

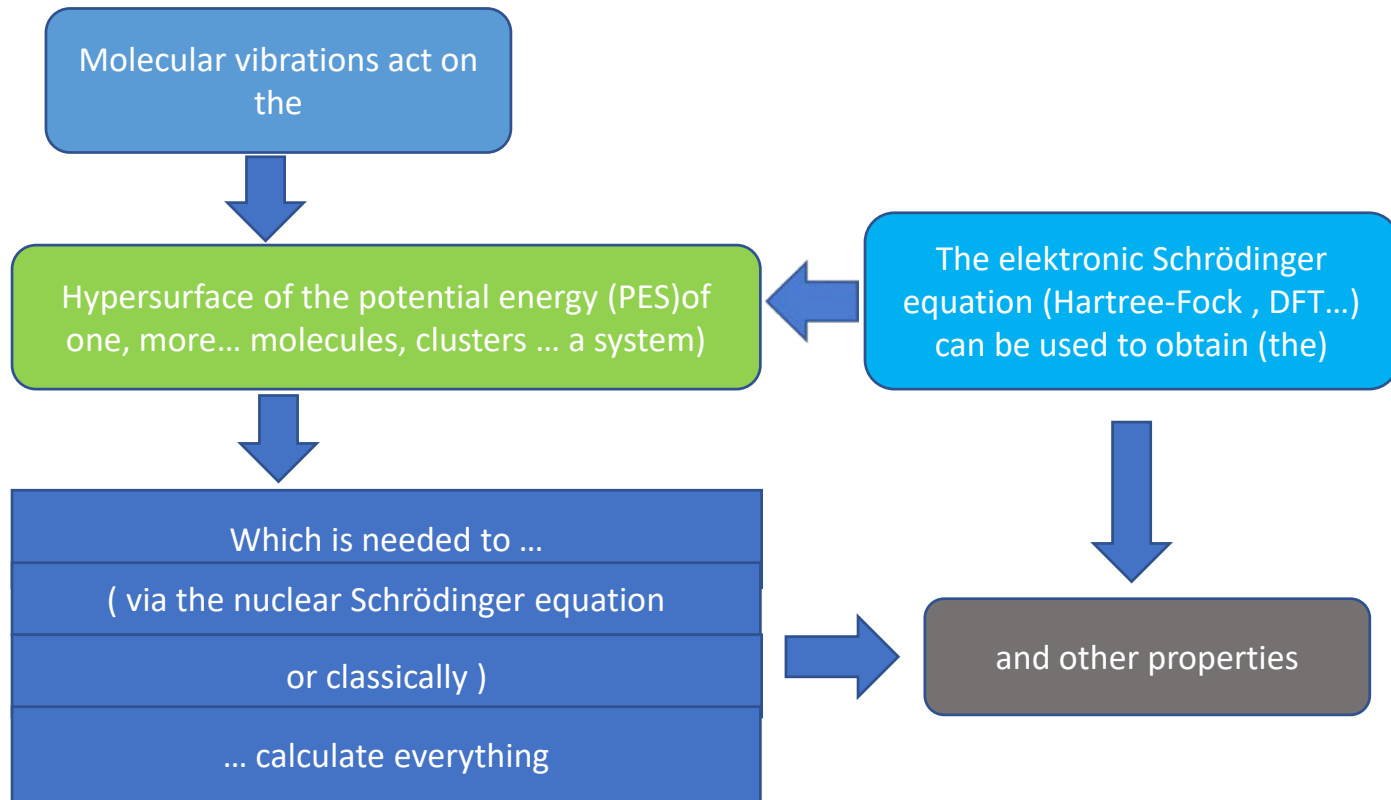
Topic:

Modeling the molecular interactions

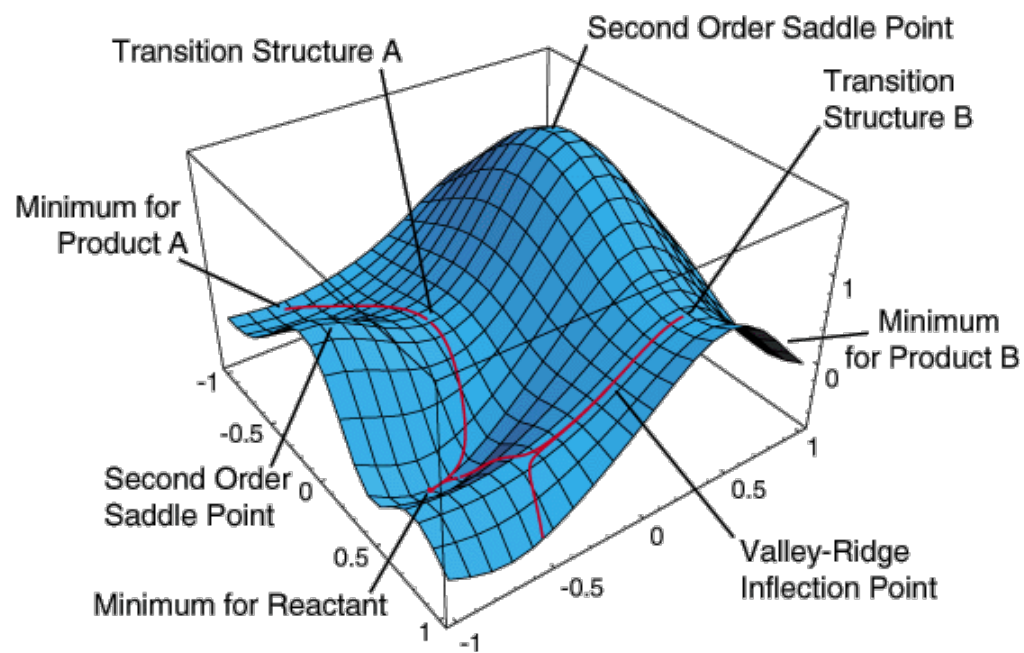
The PES of a molecule (cluster ...)

- Without PES no chemistry and no molecules:
 - every 'Molecule' would be different (its energy would not only depend on its geometry but also on its history).
- The Born-Oppenheimer – approximation states that a PES exists.
- But: one PES for each electronic state.
- The PES allows a eine topographical interpretation of reactions and conversions of molecules.
- Next slide

To remember: Relations



The PES of a molecule (cluster ...)



Modeling the molecular interactions - general

- The potential energy in an N-particle system $V = f(r)$ must be expressed in some tractable way since it needs to be *known* and it needs to be *calculable*.
- It can either be calculated quantum mechanically or be derived from a mathematical expression ('energy formula') that includes some physics.

Modeling the molecular interactions - general

Which energy formulas are useful ?

- For $r_{ij}=0$, E_{ij} should be ∞
 - ... not fulfilled by $\exp(-a r_{ij})$
- For $r_{ij}=\infty$, E_{ij} should be 0
 - ... not fulfilled by a polynomial in r_{ij}
- Normally, E_{ij} should have none or one minimum
 - No minimum \rightarrow repulsion
 - One minimum \rightarrow attraction
- Sometimes but rarely, E_{ij} has one maximum
(and therefore how many minima ?)

