

# Assessing Conformer Energies: Machine Learning vs Conventional Quantum Chemistry

Dakota Folmsbee and Geoffrey Hutchison  
Department of Chemistry, University of Pittsburgh

Funding: NSF CHE-1800435

## Motivation

Understanding and predicting thermodynamically accessible conformations is a key task as the number of possible minima increase exponentially for each rotatable bond. This becomes expensive for wavefunction and density functional first principles methods leading to a desire of a fast and accurate method, such as machine learning (ML), to be used as a surrogate for these time-consuming calculations.

## Methodology

- Conformer dataset consists of ~6000 DLPNO-CCSD(T) single points of ~700 small molecules with up to 10 conformers each containing atoms of H, C, N, O, F, P, S, Cl, and/or Br.
- Evaluated 39 methods including dispersion and non-dispersion corrected DFT and normalized ML methods.

## Performance

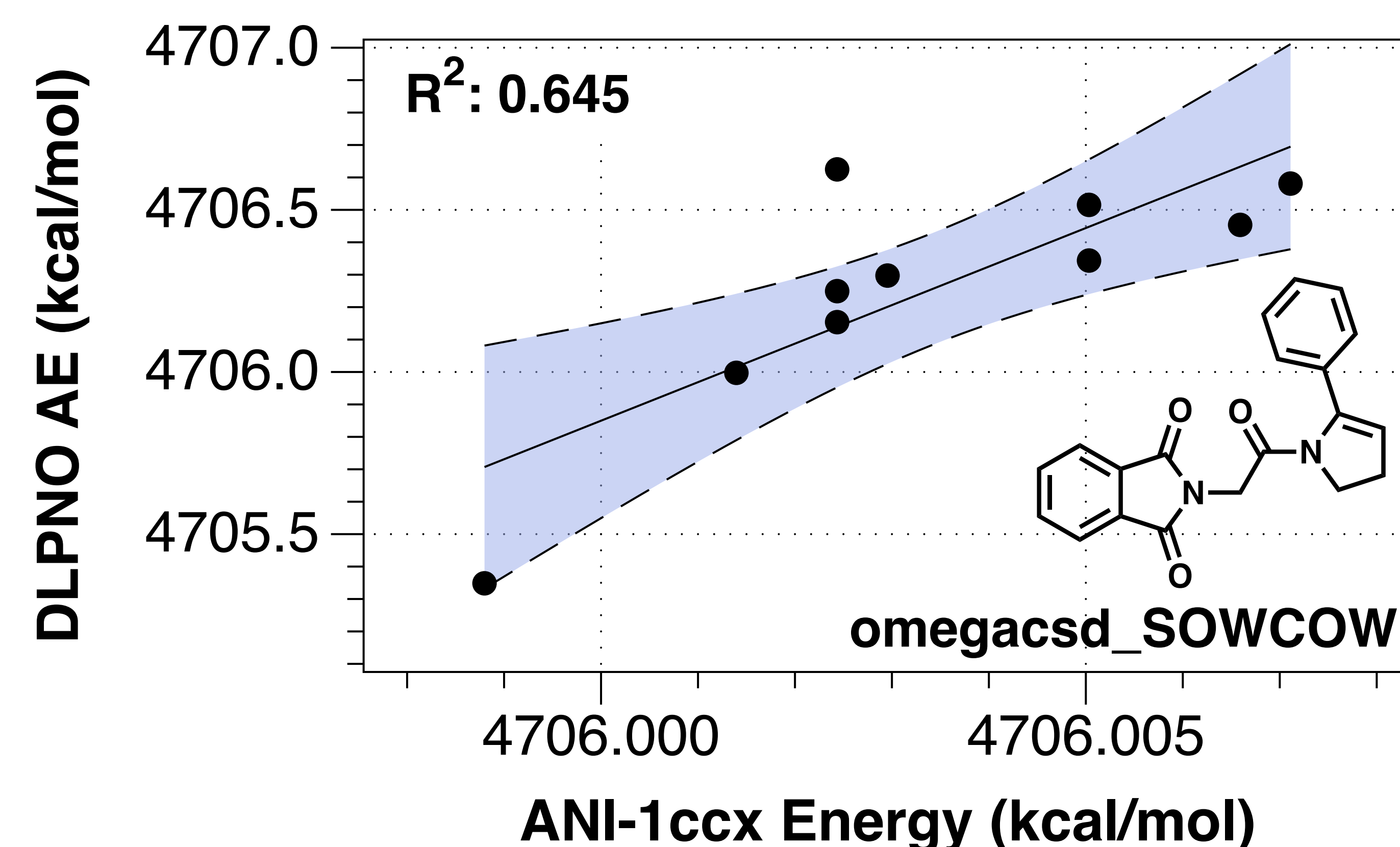
- RI-MP2 and DFT dispersion methods perform best at the cost of time while ML and semiempirical methods look to be an even ground between accuracy and time.
- Most methods are dominated by random errors as measured with  $R^2$ .
- A few force field and ML methods show increased MARE from systematic errors.

