

Accessing large-scale molecular simulations through machine learning

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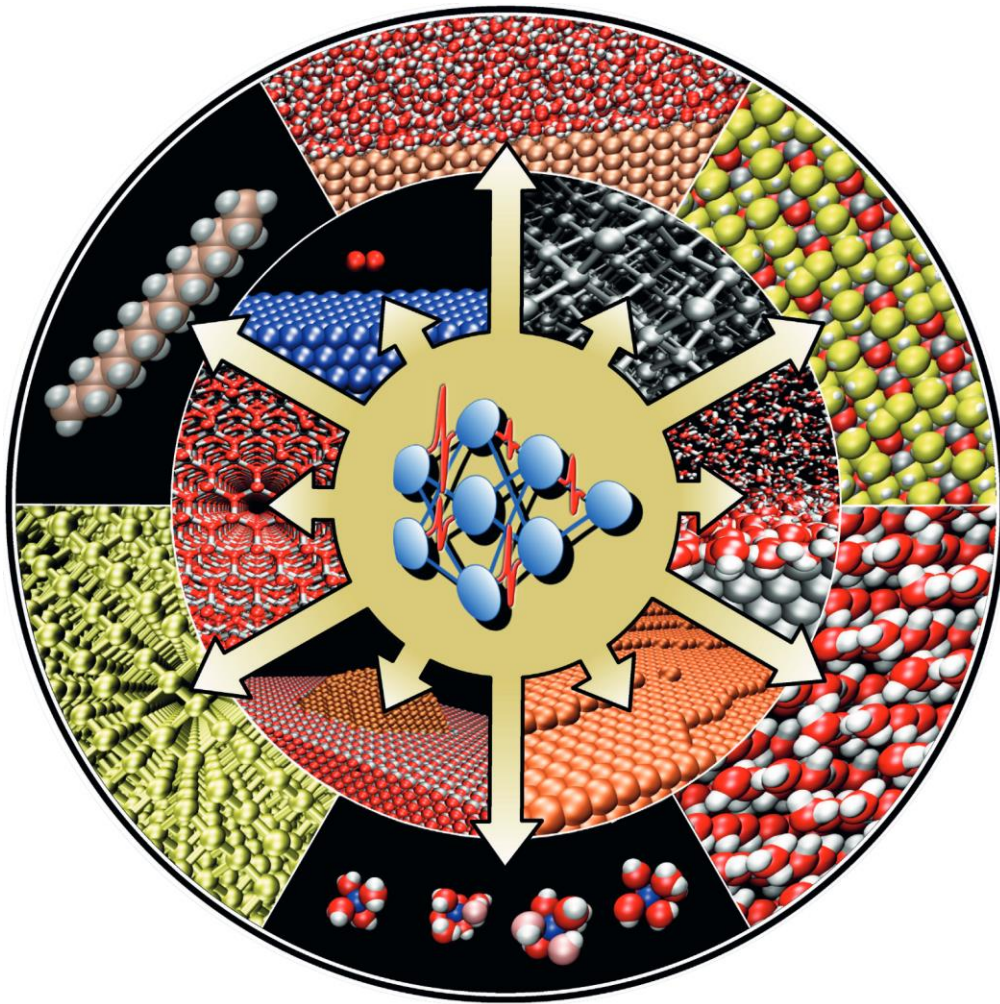
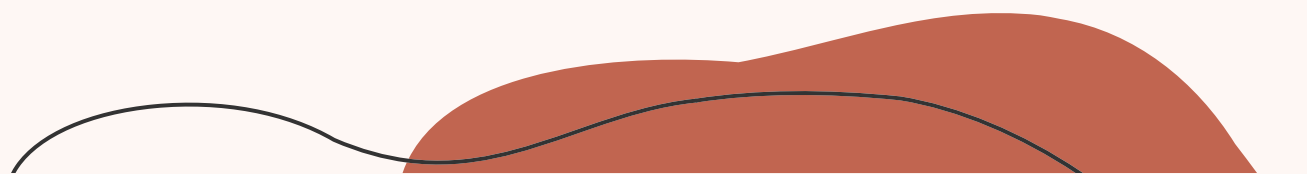


Image credits: [1]



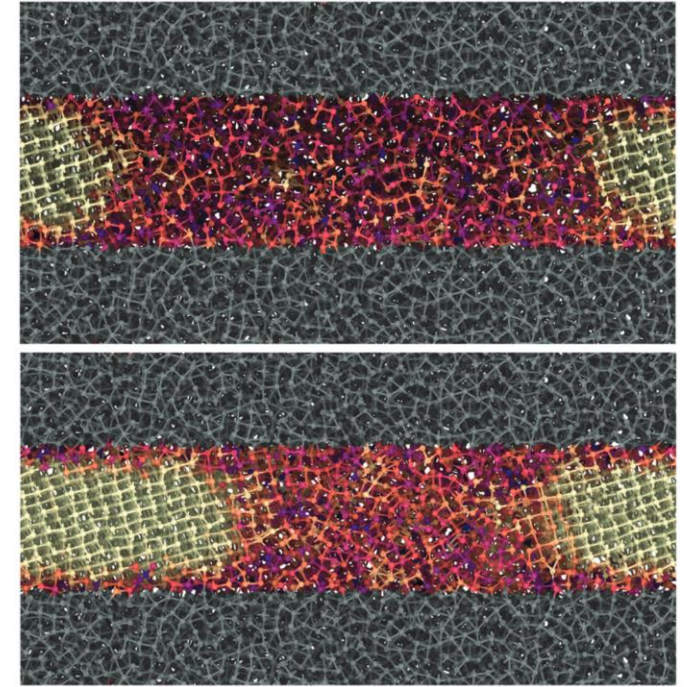
Overview

- Why large-scale?
 - The bottleneck of molecular simulations.
 - Short introduction to Neural Networks.
 - Neural-Network methods for molecular simulations.
 - My research topic: application to *Phase-Change materials*.
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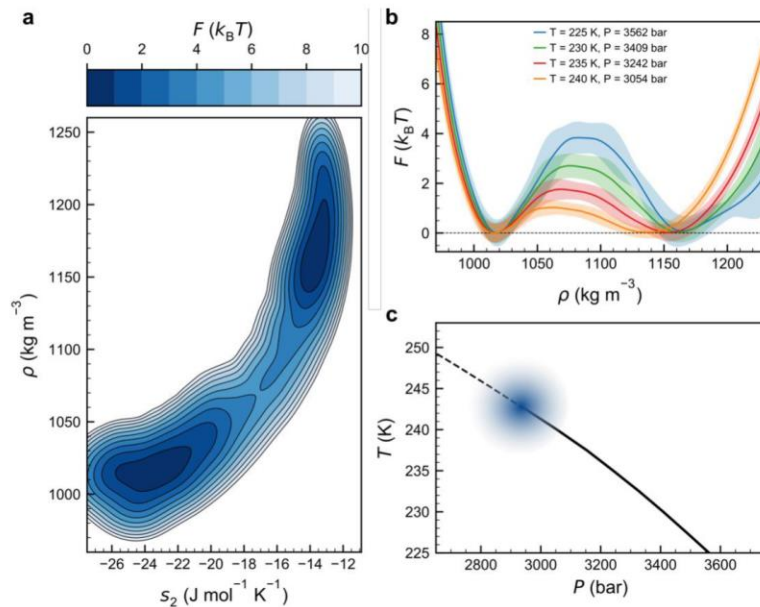
Why large scale?

Simulation of crystals can be done on small scale (unit cell), because of their spatial symmetry.

But amorphous/disordered systems (liquids, glasses, biomolecules, complex systems) require large spatial and time scales:



Growth of the crystal front in thin Sb. Dragoni et al., *Nanoscale* (2021).



Liquid-Liquid critical point for a water model. Gartner et al., *Phys. Rev. Lett.* (2022).

- Long-wavelength vibrational modes.
- Slow relaxation dynamics near criticality.
- Study of collective phenomena.
- Free energy computations.
- Reduction of finite-size effects.

Born-Oppenheimer approximation

Legend

Electronic+nuclear (e+n) Schrödinger equation:

$$H_{e+n}\psi_E = E\psi_E$$

$$H_{e+n} = [T_e + V_{ee} + V_{en} + V_{nn}] + T_n$$

T_a = Kinetic energy of system a.

V_{ab} = Coulomb interaction between systems a,b.

N_e, N = Number of electrons/nuclei.