Modeling the molecular interactions - general

(discuss Electron-electron interaction
Long- and short range, Energy vs. force ...)

The potential energy in the pair approximation:

In the pair approximation one assumes that the total energy is a sum of pairwise interactions. For example, if E_{ij} is set to be a sum of a_k/r^k terms:

- **The total energy is then:**
(*ij* ... pairs of molecules
lm ... interaction centers in pair *ij*
k ... term in sum over $1/r$ powers)

$$V_{tot} = \sum_{ij} \sum_{lm} \sum_k \frac{a_k}{r_{lm}^k}$$

The potential Energy V

Functional forms:

Sutmann chapter 2

- The pair approximation needs distances as input:

Angular

→

Radial

$$V = \sum_{ij} f_{kl}(C_{ij})$$

$$V = \sum_{ij} \sum_{kl} f(\mathbf{r}_{kl})$$

$$f(\mathbf{r}_{ij}) = \sum_{k=1}^n \frac{a_k}{r_{ij}^k}$$

Modeling the molecular interactions

Deficiencies of the pair approximation:

The pair approximation is a truncation of the energy expression:

$$E=f(i)+f(ij)+f(ijk)+f(ijkl)\dots$$

the pair term is always the most important ...

... but unfortunately the series converges slowly

- Polarisability normally what causes the largest deviation from the pair approximation
- Charge transfer
- Directional effects (bonds)

Polarisability concepts, examples for going beyond the pair approximation, ...

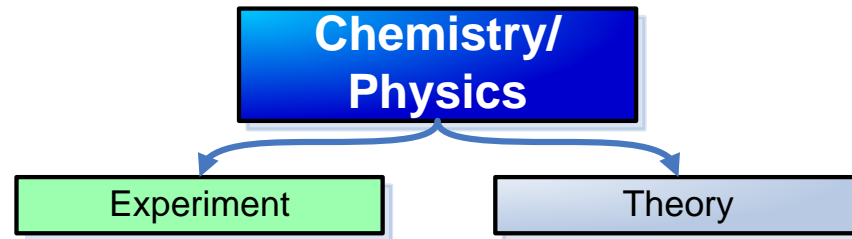
Discussion: polarisation

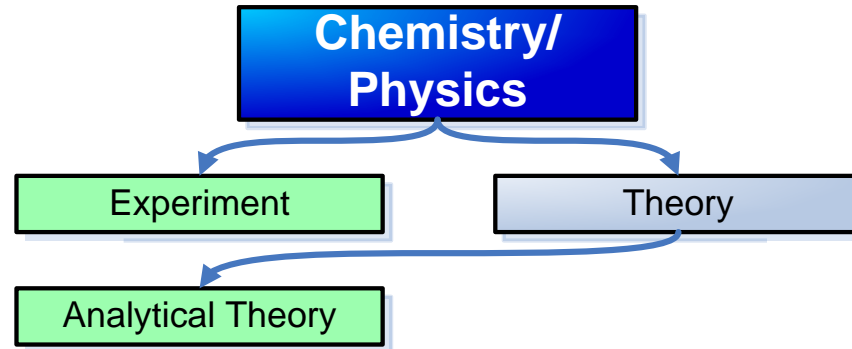
Potential Energy Functions II: step-by-step

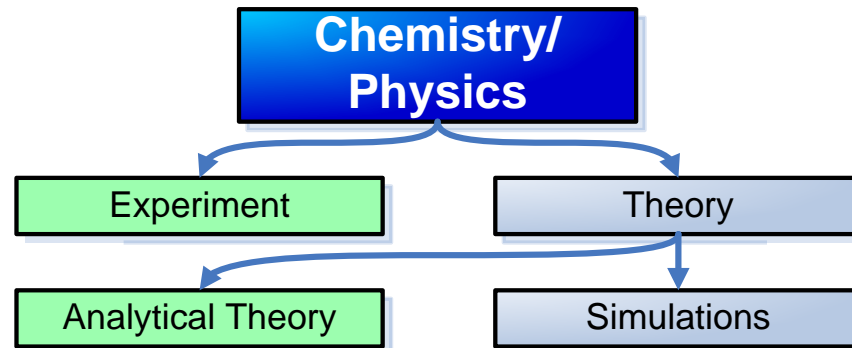
Potential functions

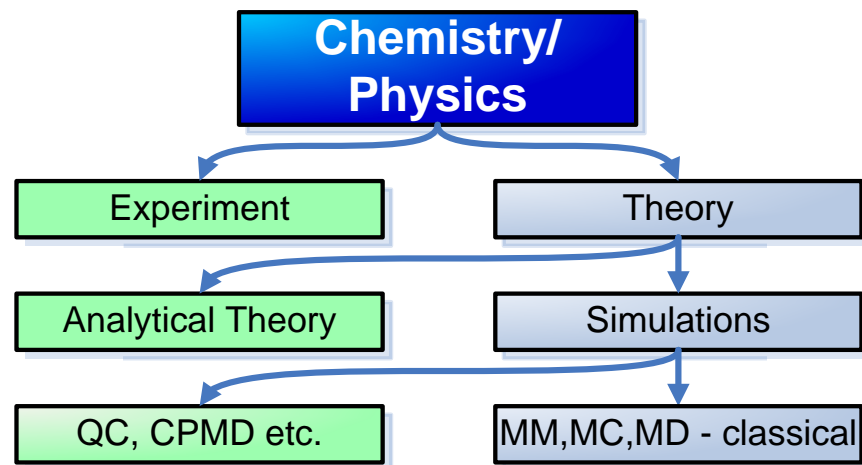
(1) What it is about ...

