Motivation

Deep Potential Molecular Dynamics: a scalable model with the accuracy of quantum mechanics - A summary [ZHW+18]



AIMD: First principles, quantum mechanics, DFT. Computational costs: 100 ps, 100s of atoms, empirica FF: based on approximations and experimental dataData: Atomic configurations and corresponding potential energies, learn this functional dependencereal system invariant to permutations, translations, rotations, so model needs too. Auxiliary quantities: Symmetry functions (describe local geometric environment, problem: ad hoc), coulomb matrix entries: distinct inverse distances between all atoms, problem: extened periodic systems

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Convection: transfer of heat by the movement of the fluid Natural: motion generated by density differences, not by external sources Applications: Nuclear reactor design, cooling of electronic eqiupment, heating and ventialtion in the design of buildings Goal: Identify different flow patterns and investigate which method is suited better

-The model

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Embedded Atom Concept: Energy sum of functions of seperation of atom and neighbors. Reciprocal distances: Reduce weight of distant neighbors-Figure: Water, e  $\times$  O-H bond, e z perpendicular to plane of water molecule other option for D i j angular information as well  $\times$  ij cartesian coordinates of R i j

The model

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Local coordinate frame for each atom with atom at center
 D<sub>I</sub> = (D<sub>ij</sub>)<sub>j</sub> input for NN, where j covers all neighbors within cutoff radius R<sub>c</sub>, sorted by chemical species, inverse distances



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➤ Theoretical foundation: Embedded Atom Concept

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2021-02-02





NN: fixed input size, append 0s potentially

 Fully connected NN consisting of 5 hidden layers with 240, 120, 60, 30, 10 units resp.
 tanh activation for hidden layers and identity for output



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Adam optimizer



NN: fixed input size, append 0s potentially

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Virial: comparison of kinetic, potential energy

Deep Potential Molecular Dynamics: a scalable modeloss function

Virial: comparison of kinetic, potential energy

 $\mathcal{L}(\rho_i,\rho_f,\rho_f) = \rho_i \Delta i_f^2 + \frac{\rho_f}{10} \sum_i |\Delta f_i^2|^2 + \frac{\rho_i}{\eta} |\Delta \Xi|^2$   $\Delta i \ \, \text{Difference between prediction and training data. N. in unboth of storm, e.; inverge per atom., <math>f_i : \text{Force on atom} i : \Xi : \text{Violal tensor} \ \Xi = -\frac{1}{2} \sum_i \beta_i \otimes f_i^2$ . However, the prediction of forces and volt leaves for regularization  $\text{Parameters} \ \rho_i, \rho_i \ \, \text{Standily increased during training}, \rho_i \ \, \text{standily discussed}$ 

Virial: comparison of kinetic, potential energy

Deep Potential Molecular Dynamics: a scalable modelos function

Deep Potential Molecular Dynamics: a scalable model

Deep Potential Molecular Dynamics: a scalable model with the accuracy of quantum mechanics - A summary [ZHW+18]

Advantages:

High accuracy: Similar performance to AIMD simulations for water, slightly better than GDML benchmark for molecules

Summary

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Deep Potential Molecular Dynamics: a scalable model	
with the accuracy of quantum mechanics - A	Advantages:  High accuracy: Similar performance to AIMD simulations for water, slightly better than CDML benchmark for molecules
summary [ZHW <sup>+</sup> 18]	Scalable and parallelisable: Computational costs increase only linearly with the number of atoms     Simplicity: No auxiliary quantities as opposed to other ML methods
Summary	► 3 implicity: two autoriary quantities as opposed to other rest. methods

Deep Potential Molecular Dynamics: a scalable model	
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• High sociols' Smiller performance to AIMD simulations for water, slightly letter than CDML benchmark for molecular.

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