$r^N = \{\mathbf{r}_i\}_{i=1}^N$ Spatial coordinates.

separation of energy/time scales

$$m_n/m_e > 200$$



separate the wavefunction:

$$\psi_E(r^{N_e}, r^N) = \sum_{\nu} \phi_{E\nu}(r^{N_e}|r^N) F_{\nu}(r^N)$$

Solve the electron problem first, at fixed nuclear positions:

$$[T_e + V_{ee} + V_{en} + V_{nn}] \phi_E(r^{N_e}|r^N) = U(r^N) \phi_E(r^{N_e}|r^N)$$

Classical Dynamics of the nuclei

The electrons induce an **effective interaction potential** on the nuclear problem:

$$[T_n + U(r^N)] F(r^N) = EF(r^N)$$

Effective Hamiltonian for the nuclei: $H = T_n + U(r^N) = \sum_{i=1}^N \frac{\mathbf{p}_i^2}{2m_i} + U(\mathbf{r}_1, \dots, \mathbf{r}_N)$

In a classical treatment, the dynamics follows Hamilton equations:

$$\begin{cases} \dot{r}_{i,\alpha} = \frac{\partial H}{\partial p_{i,\alpha}} = \frac{p_{i,\alpha}}{m_i} \\ \dot{p}_{i,\alpha} = -\frac{\partial H}{\partial r_{i,\alpha}} = -\frac{\partial U}{\partial r_{i,\alpha}} \end{cases}$$

Classical Molecular Dynamics algorithm

Pseudocode (from [2])

```
program md
```

simple MD program

```
call init
```

initialization

```
t=0
```

```
do while (t.lt.tmax)
```

MD loop

```
    call force(f,en)
```

determine the forces

```
    call integrate(f,en)
```

integrate equations of motion

```
    t=t+delt
```

```
    call sample
```

sample averages

```
enddo
```

```
stop
```

```
end
```

Classical Molecular Dynamics algorithm

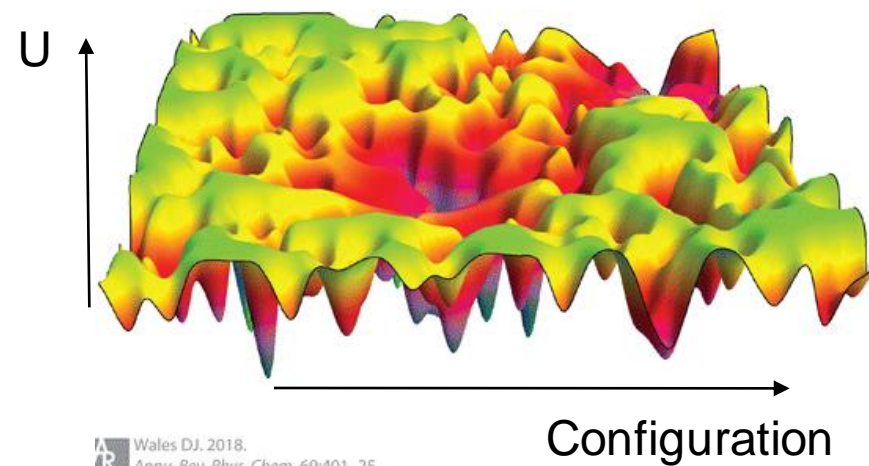
Pseudocode (from [2])

<code>program md</code>	simple MD program
<code>call init</code>	initialization
<code>t=0</code>	
<code>do while (t.lt.tmax)</code>	MD loop
<code> call force(f,en)</code>	determine the forces
<code> call integrate(f,en)</code>	integrate equations of motion
<code> t=t+delt</code>	
<code> call sample</code>	sample averages
<code>enddo</code>	
<code>stop</code>	
<code>end</code>	

Bottleneck!

Interaction Potential

a.k.a. Potential Energy Surface (PES)



PES $U(r)$

Ab-Initio
method

For example,
Density Functional
Theory (DFT)



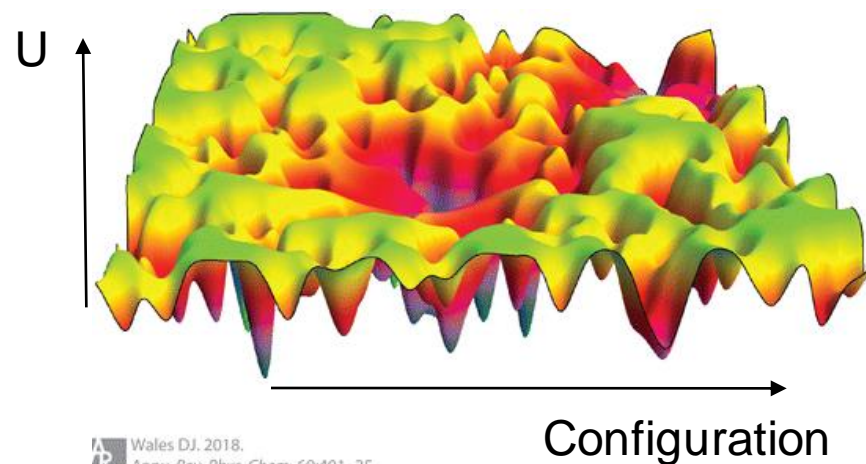
Slow implementation.



Accurate PES.
Dynamical chemical bonds.

Interaction Potential

a.k.a. Potential Energy Surface (PES)



PES $U(r)$

Ab-Initio
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Empirical
parametrization

For example,
Density Functional
Theory (DFT)

For example,
Lennard-Jones
pairwise potential.



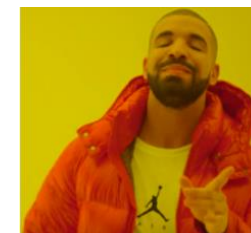
Slow implementation.

Fast implementation.



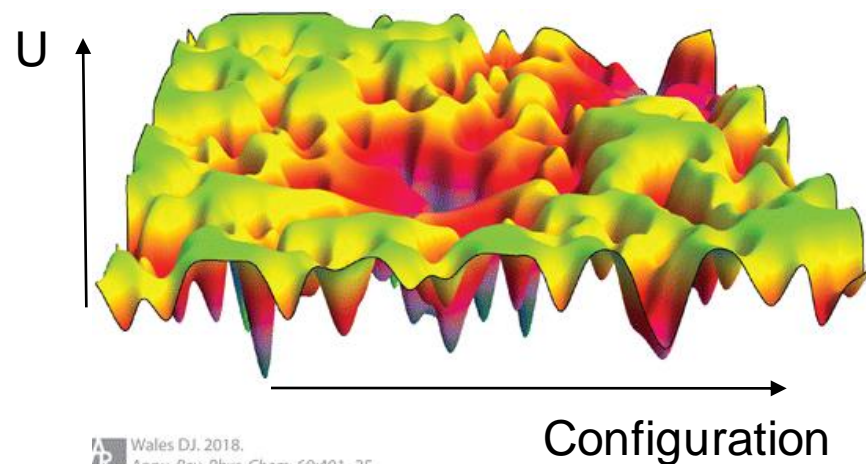
Accurate PES.
Dynamical chemical bonds.

Inaccurate PES (usually).
Static chemical bonds.



Interaction Potential

a.k.a. Potential Energy Surface (PES)



PES $U(r)$

Ab-Initio
method

Neural-Network
parametrization

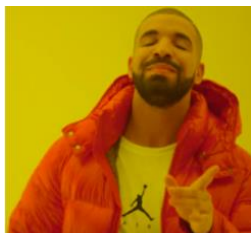
Empirical
parametrization

For example,
Density Functional
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For example,
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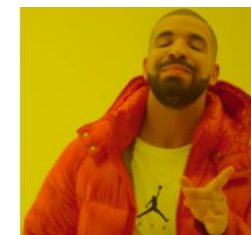
Slow implementation.

Fast implementation.

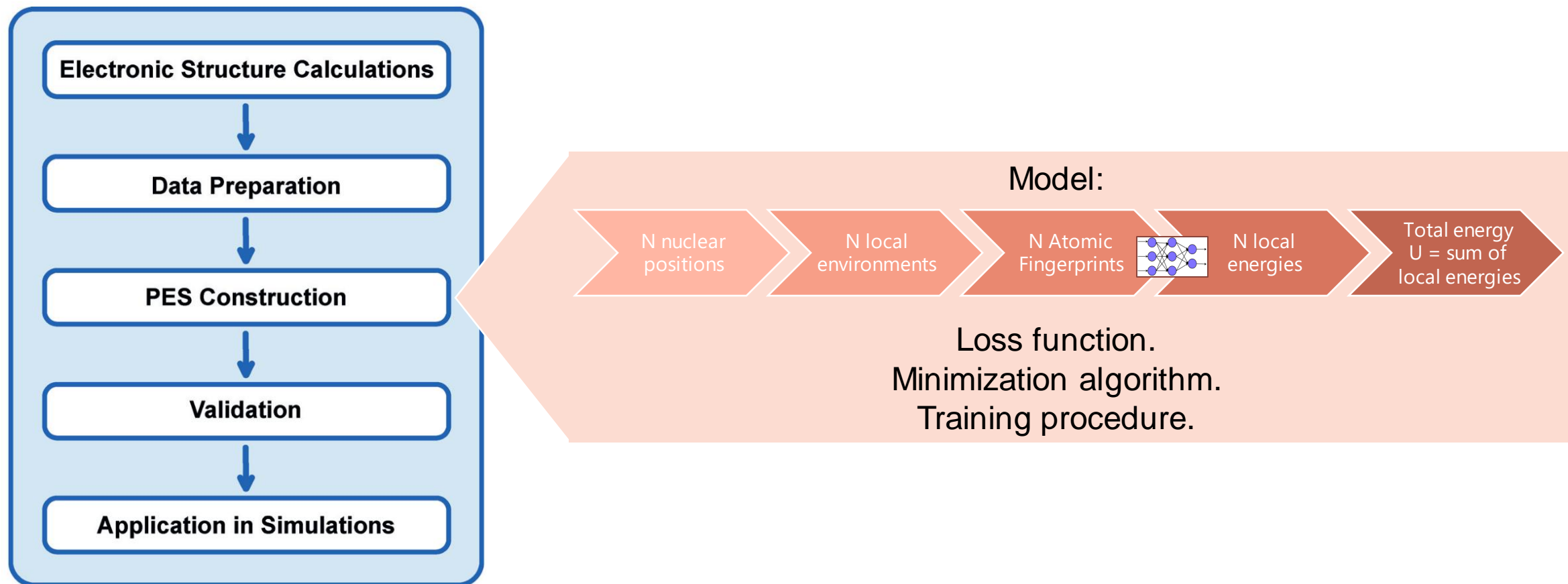


Accurate.
Dynamical chemical bonds.