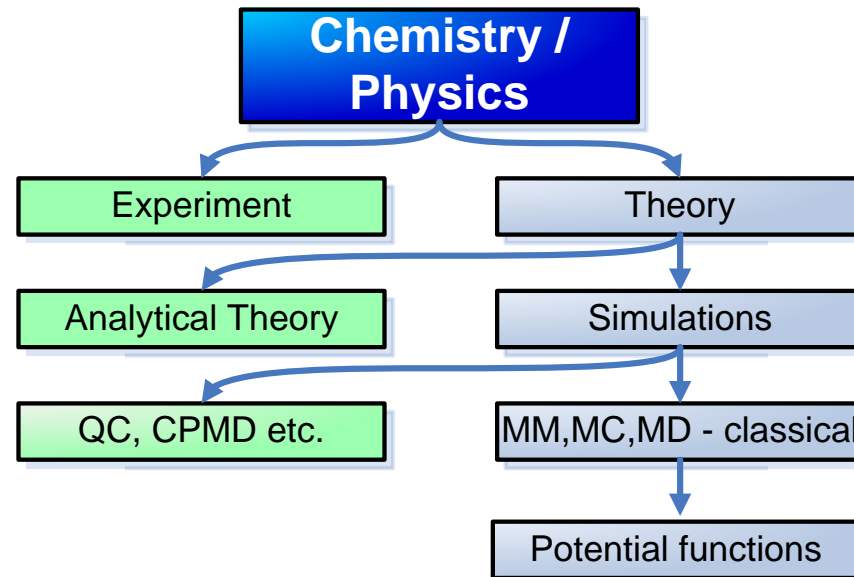
**Chemistry/
Physics**



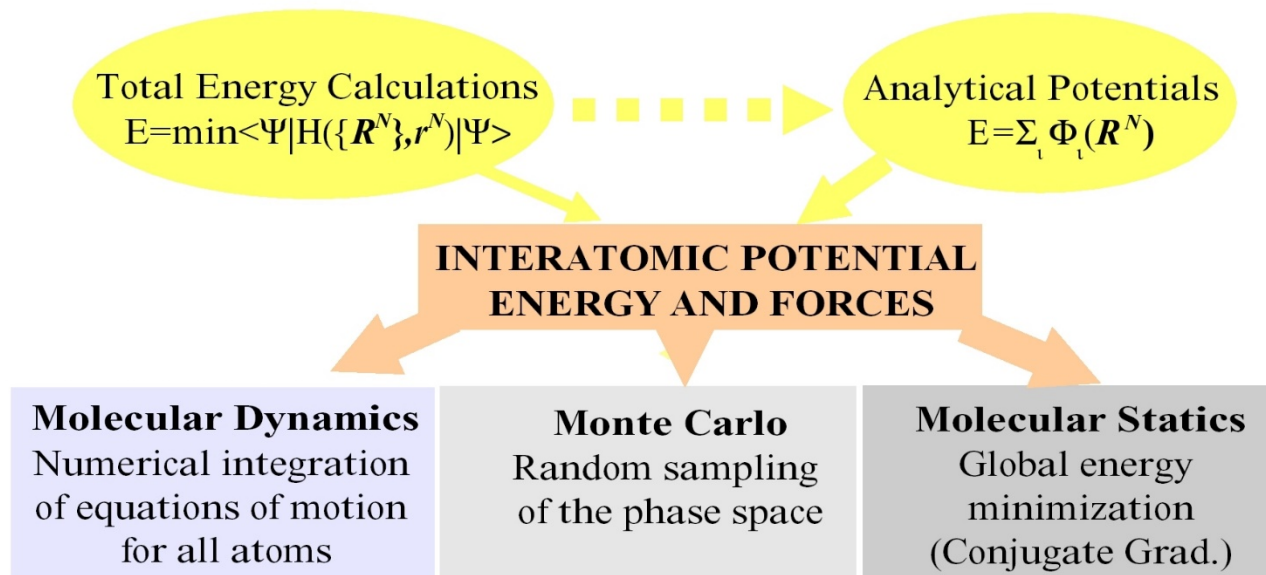


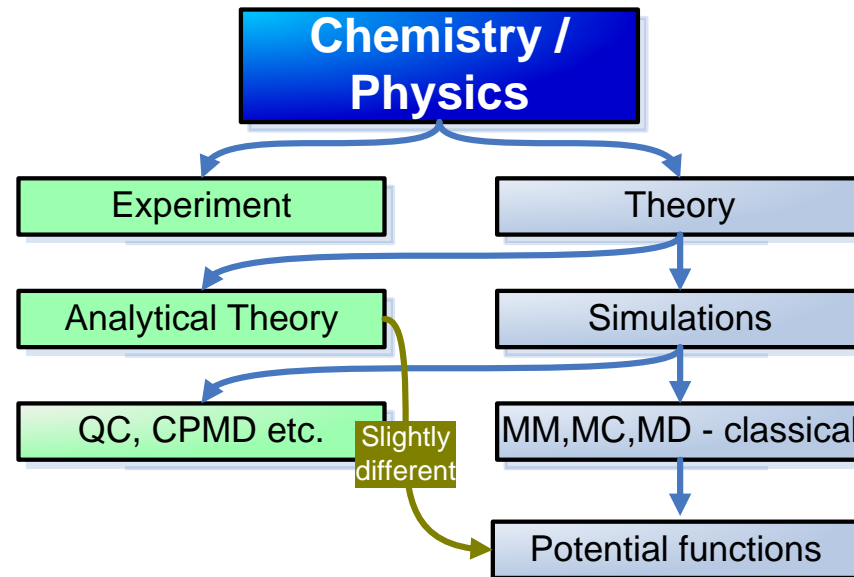






Atomistic Computer Simulation Methods





Potential functions

An analytical potential function is an expression for the total energy of a system as function of the atomic coordinates.

Potential functions

Why does one need them ?
(in the context of molecular simulations)

Properties via Simulations ?

- A system consists of n particles. (Atoms, molecules, ions ...).
Their behavior results from their electronic structure which governs the interaction between them (their attraction and repulsion).
- A potential energy function E which is a scalar function of the coordinates of the particles is containing all this in is containing all information about these interactions.
- How can one calculate macroscopic properties from E ?

(Macroscopic properties could be, for example:

- the pressure $p=f(V)$,
- average geometries at a given temperature or
- (vibrational, electronic) spectra ?

- This is normally not possible in a simple way.
- One needs MD simulations because of the lack of analytical formulas.
There are no formulas like ' $p=nRT / V$ ' for properties of real systems.

- Therefore, there are two problems
- The first one is the **potential energy function** $E_{\text{pot}}=f(X)$.
- *Experience (of the last 30 years) has shown that quantum chemistry is a powerful tool to tackle the problem of the potential energy function.*
- The second problem is the **calculation of properties** if $E_{\text{pot}}=f(X)$ is known. Computer simulations (especially MD) can do this numerically.

- **(2)**

The general scheme of fitting quantum chemically derived data to analytical expressions:

How to get a potential function

Potential functions

i.E. Lennard-Jones

$$E = E_{LJ} + E_{qq}$$

$$E_{LJ}(r) = \left\{ 4\varepsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right] \right\}$$

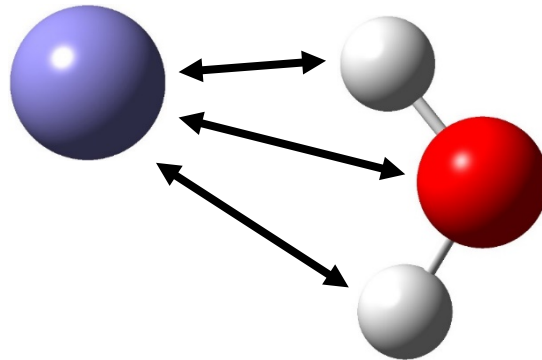
$$E_{qq}(r) = \frac{q_i q_j}{r_{ij}}$$

r is the distance between 2 atoms

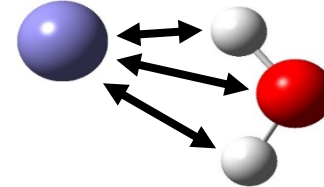
Potential functions

$$E(r_{ij}) = \left\{ 4\epsilon \left[\left(\frac{\sigma}{r_{ij}} \right)^{12} - \left(\frac{\sigma}{r_{ij}} \right)^6 \right] \right\} + \frac{q_i q_j}{r_{ij}}$$

