

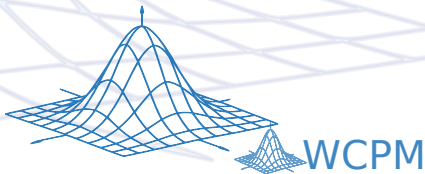
# Uncertainties in classical effective potentials: sources and quantification strategies

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WCPM Seminar  
14 May 2015

THE UNIVERSITY OF  
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# Motivation

## Quantum theory of electrons and nuclei

QM (QED) is a theory with enormous predictive power:

- Energy levels of hydrogen atom to a few ppm.

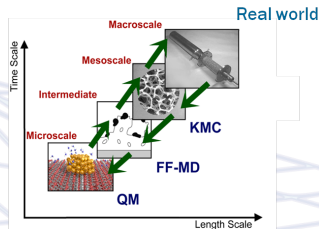
Solution to all our modelling needs? ... No.

## “Sequential” Multiscale Modelling

Use proper simulation tool for each scale.

- Parameterise from small to large.
- No direct coupling between models.

Propagate uncertainty from one scale to the next.



[essenceofscience.se/nobel-2013/](http://essenceofscience.se/nobel-2013/)

## 1 Standard Sequential QM/MM Multiscale Modelling

- Density Functional Theory
- Molecular Dynamics
- Effective Potentials
- Force Matching

## 2 Uncertainty

- Sources
- Quantification
- Examples

# (Plane Wave) Density Functional Theory

## A sequence of approximations

Solve Schrödinger (Dirac) equation of electrons and nuclei

- for stationary nuclei (Born-Oppenheimer approximation),
- mapping the many-electron problem to many one-body problems,
- which use approximative functionals to represent XC,
- while core electrons are treated by pseudopotentials;
- wave functions are represented using plane wave basis set,
- cut off at finite energy and sampled on a finite grid;
- the problem is then solved by iteration to self-consistency.

Depending on some of the choices, further corrections are necessary.

## Uncertainty and errors?

Not all errors are controllable.

- Would merit a talk of its own.



## What is Molecular Dynamics?

Equations of motion of a system of interacting particles are integrated numerically.

- **Direct simulation** of the basic laws of physics: Newton's (or Hamilton's) equations.

## Needed

- Initial condition: **structure model**
- Equation of motion: **model of the interactions**

Big systems or long simulation times are feasible only with **classical effective potentials**.

## Eliminate electronic degrees of freedom

Re-write equation of motion

$$\frac{\partial \mathbf{P}_I}{\partial t} = -\nabla_I(\epsilon_0(\mathbf{R}) + V_{NN}(\mathbf{R})) = \nabla_I(\tilde{V}_{\text{eff}}(\mathbf{R}))$$

Usually,  $V_{\text{eff}}$  much simpler. Expand in manybody contributions:

$$V_{\text{eff}}(\mathbf{R}) = \sum_i \phi_1(\mathbf{r}_i) + \frac{1}{2} \sum_{\substack{i,j \\ i \neq j}} \phi_2(\mathbf{r}_i, \mathbf{r}_j) + \frac{1}{6} \sum_{\substack{i,j,k \\ i \neq j \neq k \neq i}} \phi_3(\mathbf{r}_i, \mathbf{r}_j, \mathbf{r}_k) + \dots$$

Even simpler: Central (homogeneous and isotropic) pair potentials:

$$V(\mathbf{R}) = \frac{1}{2} \sum_{\substack{i,j \\ i \neq j}} \phi_{ij}(r_{ij}), \quad \text{with } r_{ij} = |\mathbf{r}_i - \mathbf{r}_j|$$

## Choice of interaction model depends on material to simulate

(different ways to truncate manybody expansion)

- Central **pair** potentials, **EAM** potentials for metals.
- **Angular** dependent potential (ADP), MEAM.
- **Covalent** potentials ( Tersoff, Stillinger-Weber, ... ) for ceramics.
- **Coulomb** potential (Ewald method, Wolf summation).
- **Dipolar** interaction for oxides.
- Simulation of **organic molecules**: Force fields for polymer chains, water, amino acids,...

