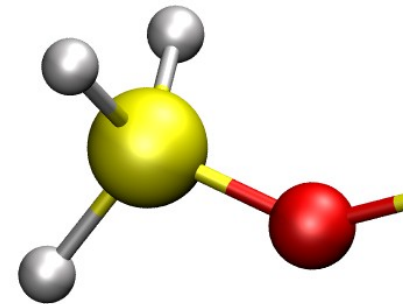
Improved least-squares approach

- * Still no exclusion lists
- * Try to fix all the issues encountered in B

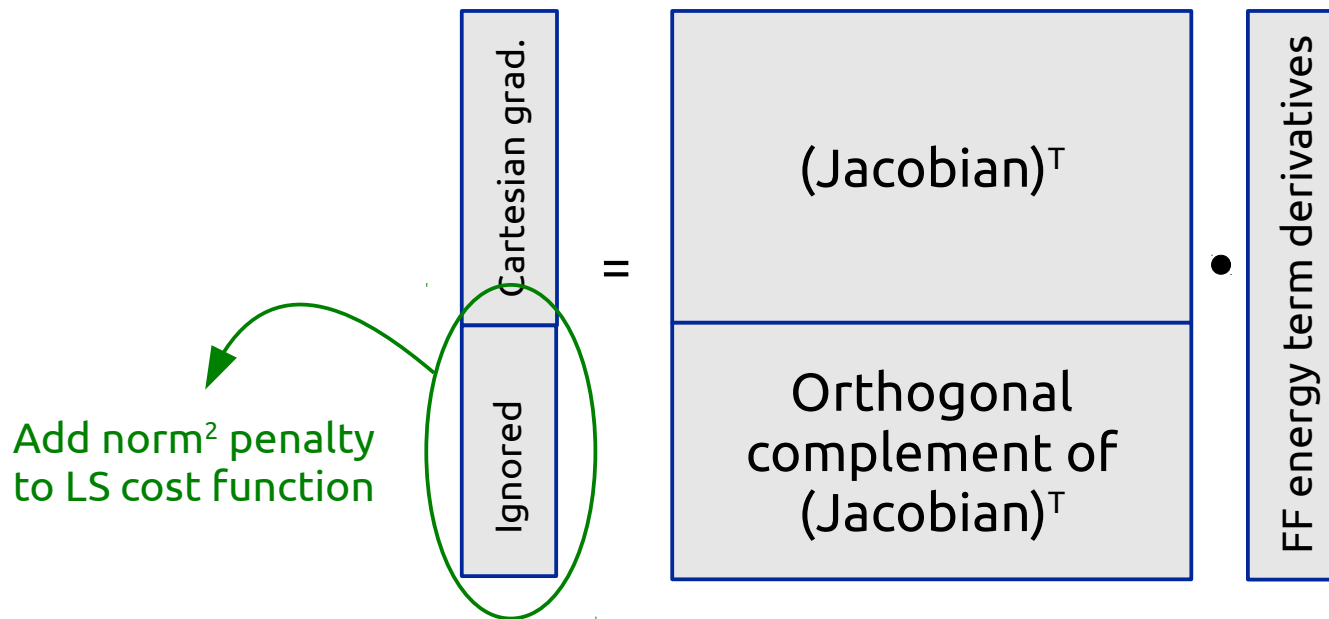
Problem 1: the computation of the Cartesian gradient is a projection

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\Rightarrow rest values for **H-Si-H** and **H-Si-O** are ill-defined



6 angles
5 independent



(paper in preparation)

Improved least-squares approach

- * Still no exclusion lists
- * Try to fix all the issues encountered in B

Problem 2: gradient along low modes should be more accurate

Solution: use a transformed gradient in LS fit, as follows

$$\Delta X = H_{\text{QM}}^{-1} G$$

Related transformation for Hessian fit:

$$H' = H_{\text{QM}}^{-\frac{1}{2}} H H_{\text{QM}}^{-\frac{1}{2}}$$

(paper in preparation)

Improved least-squares approach

- * Still no exclusion lists
- * Try to fix all the issues encountered in B

Problem 3: not entirely optimal

The last step (3) should optimize all parameters with one total cost function

$$X_{\text{total}} = \frac{X_{\text{grad}} + \lambda_1 X_{\text{hess}}}{1 + \lambda_1} + \lambda_2 X_{\text{penalty}}$$