For a water molecule interacting with i.e. an atomic ion I^+ there would be 3 such terms:



We can write the formula slightly different so that it becomes a polynomial in $1/r$:



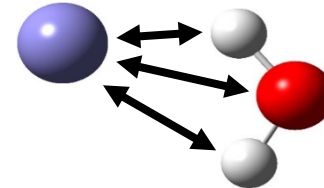
$$E(r_{ij}) = \frac{A}{r_{ij}^{12}} + \frac{B}{r_{ij}^6} + \frac{q_i q_j}{r_{ij}}$$

We need different parameters A,B for the interaction between I^+-O and I^+-H but the two I^+-H interactions have the same parameters A,B.
The total interaction energy is:

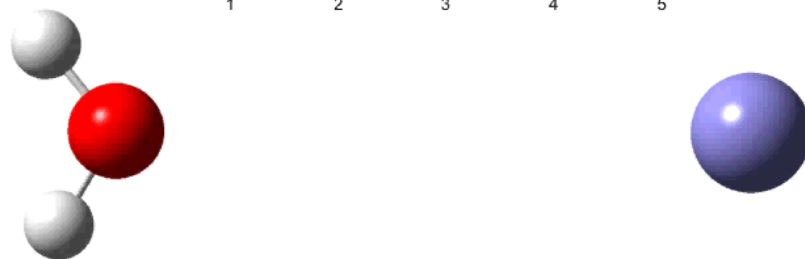
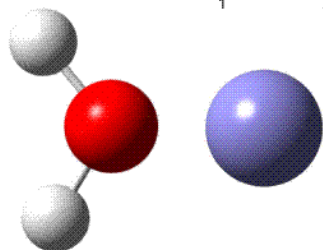
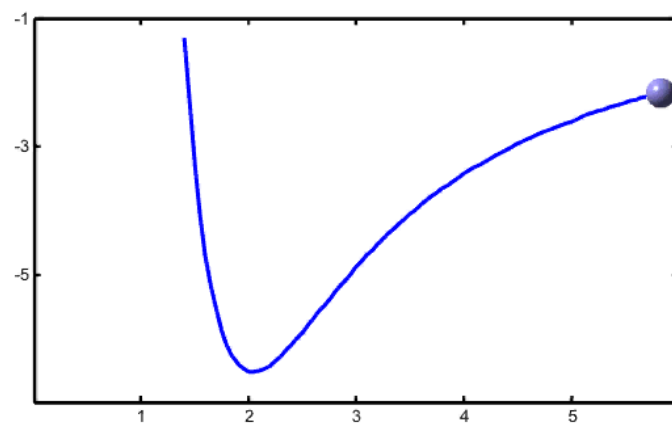
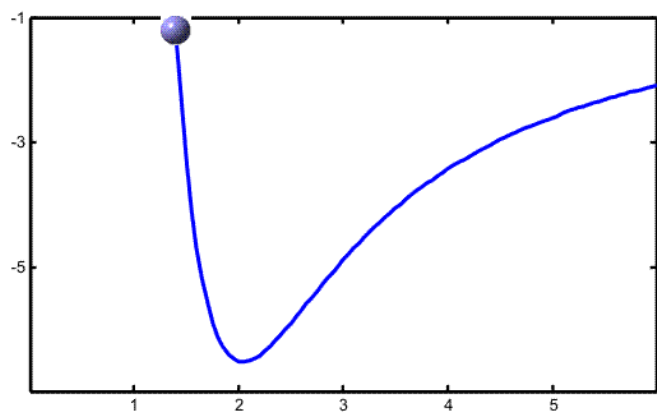
$$\begin{aligned} E_{I^+-W} = & \frac{A_{I^+-O}}{r_{I^+-O}^{12}} + \frac{B_{I^+-O}}{r_{I^+-O}^6} + \frac{q_{I^+} q_O}{r_{I^+-O}} + \\ & + \frac{A_{I^+-H}}{r_{I^+-H^1}^{12}} + \frac{B_{I^+-H}}{r_{I^+-H^1}^6} + \frac{q_{I^+} q_H}{r_{I^+-H^1}} + \\ & + \frac{A_{I^+-H}}{r_{I^+-H^2}^{12}} + \frac{B_{I^+-H}}{r_{I^+-H^2}^6} + \frac{q_{I^+} q_H}{r_{I^+-H^2}} \end{aligned}$$

We have now the 7 parameters

$$A_{I^+-O}, B_{I^+-O}, A_{I^+-H}, B_{I^+-H}, q_{I^+}, q_O, q_H$$



- In general, a set of atoms in a certain environment (like H in H₂O which, for example, is different from H of CH₄) is called a 'class'. In this example, atoms and classes are the same.
- The q – parameters are *class-specific* parameters for which good values often can be calculated via so-called 'population analysis', (atomic partial charge analysis), a standard quantum chemical method.
- The A and B – parameters are *class-pair specific* parameters that can be calculated via fitting to quantum chemical energies.



To consider:

- Many orientations and distances
- Scanning a 3-dimensional (in the example) or 6-dimensional (in the general case) space.
- Computationally demanding
- The energy is linear in the A, B – parameters in a polynomial expression. Expressions like ' $\exp(-C r)$ ' are also commonly used, where this is not the case.

(3) The (non)additivity of interactions

- A formula like
$$E_{I^+-H_2O} = \sum_{O,H,H} \frac{A}{r^{12}} + \frac{B}{r^6} + \frac{q_i q_j}{r}$$

describes the interaction between the ion and water. If we have many water molecules, can we simply say:

$$E_{total} = \sum_{(H_2O_i)} \sum_{O_i, H_i, H_i} \frac{A}{r^{12}} + \frac{B}{r^6} + \frac{q_i q_j}{r} + E_{H_2O} \quad ?$$

- If we do so, we use the so-called pair approximation.

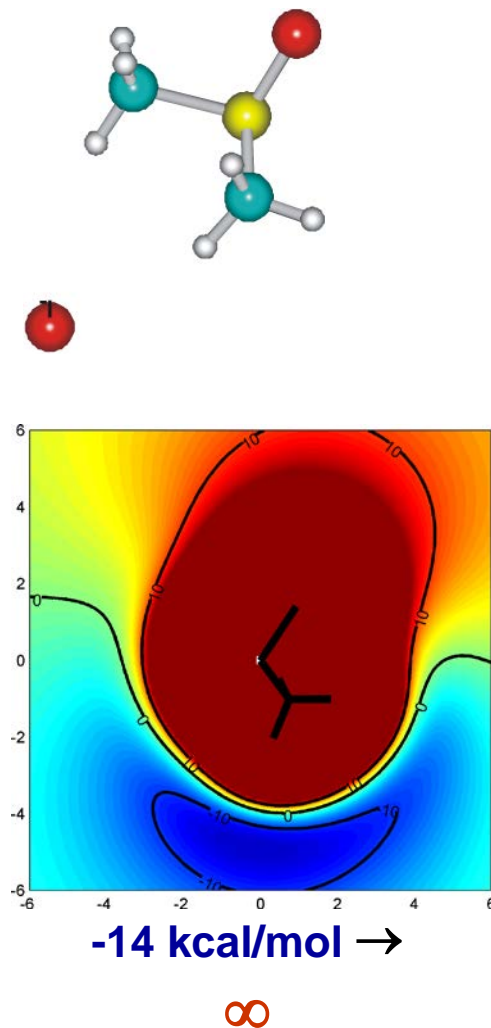
(4)

An example (where the pair approximation is used):

Interaction between Be^{2+} / I^-
and DMSO:

After the construction of the potential energy function, it must be checked.

