

Physical Chemistry Seminar

Accurate predictions of QM/MM energies using a modified long-range ANI/MM neural network potential

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ANI is one of the promising machine-learned Neural Network Potentials (NNPs) that replace computationally expensive ab initio calculations by learning the complex correlations between atomic coordinates and electronic structure properties¹. Such NNPs calculate the final energy as a sum of local environmental energies of each atom, leading them to be widely transferable to almost any molecule but at the cost of losing long-range interactions like electrostatic effects. This is due to the usage of Behler-Parinello symmetry functions to vectorize the molecules, which have a cutoff that ignores all interactions further than 5.2 Angstroms. In our previous work, we designed an ANI-1x-based network that could predict the Coulomb and polarization components of the solute-solvent interaction energies and trained it on a set of small, solvated molecules². This new ANI/MM network takes electrostatic potentials on each atom computed with the Effective Fragment Potential as an additional input feature. We extend the previous work to show how to utilize the ANI/MM developed network for predicting protein-ligand interaction energies. Here, we trained it on QM/MM energies for two protein-ligand systems, 2VTL/LZ5 and 4EK5/03K. The ANI/MM network predicts both energies and forces with singular kcal/mol accuracy.

References:

1. Gao, X.; Ramezanghorbani, F.; Isayev, O.; Smith, J. S.; Roitberg, A. E. TorchANI: A Free and Open Source PyTorch-Based Deep Learning Implementation of the ANI Neural Network Potentials. *J. Chem. Inf. Model.* **2020**, 60 (7), 3408–3415. <https://doi.org/10.1021/acs.jcim.0c00451>.
2. Haghiri, S.; Viquez Rojas, C.; Bhat, S.; Isayev, O.; Slipchenko, L., ANI/EFP: Modeling Long-Range Interactions in ANI Neural Network with Effective Fragment Potentials, *J. Chem. Theory Comp.* **2024**. DOI: 10.1021/acs.jctc.4c01052.



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