Inaccurate in most cases.
Static chemical bonds.

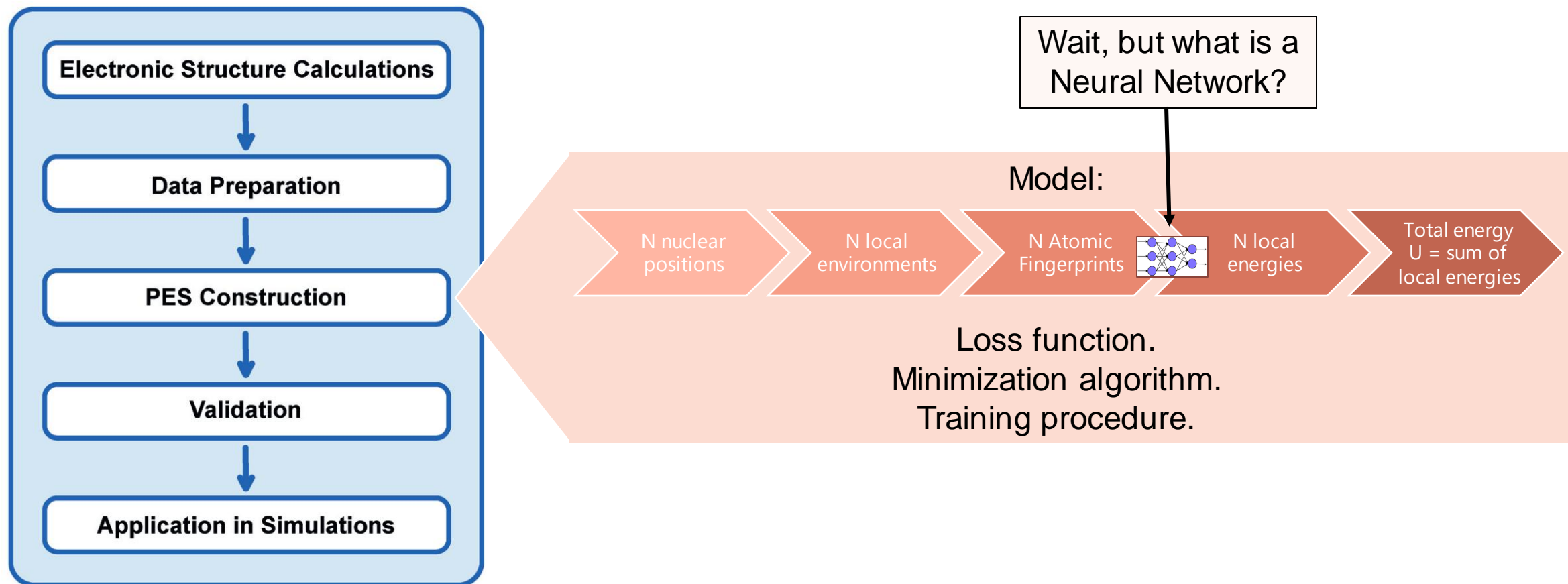


How to build a Neural-Network model for the PES



Flow chart from [1].

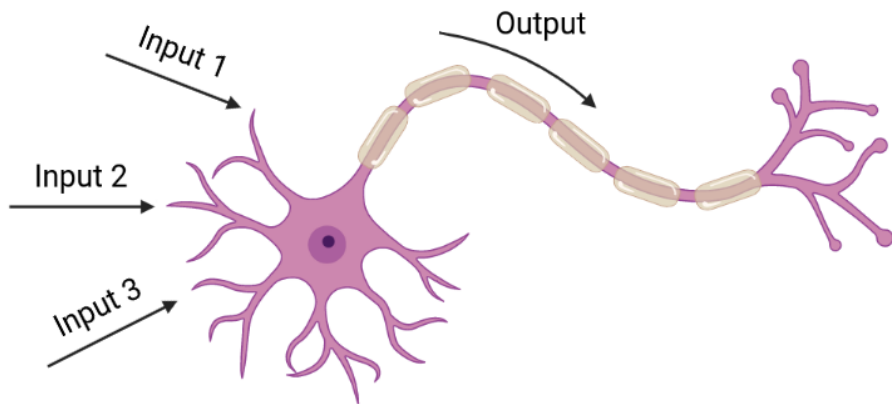
How to build a Neural-Network model for the PES



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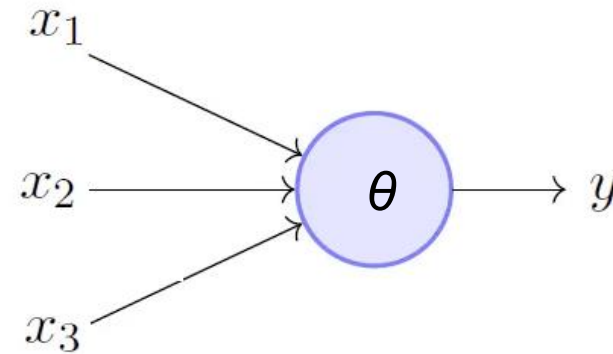
Basics of Artificial Neural Networks

Biological neuron



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McCulloch & Pitts (1943) model

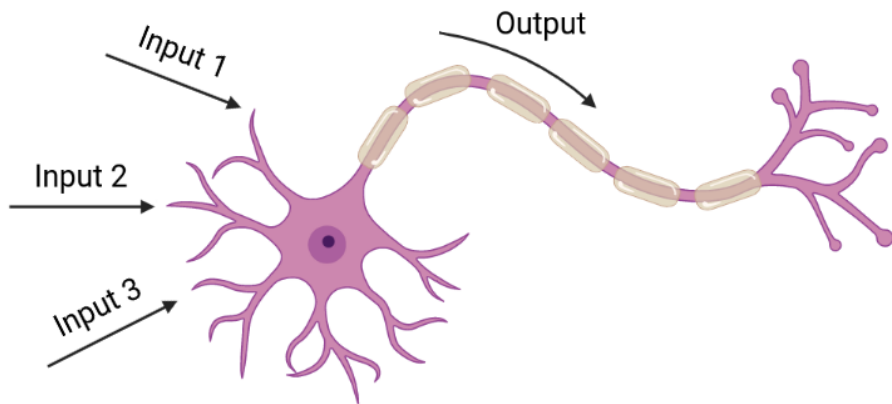


$$y = f(\mathbf{x}|\theta) = \begin{cases} 1 & \text{if } \sum_k x_k > \theta \text{ and neuron is active} \\ 0 & \text{otherwise} \end{cases}$$

binary input/output with a sum threshold θ

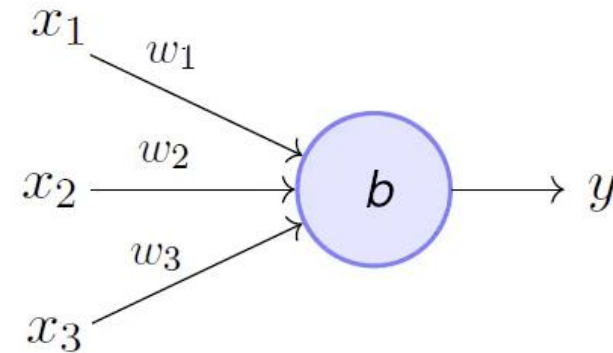
Basics of Artificial Neural Networks

Biological neuron



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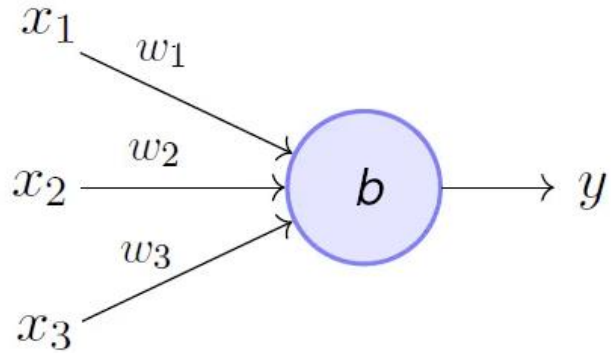
Rosenblatt's *Perceptron* (1958) model



$$y = f(\mathbf{x}|\mathbf{w}, b) = \begin{cases} 1 & \text{if } \sum_k w_k x_k + b > 0 \\ 0 & \text{otherwise} \end{cases}$$

linear mapping + non-linear threshold

Perceptron



$$y = f(\mathbf{x}|\mathbf{w}, b) = \sigma(\underbrace{\mathbf{w}^T \cdot \mathbf{x} + b}_{\text{linear mapping with weights } (\mathbf{w}, b)})$$

non-linear activation function σ
(e.g. tanh, sigmoid, ...)