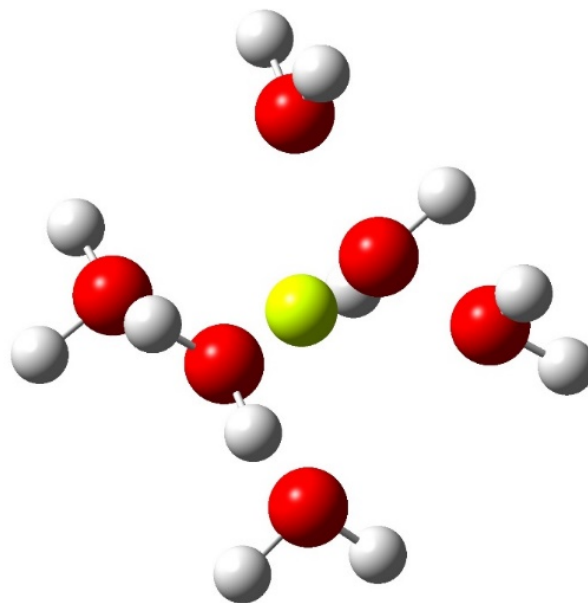
- Example for I^- :



(5)

Another example (beyond
the pair approximation)

The pair approximation is
often not good enough,
especially for
ion-ligand
interactions:

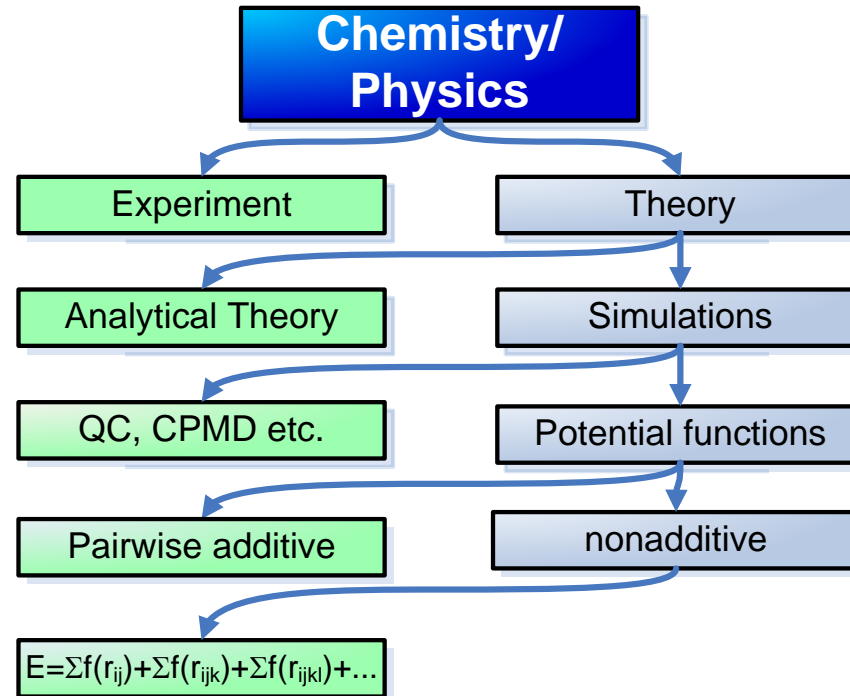


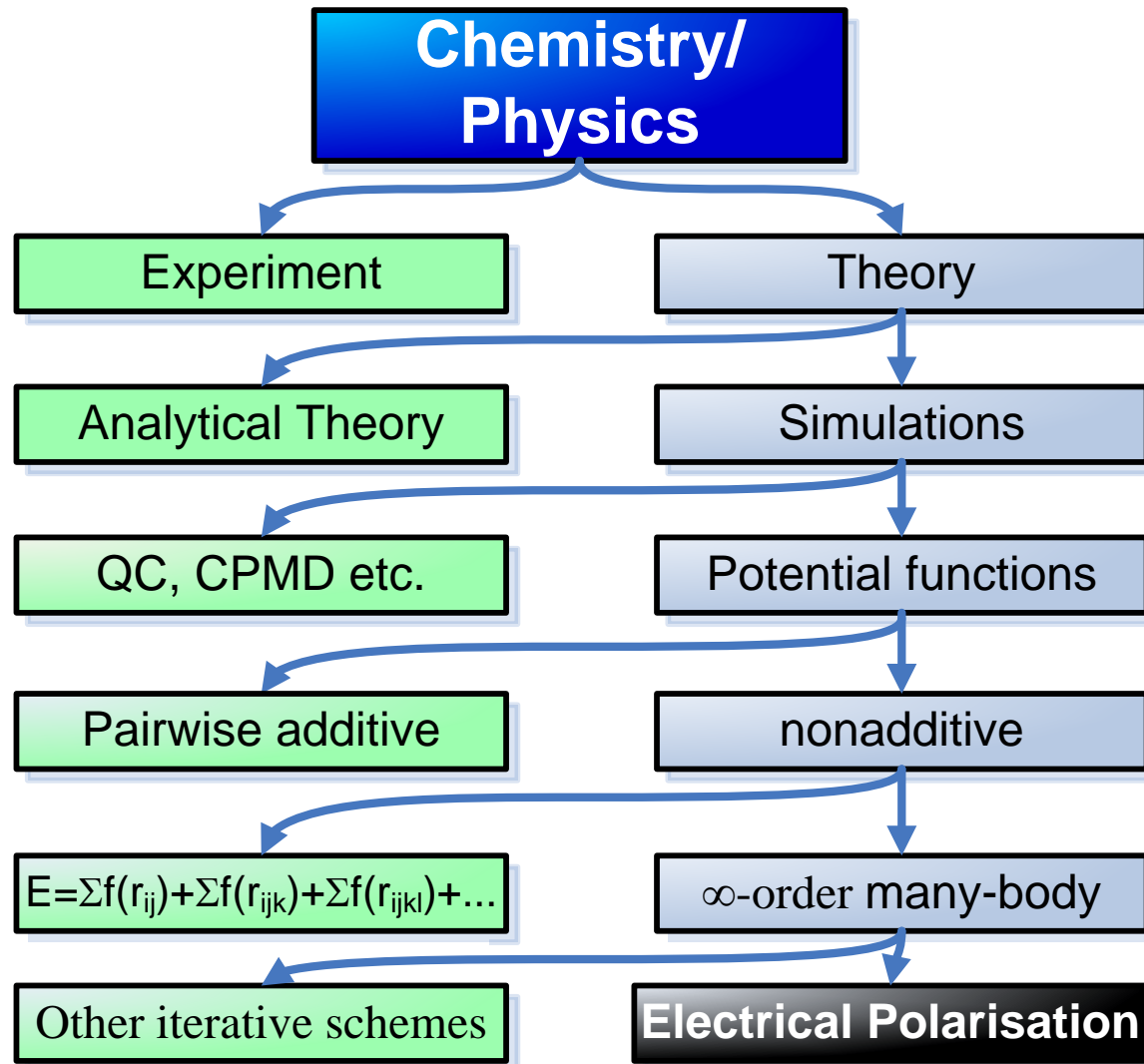
AlCl_3 in water with the following potential energy function:

Interatomic potentials for the intermolecular interactions (in Å and kJ mol^{-1}).^a

$$\begin{aligned}V_{\text{DO}}(r) &= 604.620/r + 111.91 \times 10^3 / r^{8.8591} - 1.045(\exp(-4(r - 3.4)^2) + \exp(-1.5(r - 4.5)^2)) \\V_{\text{OH}}(r) &= -302.31/r + 26.0725/r^{9.19912} - 41.8229/(1 + \exp(40(r - 1.05))) - 16.7292/(1 + \exp(5.49305(r - 2.2))) \\V_{\text{HH}}(r) &= 151.155/r + 418.395(1 + \exp(29.9(r - 1.968))) \\V_{\text{AlO}}(r) &= -2750.51/r - 2495.69/r^2 + 266001 \exp(-3.89948r) \\V_{\text{AlH}}(r) &= 1375.26/r + 160.655/r^2 + 287.458 \exp(-0.35461r) \\V_{\text{ClO}}(r) &= 916.563/r - 111.380/r^2 + 379670 \exp(-3.20906r) \\V_{\text{ClH}}(r) &= -458.281/r + 1.88974 \times 10^{26} \exp(-33.977r) \\V_{\text{AlAl}}(r) &= 12505.0/r - 2360.20/r^2 + 5228.69 \exp(-1.038r) \\V_{\text{AlCl}}(r) &= -4168.14/r - 1115.36/r^2 + 373934 \exp(-3.68r) \\V_{\text{ClCl}}(r) &= 1389.29/r + 28674.4/r^6 + 917099 \exp(-3.39r) \\V_{\text{OAlO}}(r_1, r_2, \alpha) &= 74.8588(0.0641288 + (\pi - \alpha)^2) \exp(-0.246481(r_1^2 + r_2^2))\end{aligned}$$

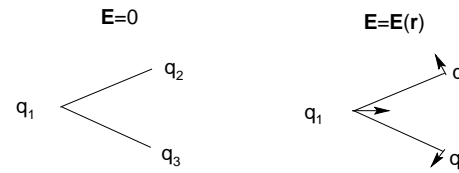
^a In the last line, α is the O-Al-O angle.





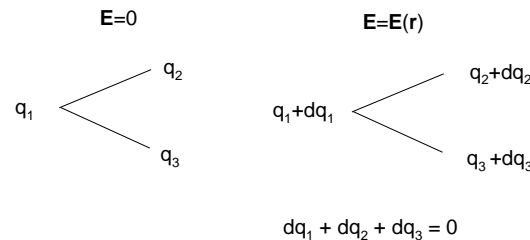
Ways to deal with nonadditive interactions: The point-dipole model (PDM):

Atomic polarizabilities α_i are assigned to some molecular site and the electric field induces the formation of a point dipole μ_i



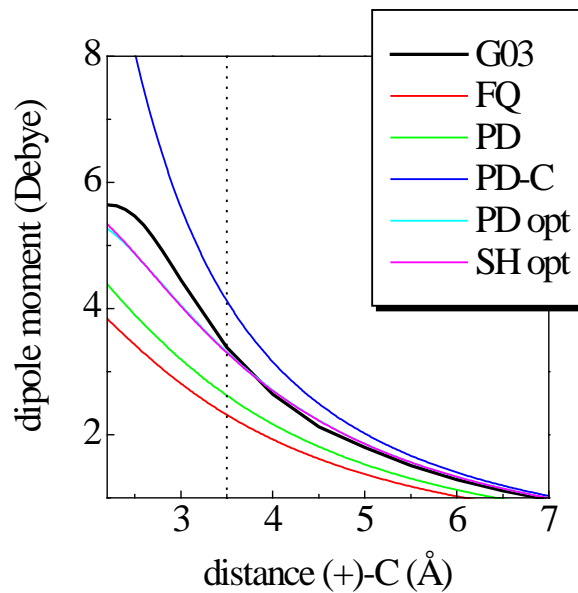
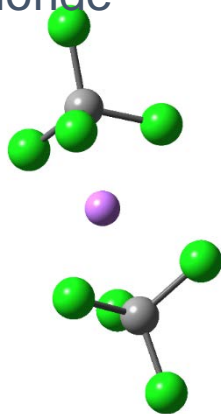
The fluctuating charge model (FQ):

Charges are allowed to fluctuate according to the electronic properties of the molecule as **atomic electronegativity** and **atomic hardness**.



Comparison of the behaviour of these 'polarisation models':

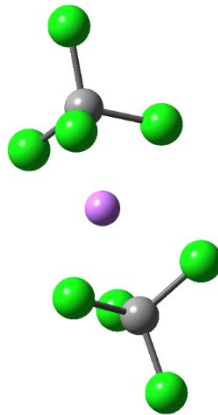
Example:
 Li^+ ion near
Carbon
Tetrachloride



Conclusion:

Induced dipole moments (=polarization) can be large.
The effect on the electrostatic energy can be large

(keep in mind, however, that Li^+ ion near carbon tetrachloride is an extreme example.)



Modeling the molecular interactions: The reality for everyday systems

Normally one divides between:

- Inter- and
- intramolecular interactions

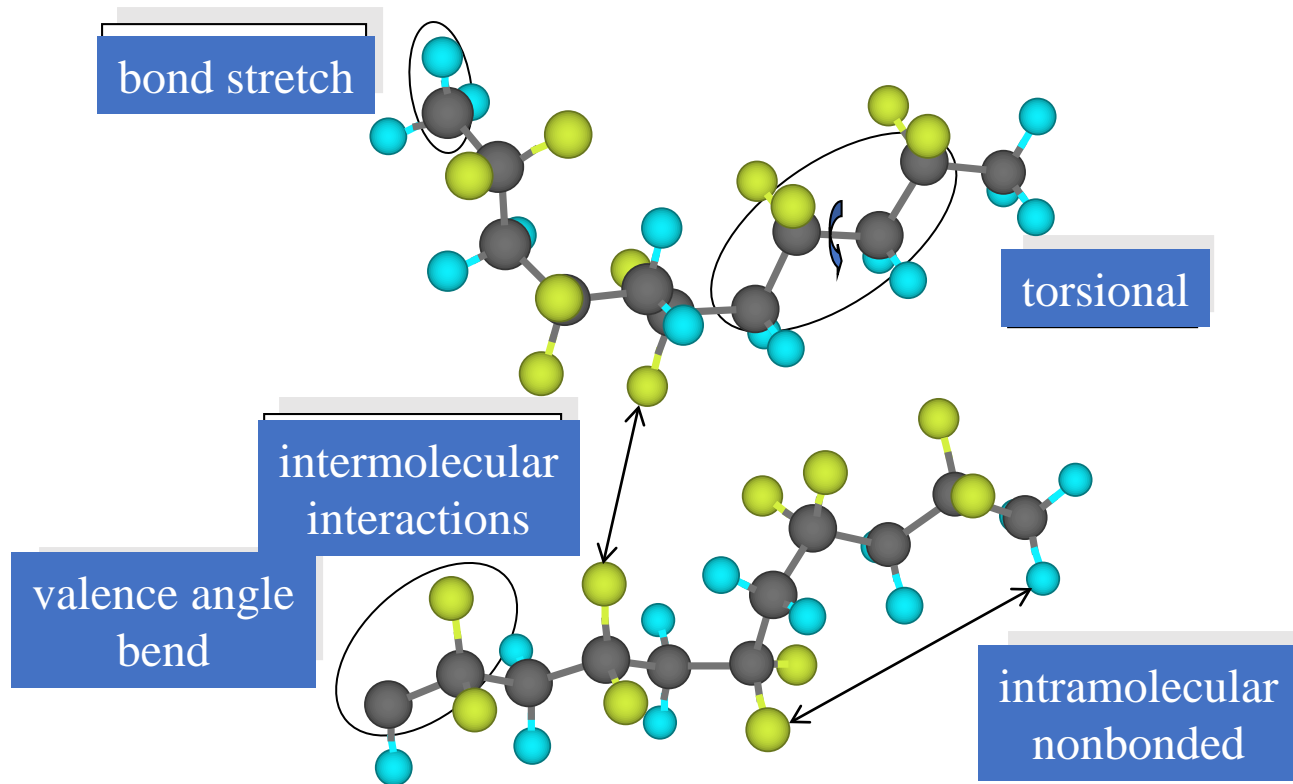
Modeling the molecular interactions

Inter- and intramolecular interactions

Why distinguish between them ?

- In the ideal case, no difference
- In reality:
 - Equal treatment leads is difficult/expensive
 - Examples
 - UFF
 - Central force model of water
 - Reactive potentials
- Why is it advantageous to differentiate ?
 - Bond breaking is difficult to describe
 - Angular (3body) and 4-body terms are natural and needed in a network of bonded atoms (A-B-C-D)
 - but difficult and less necessary between non-bonded atoms (H₂O-M⁺⁺-OH₂)

1) Typical intermolecular energy functions (=force field)



Every atom will be affected by the potential energy functions of every atom in the system. Either from

- Bonded Neighbors
- Non-Bonded Atoms
(=other atoms in the same molecule or atoms from different molecules)

$$V(R) = E_{bonded} + E_{non-bonded}$$

2) Non-Bonded Atoms

In the simplest case, there are three types of potential energy which we need to consider for the interaction between non-bonded atoms:

- Repulsion of electron shells
- van der Waals Potential

$$E_{non-bonded} = E_{repulsion} + E_{van-der-Waals} + E_{electrostatic}$$

The Lennard-Jones form is a compromise between accuracy and fast computability.

Repulsive ($\sim r^{-12}$) and van der Waals ($\sim r^{-6}$) potential terms:

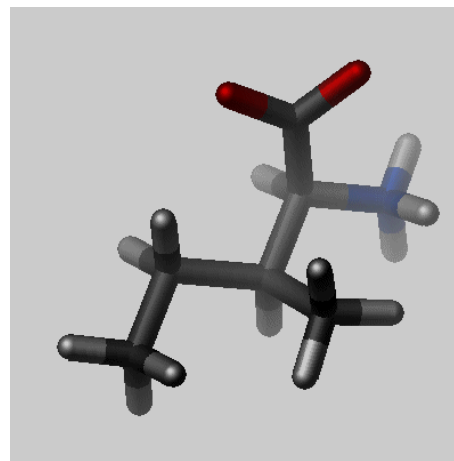
$$E_{\text{Lennard-Jones}} = \sum_{\text{nonbonded pairs}} \left(\frac{A_{ik}}{r_{ik}^{12}} - \frac{C_{ik}}{r_{ik}^6} \right)$$

Scaling, parameters, alternative versions, accuracy, deficits, improvements, physical foundations

Electrostatic Potential

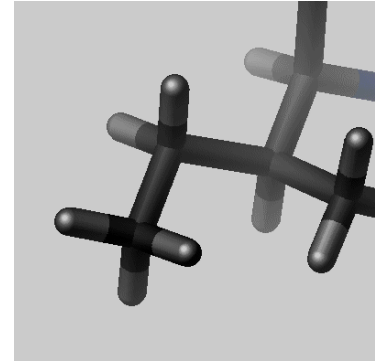
$$E_{\text{electrostatic}} = \sum_{\text{nonbonded pairs}} \frac{q_i q_k}{Dr_{ik}}$$

Bonded Atoms

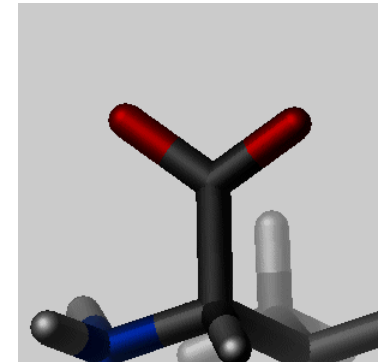


$$E_{\text{bonded}} = E_{\text{bond-stretch}} + E_{\text{angle-bend}} + E_{\text{rotate-along-bond}}$$

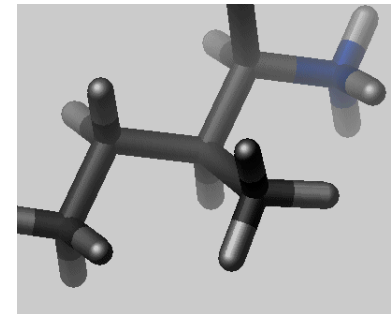
$$E_{bond-stretch} = \sum_{1,2\ pairs} K_b (b - b_0)^2$$