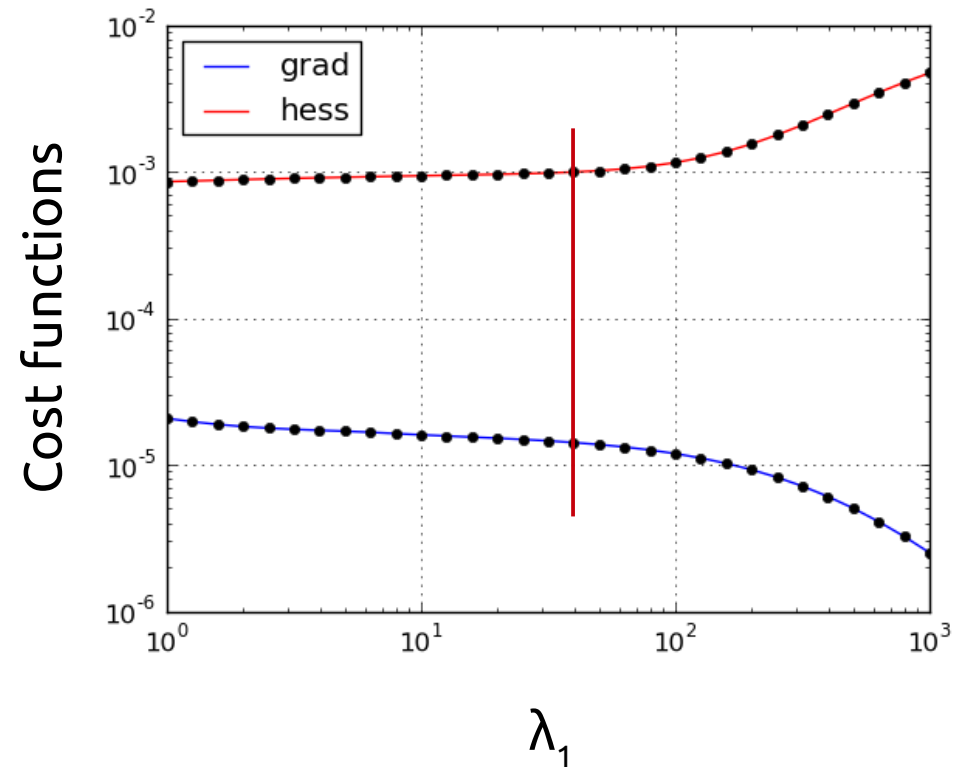
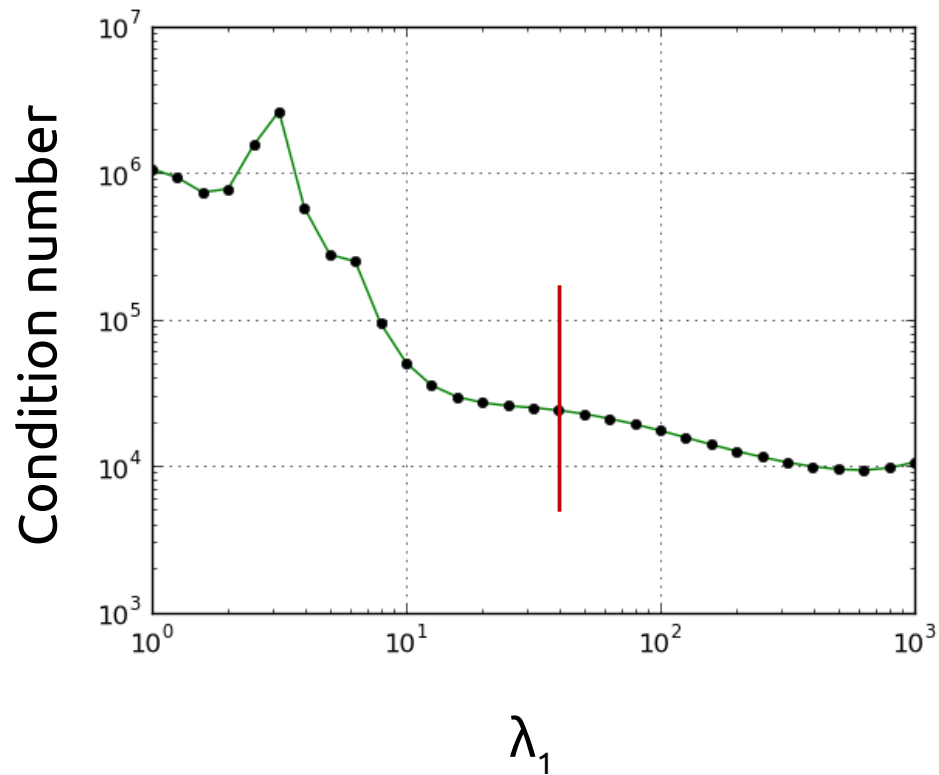
λ_2 can be set relatively high, does not hurt