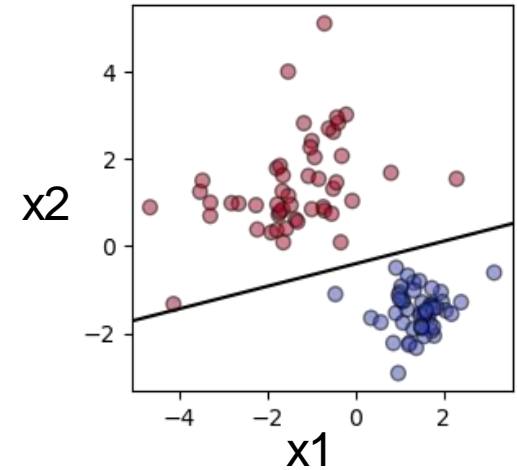
linear mapping with weights (\mathbf{w}, b)

The decision boundary is a **hyperplane**:

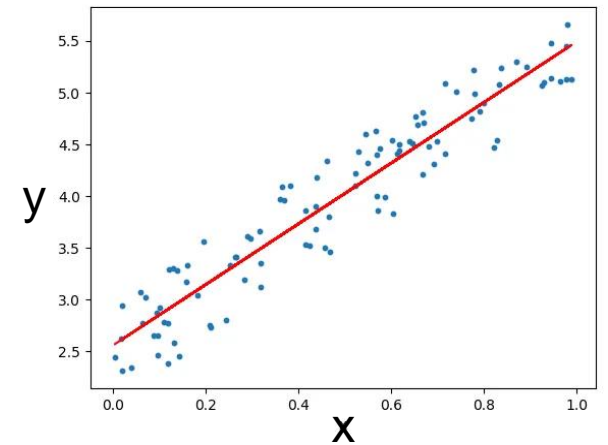
$$b + \mathbf{w}^T \mathbf{x} = 0$$

The task of finding the optimal parameters (\mathbf{w}, b) for classifying \mathbf{x} can be seen both as:

- **Linear classification:** find the best hyperplane dividing the two classes.

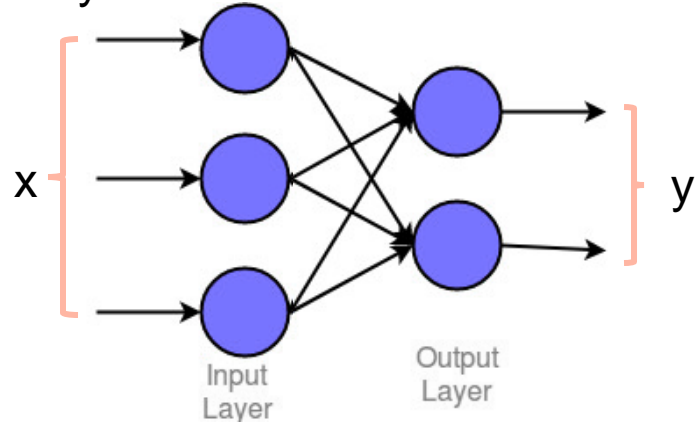


- **Linear regression:** find the best fitting hyperplane to the boundary between the two classes.



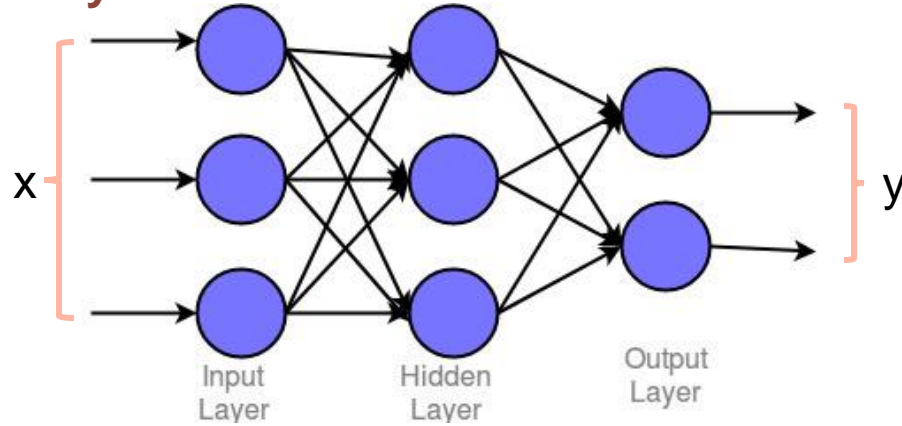
Non-linear Regression with Multi-Layer Perceptrons (MLP)

Single Layer



Non-linear Regression with Multi-Layer Perceptrons (MLP)

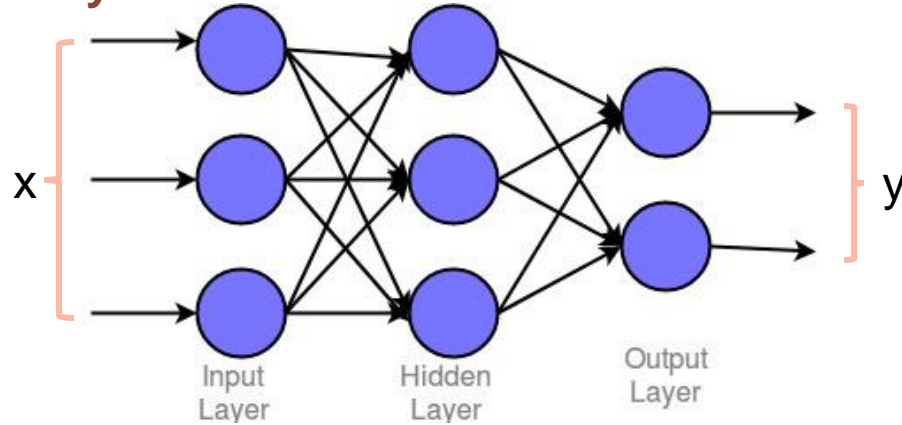
Multi-Layer



Universal approximation theorem:
A MLP is a universal approximator, if deep enough.

Non-linear Regression with Multi-Layer Perceptrons (MLP)