$$E_{bond-bend} = \sum_{angles} K_\theta (\theta - \theta_0)^2$$



$$E_{rotate-bond} = \sum_{1,4\ pairs} K_\phi (1 - \cos(n\phi))$$



Side issue: the electrostatic energy in a crystal: Ewald

$$\begin{aligned}
 U = & \underbrace{\frac{1}{4\pi\epsilon_0} \sum_{\mathbf{n}} \sum_{i=1}^N \sum_{j=i+1}^N q_i q_j \frac{\text{erfc}(\alpha|\mathbf{r}_{ij} + \mathbf{n}|)}{|\mathbf{r}_{ij} + \mathbf{n}|}}_{\text{Real-space term}} \\
 & + \underbrace{\frac{1}{\epsilon_0 V} \sum_{\mathbf{k} > 0} \frac{1}{k^2} e^{-\frac{k^2}{4\alpha^2}} \left\{ \left| \sum_{i=1}^N q_i \cos(\mathbf{k} \cdot \mathbf{r}_i) \right|^2 + \left| \sum_{i=1}^N q_i \sin(\mathbf{k} \cdot \mathbf{r}_i) \right|^2 \right\}}_{\text{Reciprocal-space term}} \\
 & - \underbrace{\frac{\alpha}{4\pi^{\frac{3}{2}}\epsilon_0} \sum_{i=1}^N q_i^2}_{\text{Point self-energy}} - \underbrace{\frac{1}{4\pi\epsilon_0} \sum_{n=1}^M \sum_{\kappa=1}^{N_m} \sum_{\lambda=\kappa+1}^{N_m} q_{n\kappa} q_{n\lambda} \frac{\text{erf}(\alpha|\mathbf{r}_{n\lambda}|)}{|\mathbf{r}_{n\lambda}|}}_{\text{Intra-molecular self energy}} \\
 & - \underbrace{\frac{1}{8\epsilon_0 V \alpha^2} \left| \sum_{i=1}^N q_i \right|^2}_{\text{Charged system term}} + \underbrace{\left[\frac{1}{6\epsilon_0 V} \left| \sum_{i=1}^N q_i \mathbf{r}_i \right|^2 \right]}_{\text{Surface dipole term}}
 \end{aligned}$$

κ, λ indices of sites within a single molecule

N total number of charged sites

M total number of molecules

N_m number of sites on molecule m

\mathbf{p}_i co-ord of site i relative to molecular centre-of-mass,

$\mathbf{r}_i - \mathbf{R}_i$

q_i charge on absolute site i

$q_{m\kappa}$ charge on site κ of molecule m

\mathbf{r}_i Cartesian co-ordinate of site i

\mathbf{r}_{ij}

$\mathbf{r}_j - \mathbf{r}_i$

α real/reciprocal space partition parameter

π_{lm} instantaneous stress tensor

δ_{lm} Kronecker delta symbol

l, m xyz tensor indices

V volume of MD cell

Ewald force

$$\begin{aligned}
 &= \underbrace{\frac{q_i}{4\pi\epsilon_0} \sum_{\mathbf{n}} \sum_{\substack{j=1 \\ j \neq i}}^N q_j \left\{ \frac{\text{erfc}(\alpha|\mathbf{r}_{ij} + \mathbf{n}|)}{|\mathbf{r}_{ij} + \mathbf{n}|} + \frac{2\alpha}{\sqrt{\pi}} e^{-\alpha^2|\mathbf{r}_{ij} + \mathbf{n}|^2} \right\} \frac{\mathbf{r}_{ij} + \mathbf{n}}{|\mathbf{r}_{ij} + \mathbf{n}|^2}}_{\text{Real-space term}} \\
 &+ \underbrace{\frac{2}{\epsilon_0 V} \sum_{\mathbf{k} > 0} q_i \frac{\mathbf{k}}{k^2} e^{-\frac{k^2}{4\alpha^2}} \left\{ \sin(\mathbf{k} \cdot \mathbf{r}_i) \sum_{j=1}^N q_j \cos(\mathbf{k} \cdot \mathbf{r}_j) - \cos(\mathbf{k} \cdot \mathbf{r}_i) \sum_{j=1}^N q_j \sin(\mathbf{k} \cdot \mathbf{r}_j) \right\}}_{\text{Reciprocal-space term}} \\
 &+ \underbrace{\left[\frac{q_i}{6\epsilon_0 V} \left(\sum_{j=1}^N q_j \mathbf{r}_j \right) \right]}_{\text{Surface dipole term}}
 \end{aligned}$$

en.wikipedia.org/wiki/Ewald_summation

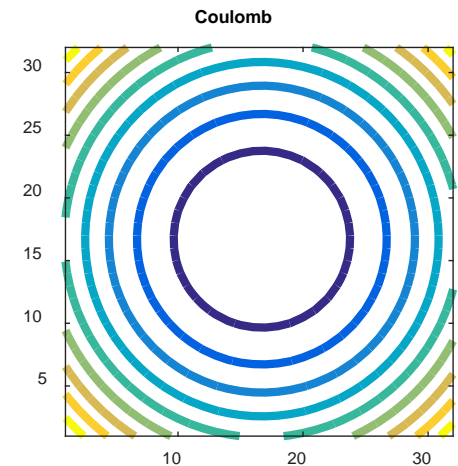
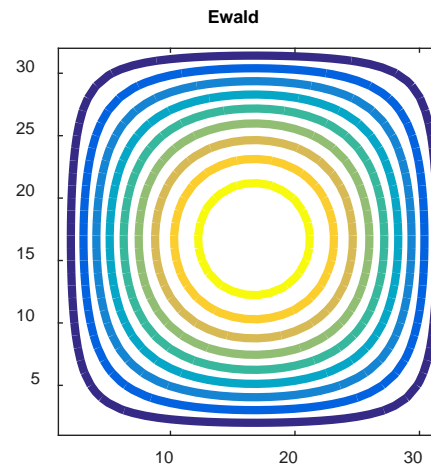
Sketching the Ewald potential

(discuss TV shape, infinity considerations, convergence,

Sketching the Ewald potential

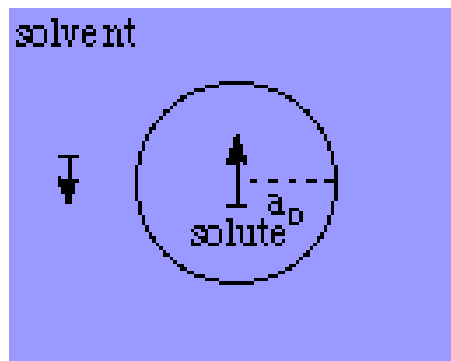
(discuss TV shape, infinity considerations, convergence,

```
[x y]=meshgrid(0:0.1:pi);  
z1=sin(x).^2.*cos(y).^2;  
subplot 121  
[a,b]=contour(z1);  
set(b,'linewidth',3); axis  
square  
title('Ewald');  
z2=(x-pi/2).^2+(y-pi/2).^2;  
subplot 122  
[c,d]=contour(z2);  
set(d,'linewidth',3); axis  
square  
title('Coulomb');
```



Reaction field:

simpler, not periodic, equally good as Ewald
Also useful in quantum chemistry



$$\vec{R} = \frac{2(\epsilon - 1) \cdot \vec{\mu}}{(2\epsilon + 1) \cdot a_0^3}$$

\vec{R} = reaction field

a_0 = cavity radius

$\vec{\mu}$ = molecular dipole moment

Bond breaking

In many instances, potentials where bonds can break are needed.

For example (the so-called Tersoff-Abel potential):

$$E = \sum_{i>j} f_{ij}(r_{ij}) \left[\phi_R^{ij}(r_{ij}) - \frac{b_{ij} + b_{ji}}{2} \phi_A^{ij}(r_{ij}) \right] .$$

$$b_{ij} = (1 + \sum_{k(\neq i,j)} f_{ik}(r_{ik}) g_{ik}(\theta_{ijk}) \exp[2\mu_{ik}(r_{ij} - r_{ik})])^{-1/2}$$