## Potentials specified

- by **values** at sampling points (tabulated potentials)
- or by **parameters** (analytic potentials).



# Example: EAM potentials

## EAM potentials

$$E = \sum_{ij} \phi_{ij}(r_{ij}) + \sum_i F_i(\rho_i), \quad \text{where} \quad \rho_i = \sum_j \psi_j(r_{ij})$$

Bond strength **depends on environment** –  
better suited to describe vacancies and other defects.

## Analytic vs. tabulated potentials

- $\phi, F, \rho$  are the potential functions.
- Can be represented by analytic functions, e.g.  $F = c\sqrt{\rho}$ .
- Alternative: Tabulated at sampling points.



## How to obtain effective potentials?

Potential serves to determine energies and forces

→ determines the **physics of the system!**

- Depending on the system (metal, oxide, etc.), a suitable **potential type** must be chosen.
- Within such a potential family, the **potential parameters** determine the physical properties a particular material.
- The parameters are chosen such that the desired **material properties** are correctly reproduced.
- The material properties to be reproduced are often **computed ab-initio**, instead of measured experimentally.

# Reference quantities

## Experimental Quantities of Interest

e.g. phase diagrams, elastic properties, diffusivities. . .

- often expensive to calculate  $\Rightarrow$  “simple” systems only.

## Literature and first-principles quantities

e.g. bond lengths, heats of formation, bond angles.

- Sampling configurational space?

Example: ReaxFF.

## First-principles forces, stresses, energies

- easily calculated (1 MD step).
- sample configurational space.

Force Matching!

Ercolessi & Adams, Europhys. Lett. **26**, 583 (1994)



# Force Matching with *potfit*

## Open source force matching code *potfit*

- Flexible and modular.
- Supports pair, (M)EAM, ADP potentials (metals).
- Oxide potentials.
- Electron-temperature dependent potentials (laser ablation).
- Interfaces to DFT and MD codes.

## Widely used code

- 40 downloads/month,
- 50 citations with potentials,
- from more than ten distinct groups around the globe.

Brommer, Gähler, Model. Simul. Mater. Sci. Eng. **15**, 295 (2007).

<http://potfit.sourceforge.net/>



# Potential Generation

- 1 Select potential **model**, starting potential.
- 2 Select **reference structures** (100–200 atoms, MD simulation at various temperatures, strained structures).
- 3 Calculate forces, stresses, energies with **ab-initio** code.
- 4 Optimize starting potential with **potfit**.
- 5 Generate reference structures with **new potential**.  
⇒ more realistic configurations.
- 6 **Test** potential.

## If results are not satisfying

- use more/different **reference configurations**,
- replace insufficient **potential model**.

and iterate procedure.



## Sources of uncertainty for force-matched potentials

Generic errors:

- “Imported” uncertainty: cannot beat DFT.
- Algorithmic uncertainty: global optimum?

Force Matching specific (structural & parameter) uncertainties:

- Bad reference data selection (parameter uncertainty).
- Wrong functional form (model bias).
- Overfitting (parameter uncertainty).
- Wrong potential model (model bias).

## Properties of force-matched potentials:

- (Generally) good representability.
- Limited transferability.

Caveat emptor! (US\$2M NSF CDI grant, cf. <https://openkim.org/>).

# Interpolated potentials

## Special case: tabulated or interpolated potential

Interpolated potentials can have many parameters ( $>100$ ).

- No bias from particular functional form.
- Parameters have no **meaning**.

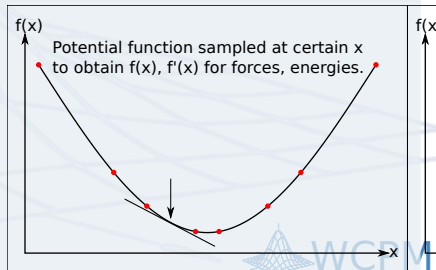
No formal uncertainty propagation in literature.

## Confidence of sampling point values

Forces: evaluate potential functions and gradients.

- Both in training and use.

Training set and application: sample similarly.