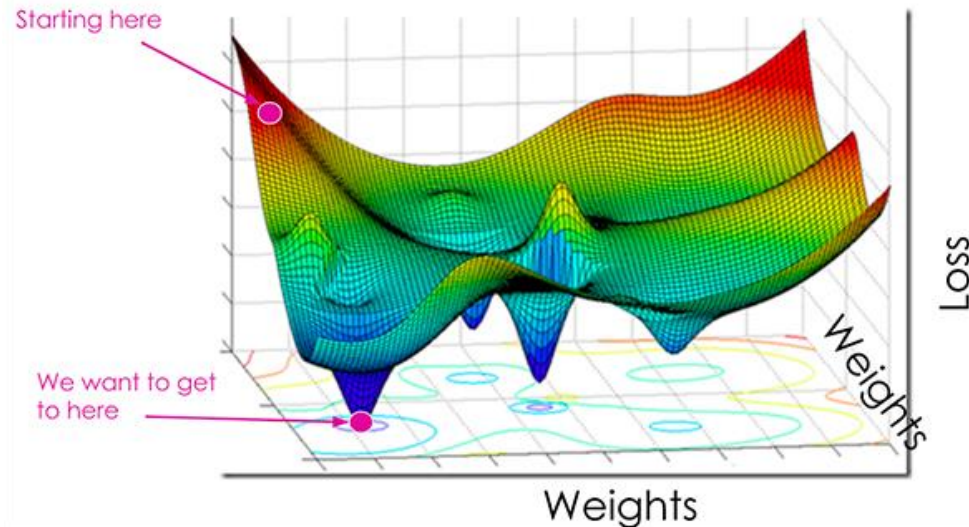
Multi-Layer



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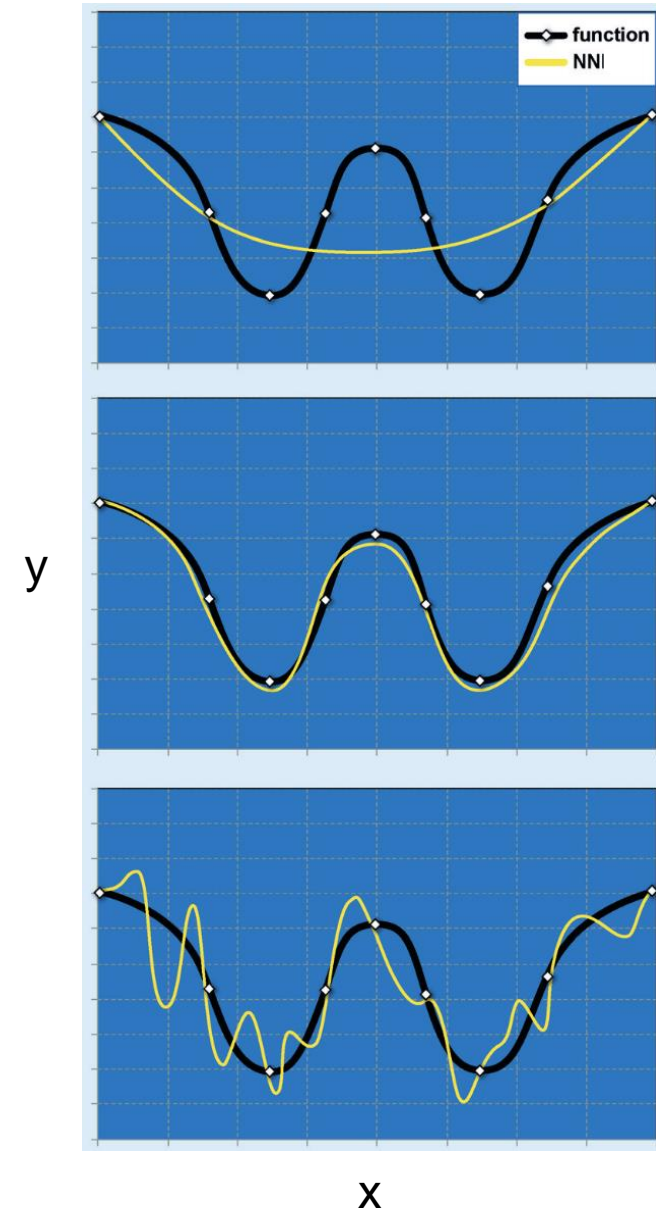
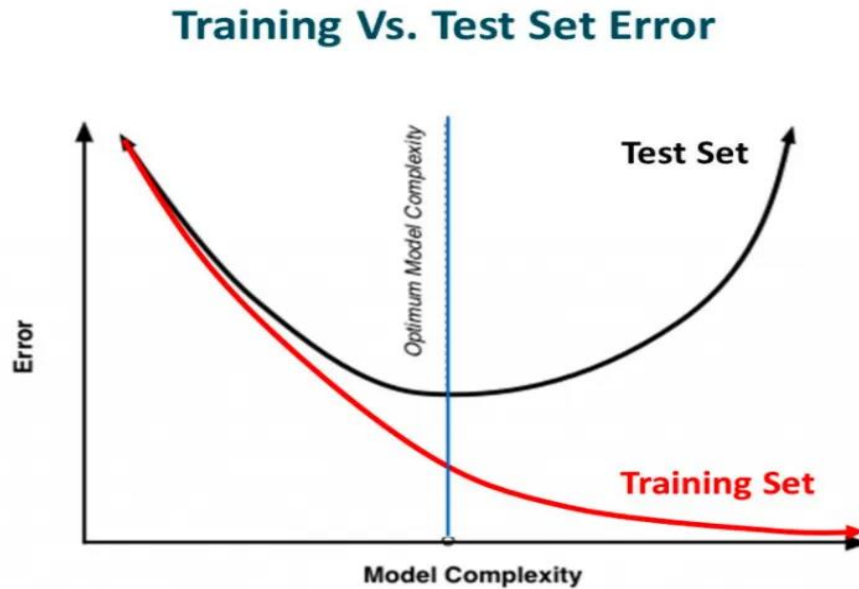
How to find the optimal parameters, at fixed network architecture?

1. Define a loss function $L(w)$ between the target y and the prediction $\hat{y} = f(x|w)$.
2. Use a minimization algorithm on $L(w)$.



Common problems in NN applications

- Overfitting and generalization problems

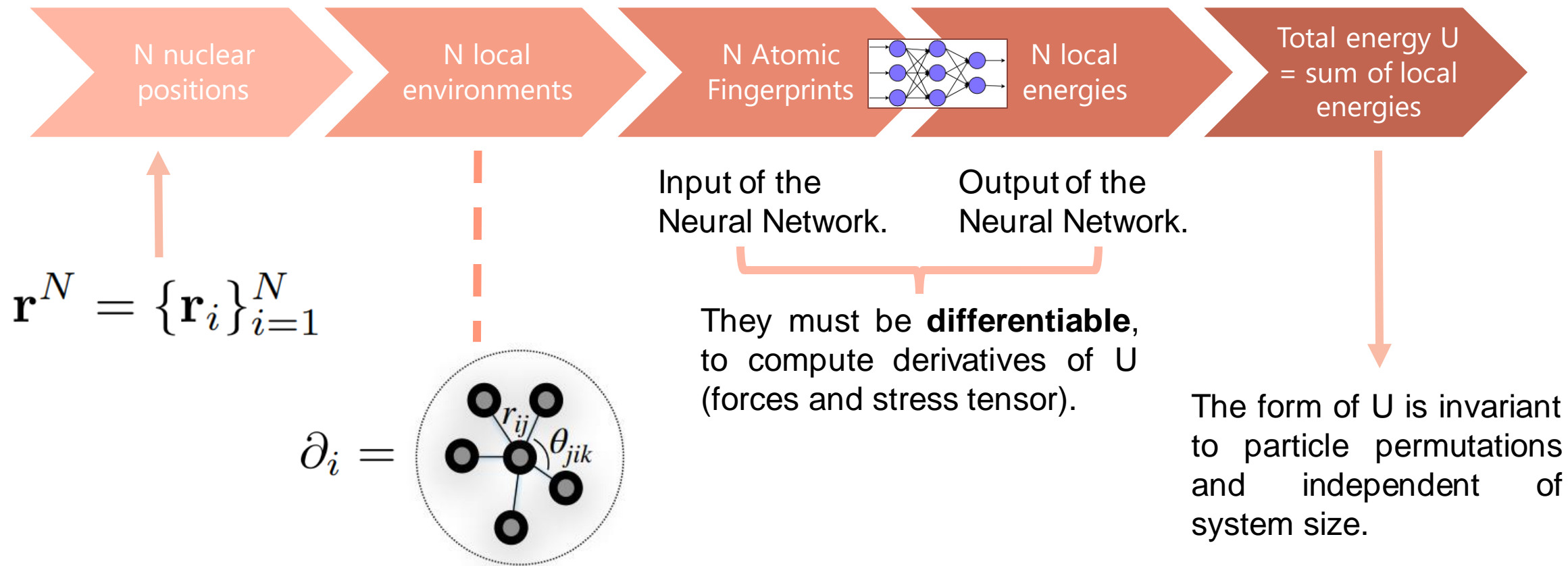




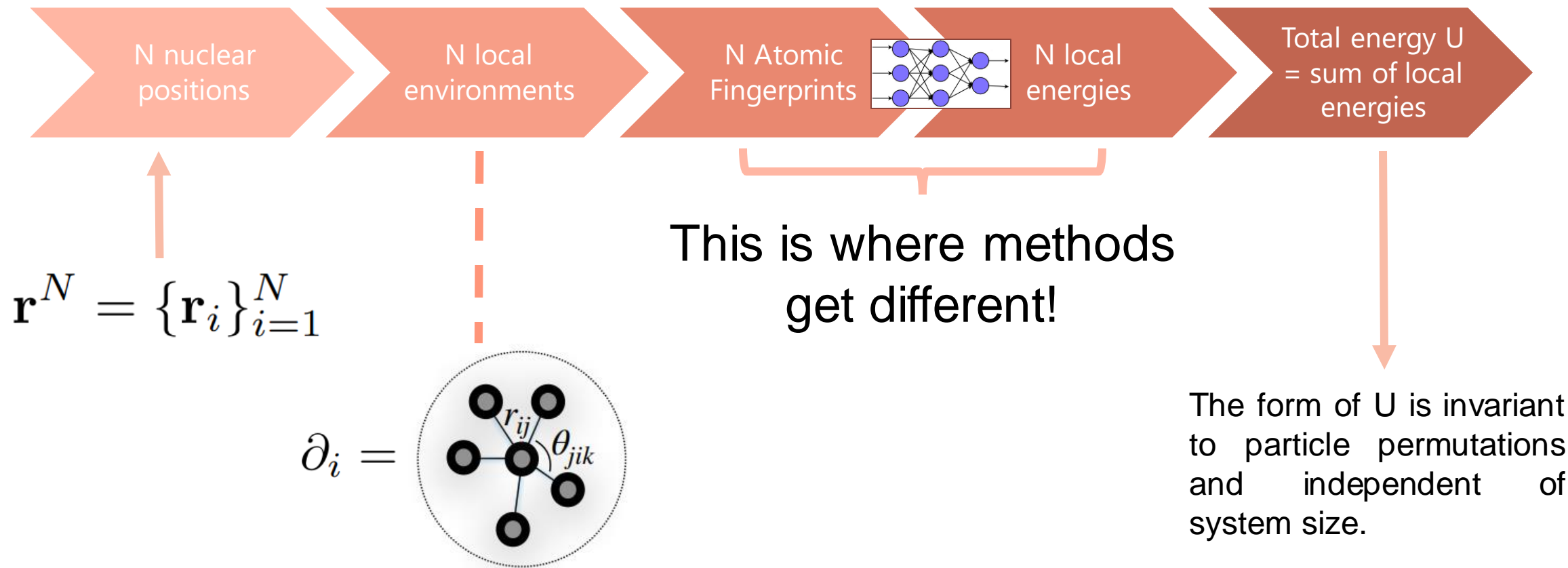
Common problems in NN applications

- Overfitting and generalization problems
- Dataset size and variability
- Choice of the input features