λ_1 must be scanned over a few orders of magnitude:

- * Good compromise between X_{grad} and X_{hess}
- * Low condition number

(paper in preparation)

Tuning λ_1 in Method C



(paper in preparation)

Parameters for Method C

Improved least-squares approach

After step 3:

	Force constant	Rest value
Si-H	1668.1 kJ mol ⁻¹ Å ⁻²	1.51 Å
Si-O	2468.4 kJ mol ⁻¹ Å ⁻²	1.70 Å
H-Si-H	269.9 kJ mol ⁻¹	110.39 °
H-Si-O	330.1 kJ mol ⁻¹	109.01 °
Si-O-Si	48.4 kJ mol ⁻¹	122.26 °

differ
traditional
values

Penalty

Differs
from
optimized
value

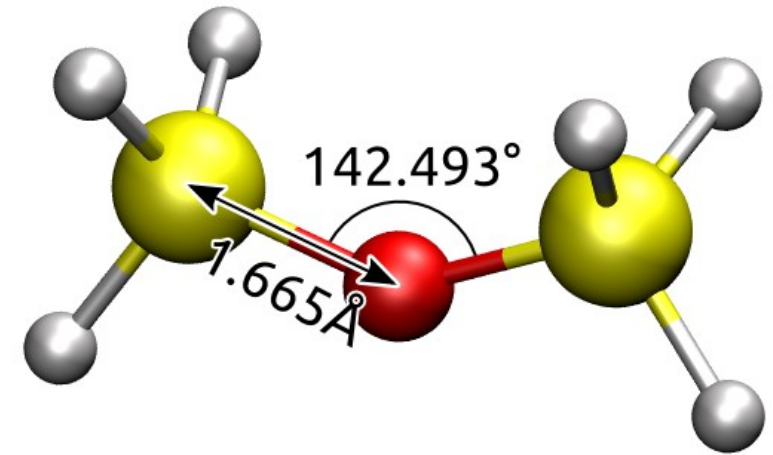
Traditional approach (method A)

	Force constant	Rest value
Si-H	1636.7 kJ mol ⁻¹ Å ⁻²	1.50 Å
Si-O	2876.1 kJ mol ⁻¹ Å ⁻²	1.67 Å
H-Si-H	344.0 kJ mol ⁻¹	109.38 °
H-Si-O	265.2 kJ mol ⁻¹	109.56 °
Si-O-Si	159.0 kJ mol ⁻¹	142.49 °

(paper in preparation)