# Potential derivation as an inverse problem

Forward problem: data (e.g. forces)  $\delta$  from parameters  $u$

$$\delta = G(u)$$

- $u$ : Parameter vector
- $\delta$ : Result vector

Well-posed problem.

Inverse problem: parameters  $u$  from noisy data  $\delta$

$$\delta = G(u) + \eta$$

Ill-posed. Optimisation problem:

$$\min_u \frac{1}{2} \|\delta - G(u)\|^2$$

# Potential derivation as a Bayesian inverse problem

Interpret  $u, \eta, \delta$  as RV/random fields

Attack problem using “standard” techniques:

- MCMC methods
  - MAP estimators
  - Sparse deterministic approximations
- ⇒ Account naturally for noisy reference data.

Computationally involved (to say the least).

In the following:

- How has this been used so far (selection)?

The problem is two-fold:

- 1 Quantifying uncertainty in potential (inverse problem).
- 2 Propagate uncertainty to MD simulations (forward problem).



## Replace best-fit effective potential by ensemble

Conditional probability of parameter set  $u$ :

$$P(u|\delta, G) \propto \exp \left[ -\frac{C(u)}{T} \right],$$

- cost function  $C(u) = \frac{1}{2} \|\delta - G(u)\|^2$
- formal temperature  $T$ , with  $T_0 = 2C_0/N_p$ ,  $C_0 = \min_u C(u)$

For any observable  $O$ , calculate mean  $\langle O \rangle|_{T,D,M}$ , variance  $\sigma_O^2|_{T,D,M}$ .

Frederiksen *et al.*, PRL **93** 165501 (2004)

# Sloppy model pros and cons

## Advantages

- Integrates seamlessly into force matching/MD software stack:
  - Generate ensemble of potentials.
  - Run  $N$  independent MD simulations.
- Accounts for model & parameter errors.
- Computational cost: factor  $N$ .

## Disadvantages

- No error propagation from DFT (DFT=truth).
- Sampling of parameter space can be tricky (anisotropic).

Open PhD position: Contact PB!

## Bayesian formulation

$$\pi(u|\delta) = \frac{1}{C} p(\delta|u) q(u),$$

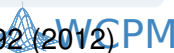
- $q(u)$ : prior on the parameters.
- $\pi(u|\delta)$ : posterior density.
- $p(\delta|u)$ : likelihood, contains diffence data/predictions.

## Direct Bayesian approach prohibitive,

if  $p$  expensive (e.g. full MD sim).

- Use surrogate models
  - non-intrusive spectral projection (NISP)
  - nondeterministic PC expansion

Rizzi *et al.*, Multiscale Modeling & Simulation **10**, 1460–1492 (2012)



# Rizzi's method pros and cons

## Demonstration on model system

TIP4P (water model, 4 parameters).

- Recover 3 parameter values (4th parameter assumed known). . .
- . . . from 30 “measurements” of density, self-diffusion, enthalpy.
- Model errors not tested (work underway).

## Open questions

- How does it scale in parameters, observables?
- Use in force matching context?

## The framework

For model class  $M$ :

$$\pi(u|\delta, M) = \frac{1}{C} p(\delta|u, M) q(u|M),$$

- Transitional MCMC algorithm.
- Surrogate models to reduce computational cost.

Angelikopoulos *et al.*, J. Chem. Phys. **137**, 144103 (2012)

Hadjidoukas *et al.*, J. Comp. Phys. **284** 1–21 (2015)

## Demonstration system

Argon, LJ potential (2 parameters).

- Analytic expressions available for many QoI.
- Functionality also works with full MD.
- Model error: additional parameter.

## Computational cost

Time to solution: 3 days (48 nodes  $\times$  16 cores).

# Conclusions

## Force Matching

Extending atomistic simulations to new materials:

- Preserve DFT precision to larger systems, longer times.
- Foundation for other atomistic and meso-scale problems.

Essential part of multi-scale modelling stack.

## Uncertainty

Work on UQ for effective potentials has just started.

- UQ in potential derivation: Bayesian inverse problem.
- UQ in MD simulations: Re-use old tools or shape new ones?

New modelling paradigm still needs work.