Neural-Network Interaction Potential



Neural-Network Interaction Potential



Behler-Parrinello (2007)



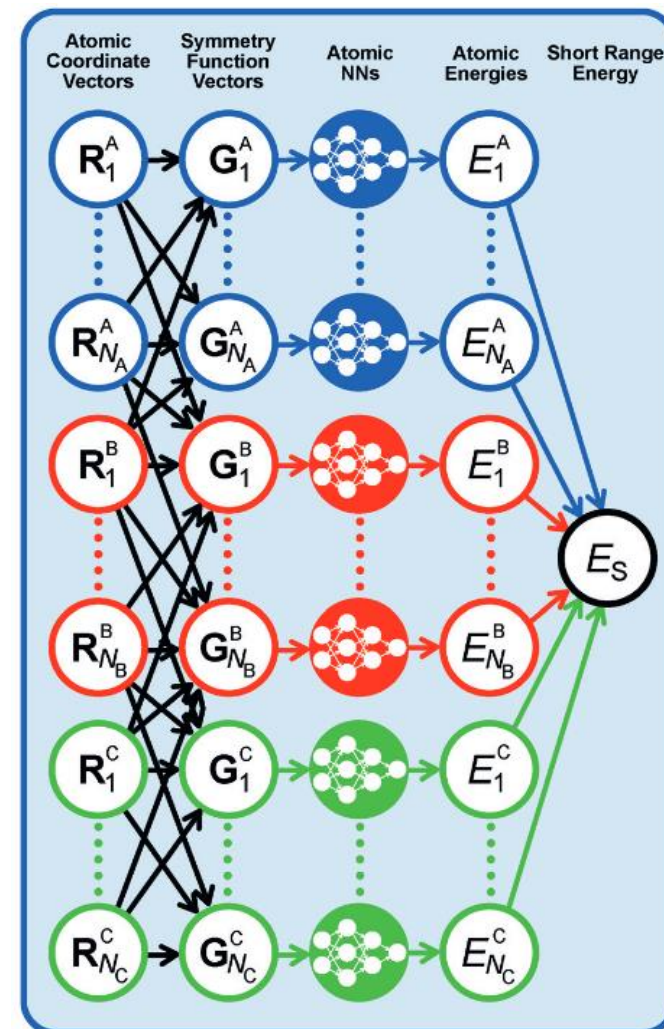
2-body and 3-body symmetry functions with **manually-tuned** parameters.

$$G_i^1 = \sum_{j \neq i}^{\text{all}} e^{-\eta(R_{ij}-R_s)^2} f_c(R_{ij})$$

$$G_i^2 = 2^{1-\zeta} \sum_{j,k \neq i}^{\text{all}} (1 + \lambda \cos \theta_{ijk})^\zeta \times e^{-\eta(R_{ij}^2 + R_{ik}^2 + R_{jk}^2)} f_c(R_{ij}) f_c(R_{ik}) f_c(R_{jk})$$

with $f_c(R_{ij}) = \begin{cases} 0.5 \times [\cos(\frac{\pi R_{ij}}{R_c}) + 1] & \text{for } R_{ij} \leq R_c \\ 0 & \text{for } R_{ij} > R_c \end{cases}$

A MLP for each species, each with 2 hidden layers of ~40 nodes.



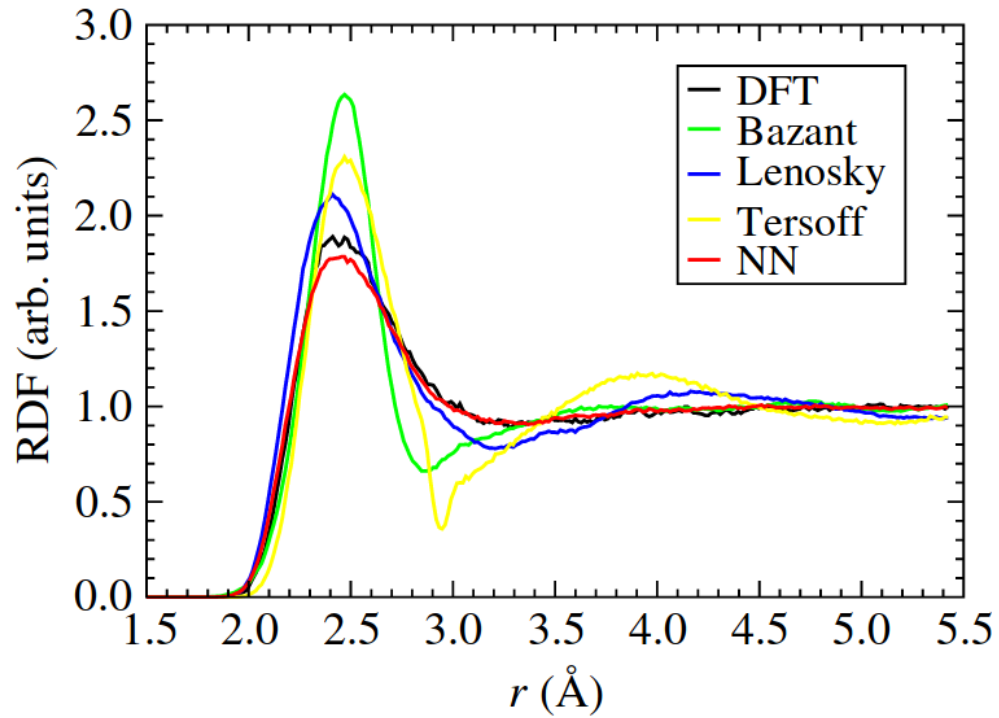
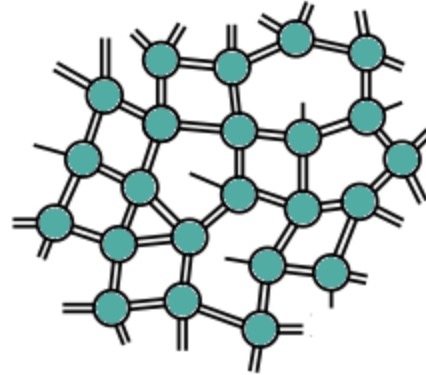
Example architecture for a 3-species system.

Good results on Silicon

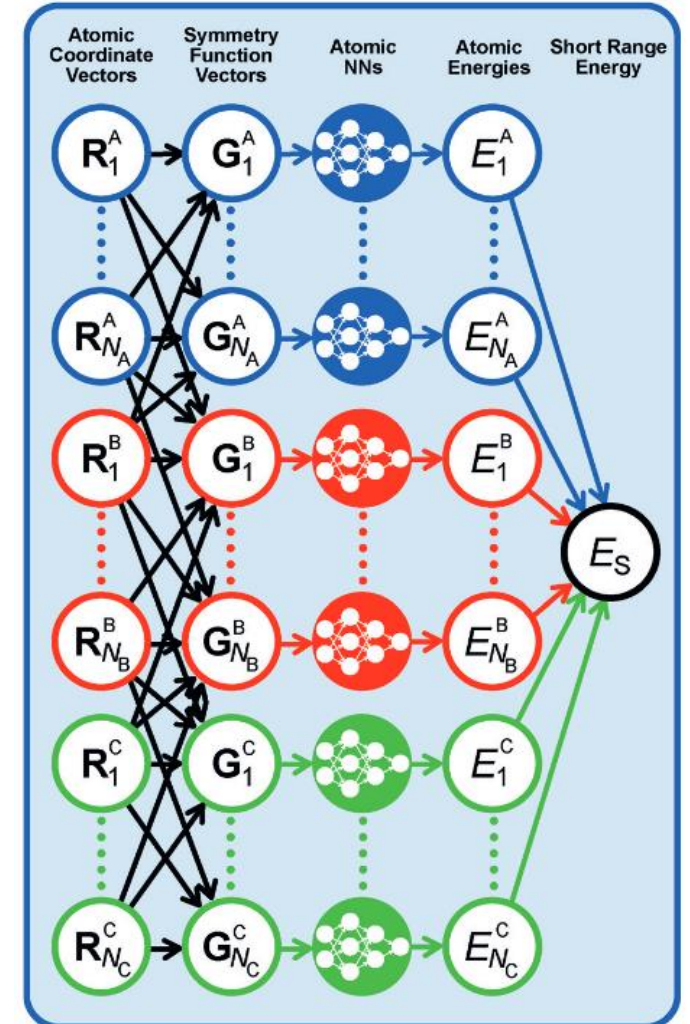
with 48 symmetry functions

Energy error: ~ 5 meV/atom

Force error: ~ 200 meV/Å



Predicted Radial pair correlation (red) compares well with the DFT one (black).