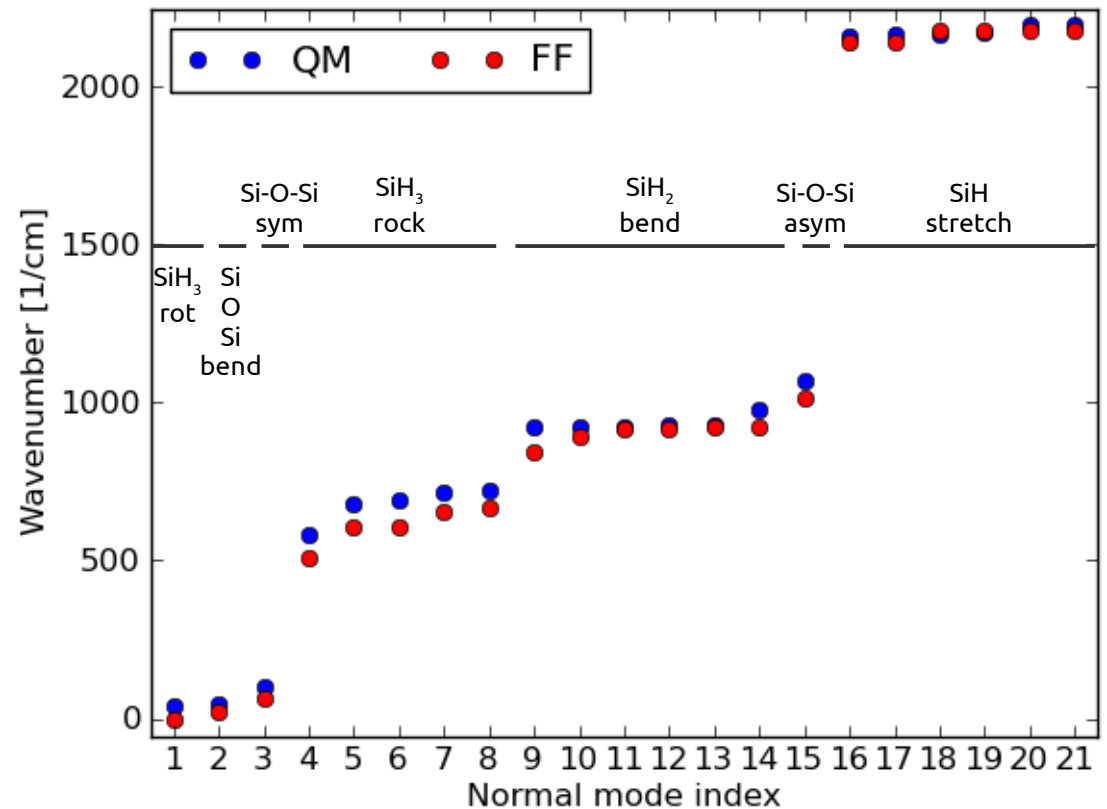
Optimized internal coordinates

Si-O-Si angle 143.146°
Si-O bond 1.663 \AA



Comments

- Best so far
- To improve further:
 1. Cross terms
 2. Improved electrostatics (e.g. atomic dipoles)
 3. Inclusion of repulsion (and dispersion)



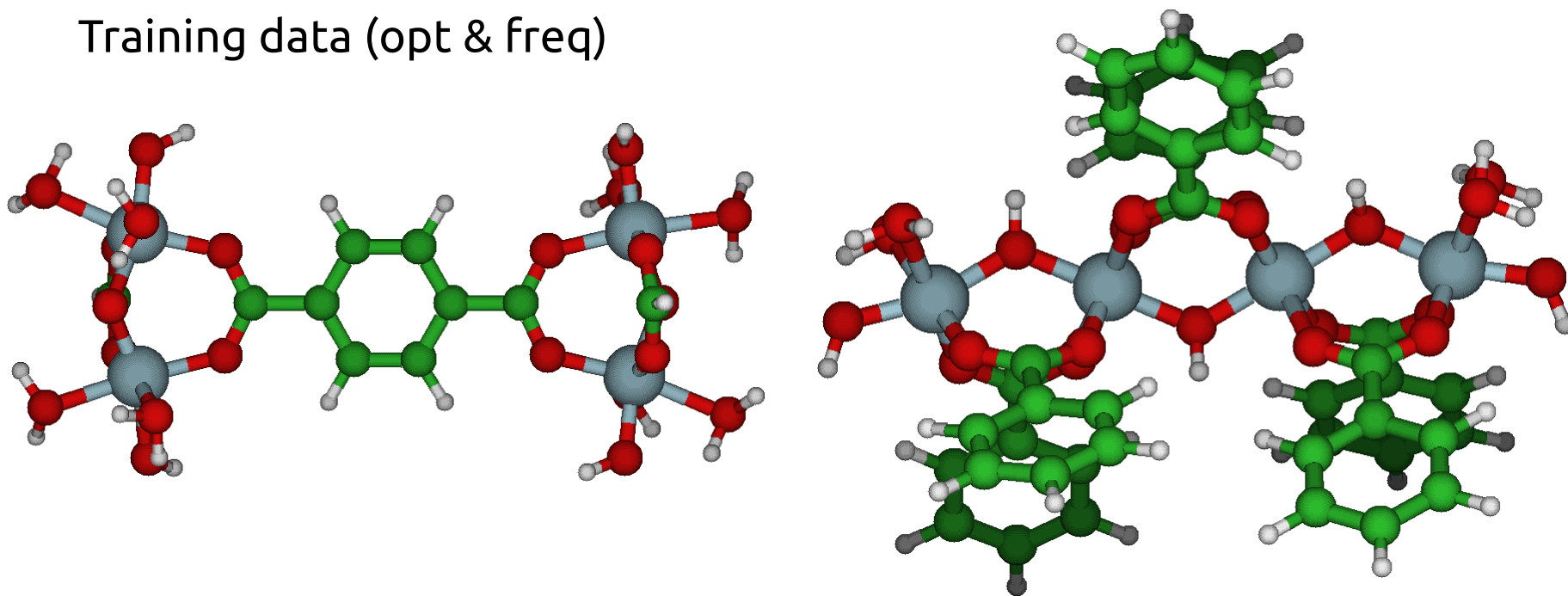
(paper in preparation)