Mattioli-Sciortino-Russo (2023)



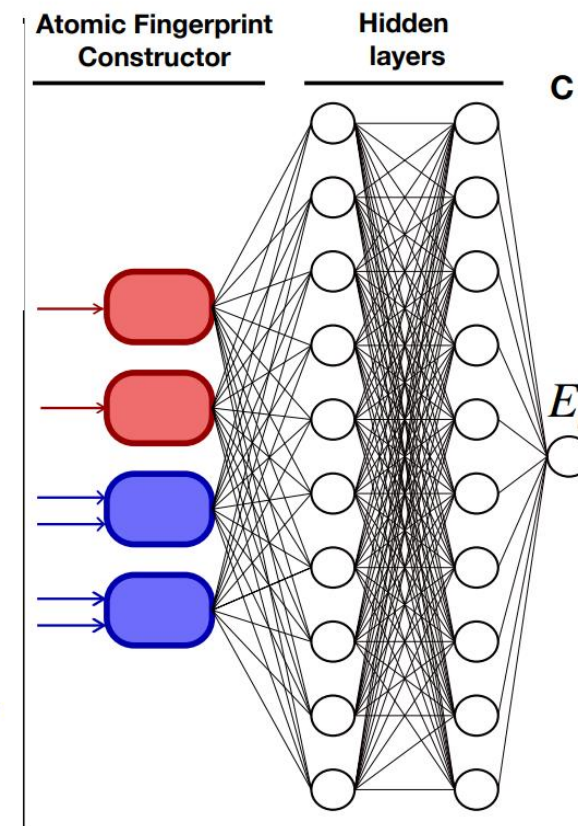
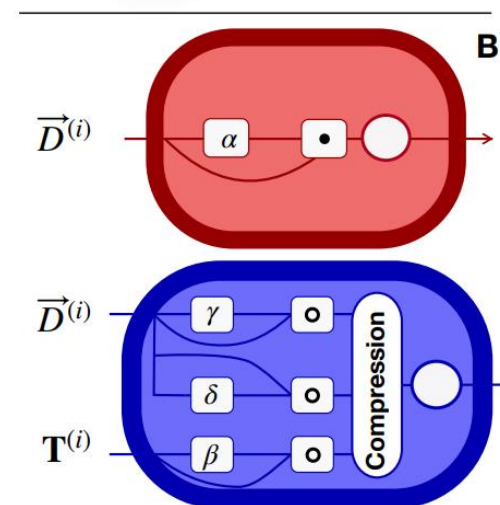
2-body and 3-body descriptors with **learnable** exponential weights.

$$D_j^{(i)}(r_{ij}; R_c) = \begin{cases} \frac{1}{2} \left[1 + \cos \left(\pi \frac{r_{ij}}{R_c} \right) \right] & r_{ij} \leq R_c \\ 0 & r_{ij} > R_c \end{cases}$$

$$T_{jk}^{(i)}(r_{ij}, r_{ik}, \theta_{jik}) = \frac{1}{2} [1 + \cos(\theta_{jik})] D_j^{(i)}(r_{ij}; R_c') D_k^{(i)}(r_{ik}; R_c')$$

Inspired by the **attention** mechanism in deep learning.

A MLP with 2 hidden layers of ~25 nodes.

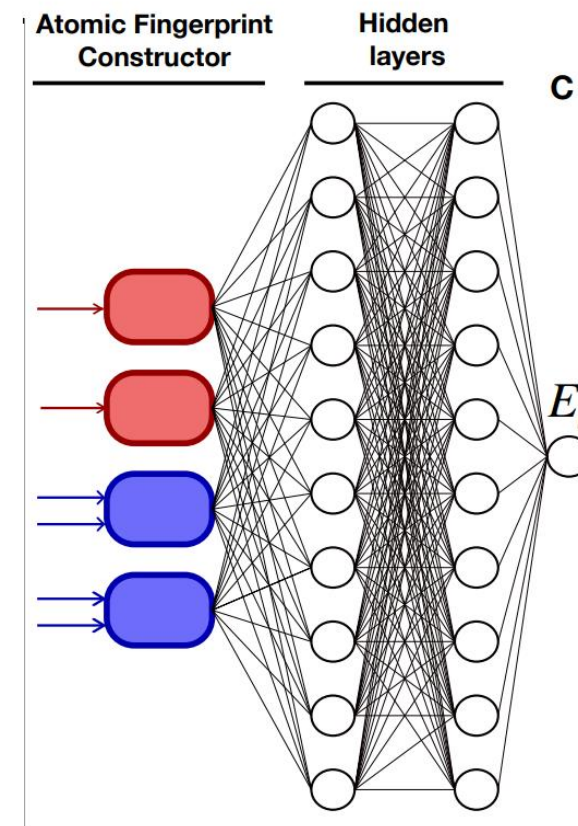
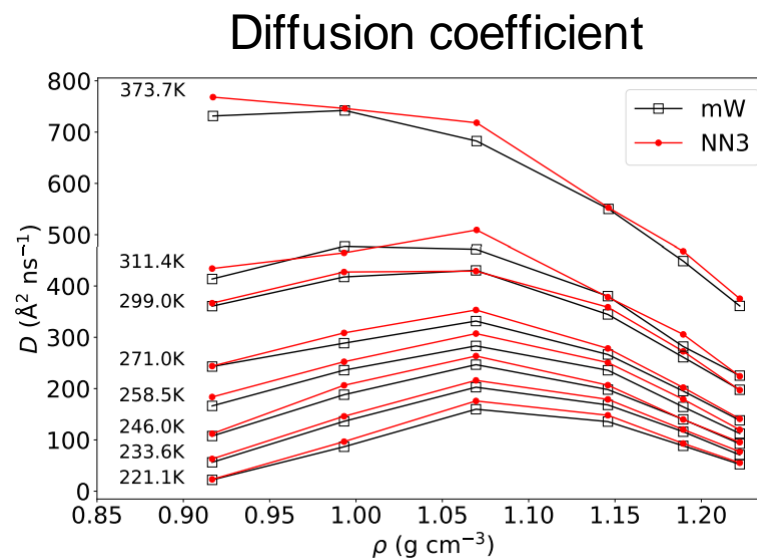
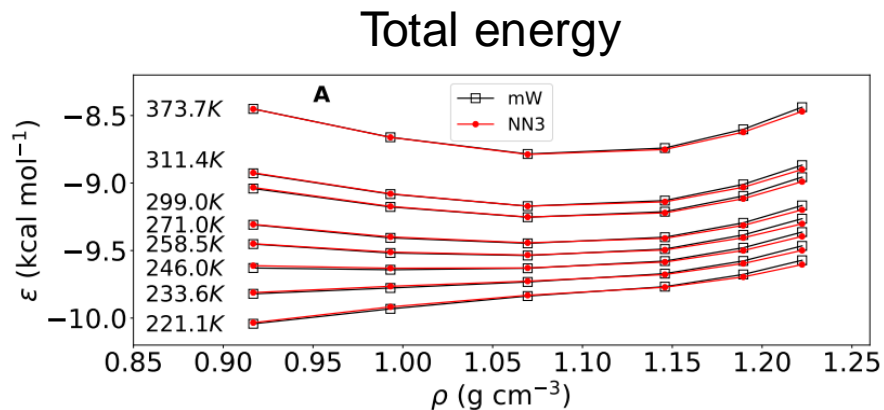
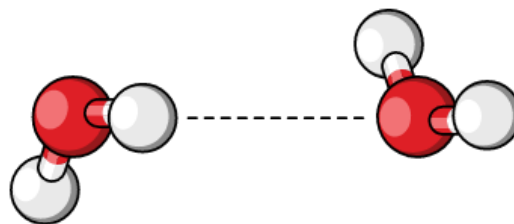


Architecture for a monospecies system

Amazing performance when trained on the “mW” water potential (a classical single-species PES but with non-trivial 3-body interaction)

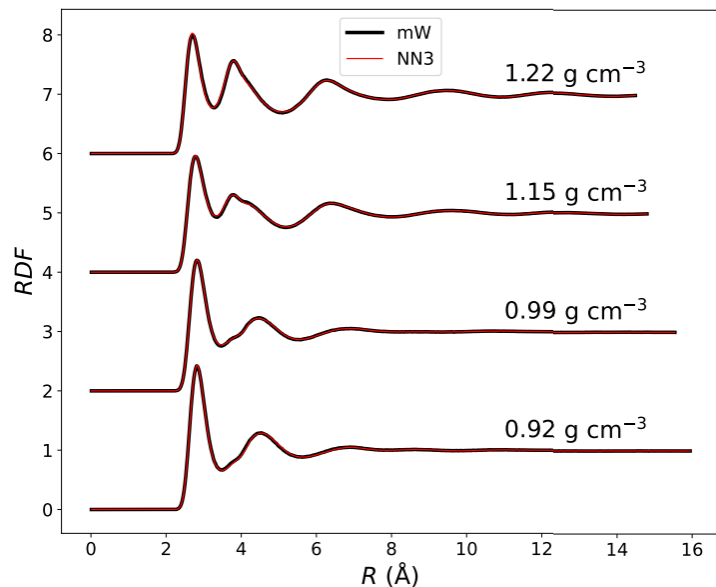
With only 10 atomic descriptors:

- Energy error $\sim 0.4 - 1.3$ meV/atom.
- Force error ~ 6.7 meV/Å.

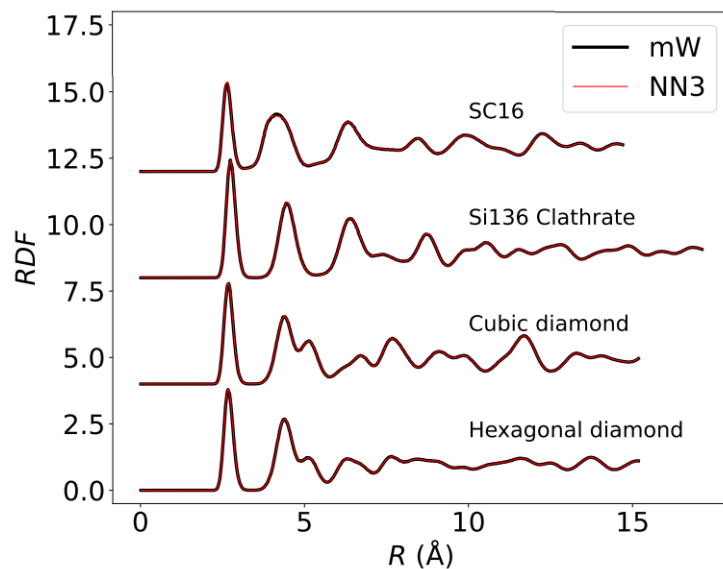


Amazing performance when trained on the “mW” water potential

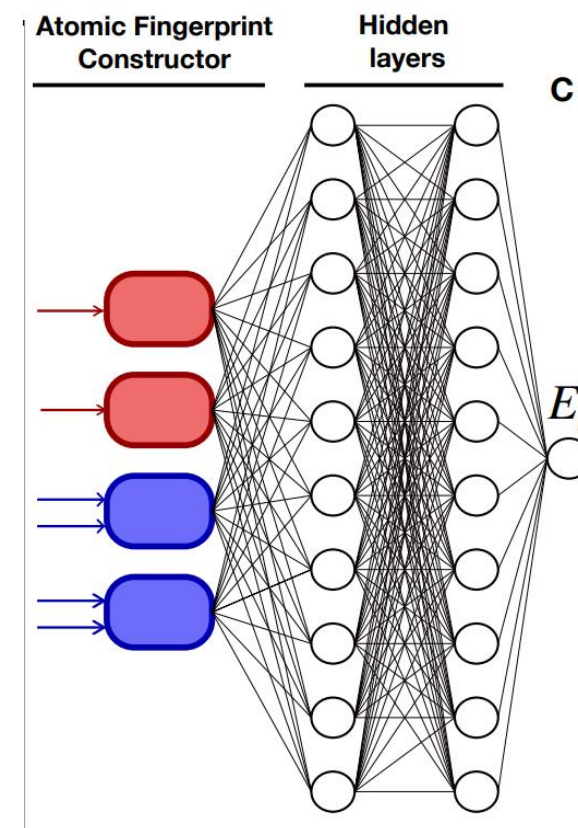
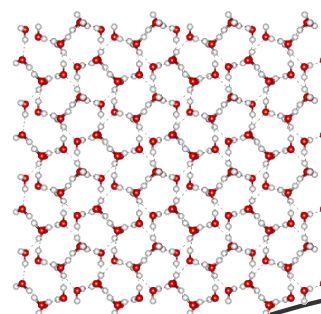
Predicted radial pair correlations perfectly match the real ones:



At different densities...



... and even in extrapolated crystal configurations!



Conclusion and Future work

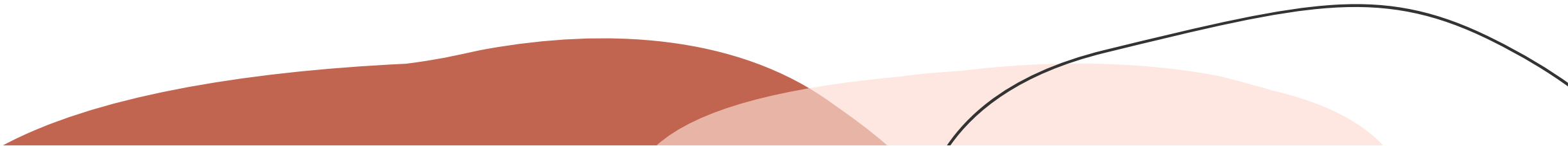
The bottleneck of large-scale quantum-accurate molecular dynamics simulations is the computation of forces.

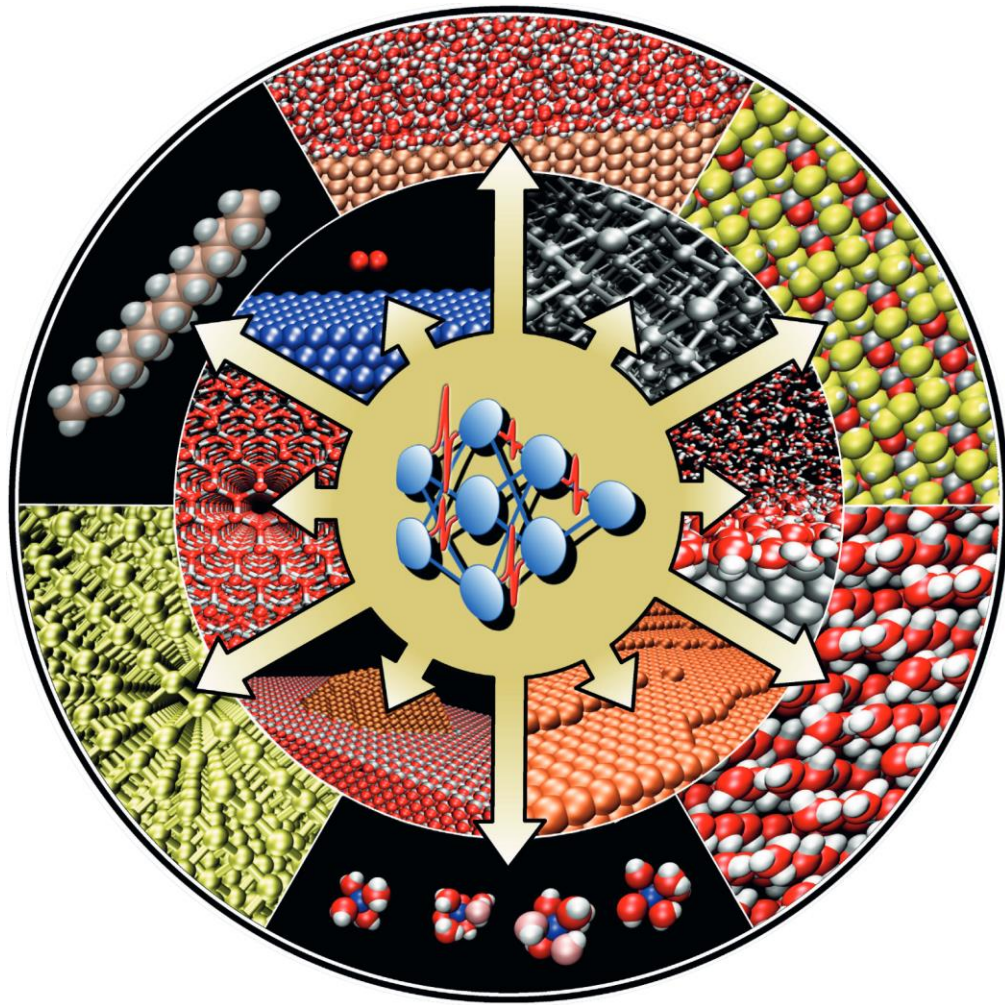
Neural-network parametrization¹ allows to compute forces both accurately and efficiently.

Part of my project is to apply the very recent method by Mattioli et al. [4] to some *Phase-Change Materials* (alloys of Ge, Sb, Te) for large-scale simulations in the supercooled phase.

¹ And other machine-learning methods which I did not cover today.

Thanks for your attention!





References

- 1) Behler, *First Principles Neural Network Potentials for Reactive Simulations of Large Molecular and Condensed Systems*. *Angew. Chem. Int. Ed.* (2017, review).
- 2) Frenkel, Smit, *Understanding Molecular Simulations*. (1996).
- 3) Behler and Parrinello, *Generalized Neural-Network Representation of High-Dimensional Potential-Energy Surfaces*. *Phys. Rev. Lett.* (2007).
- 4) Mattioli, Sciortino, Russo, *A neural network potential with self-trained atomic fingerprints: a test with the mW water potential*. *J. Chem. Phys.* (2023).