Implemented (FFit2 Code), but not demonstrated in the 2T example

- * Use Hessian & gradient data from a large set of molecules
- * Use charge equilibration models for the electrostatics
- * Account for repulsion/dispersion terms calibrated earlier
- * Possibility to ignore parts of molecules (e.g. weird terminations)

Calibration of MIL-53

Training data (opt & freq)



(paper in preparation)

Cell optimization

* Experimental (HT phase):

a=16.675 Å b=6.609 Å c=12.813 Å

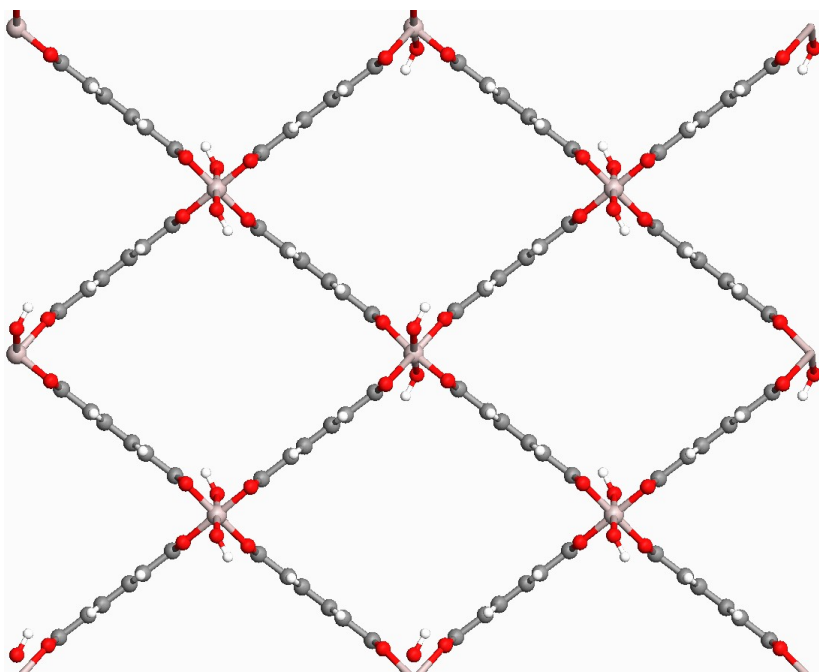
* Cell optimization with FF

a=16.120 Å b=7.417 Å c=13.093 Å

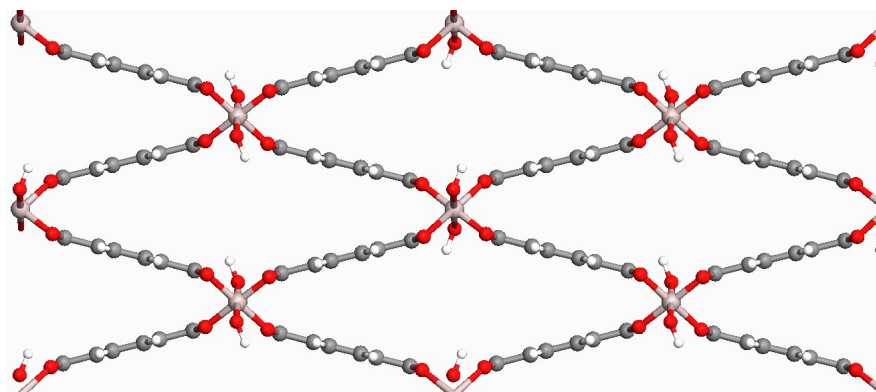
Loiseau et al, *Chem. Eur. J.* **2004**, 10, p1373

Testing sets of dispersion parameters to reproduce breathing

Open form (HT)



Closed form (LT)



(paper in preparation)

Major advances in FF's calibration are on the way!

Atoms in Molecules

Iterative Hirshfeld is a very neat partitioning scheme:

- Robust (conformational, basis)
- ESP is OK
- AIM very similar to isolated atoms

The Hydrogen Bond

AIM → FF parameters without fitting
e.g. Charge Transfer in Hydrogen Bond
(cfr. work Morokuma)

Split Charge Equilibration

e- Linear Response is ruled by charges
SQE fixes polarizability scaling in EEM
SQE only good for locally neutral systems

Harmonic Models

Improved calibration scheme
→ Accurate models,
without manual tweaking

Acknowledgements

Fine colleagues at the CMM ...



Software used: Gaussian09, **HiPart**, **FFit2**, **Yaff**, Python, VMD, POVRay

MIL-53:
L. Vanduyfhuys



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Computing facilities: Ghent University (Stevin Supercomputer Infrastructure).