## Thanks

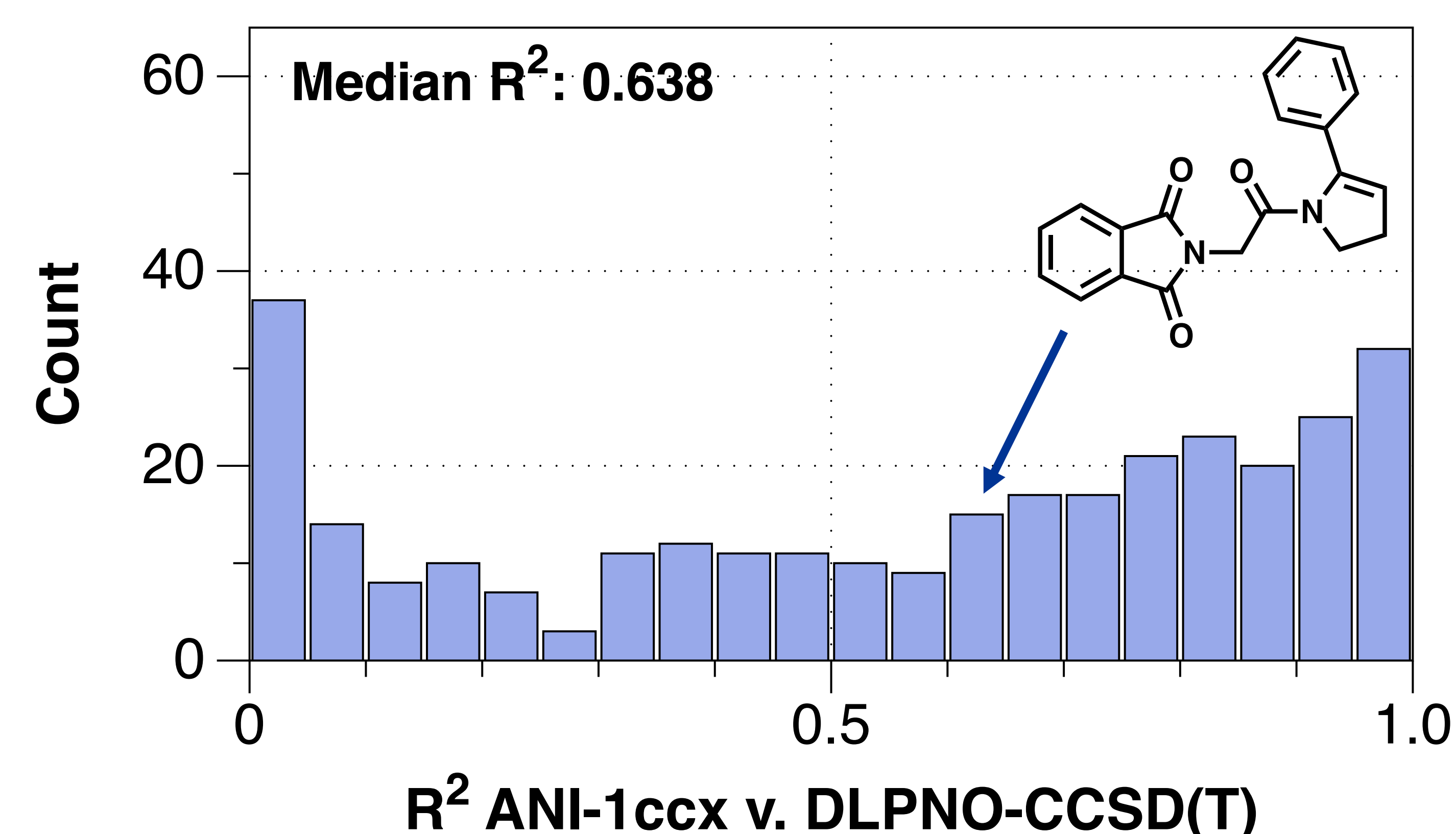
This research was supported by the NSF and the University of Pittsburgh Center for Research Computing. We also thank Olexandr Isayev and Justin Smith for access to the ANI-2x model.

**ML methods show promise with comparable accuracy to semiempirical methods and can be accelerated on GPUs. We expect these methods to provide increasing accuracy.**

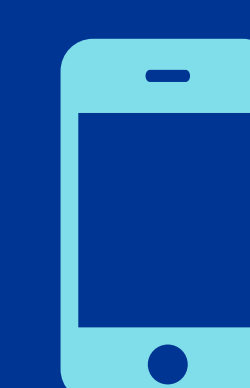
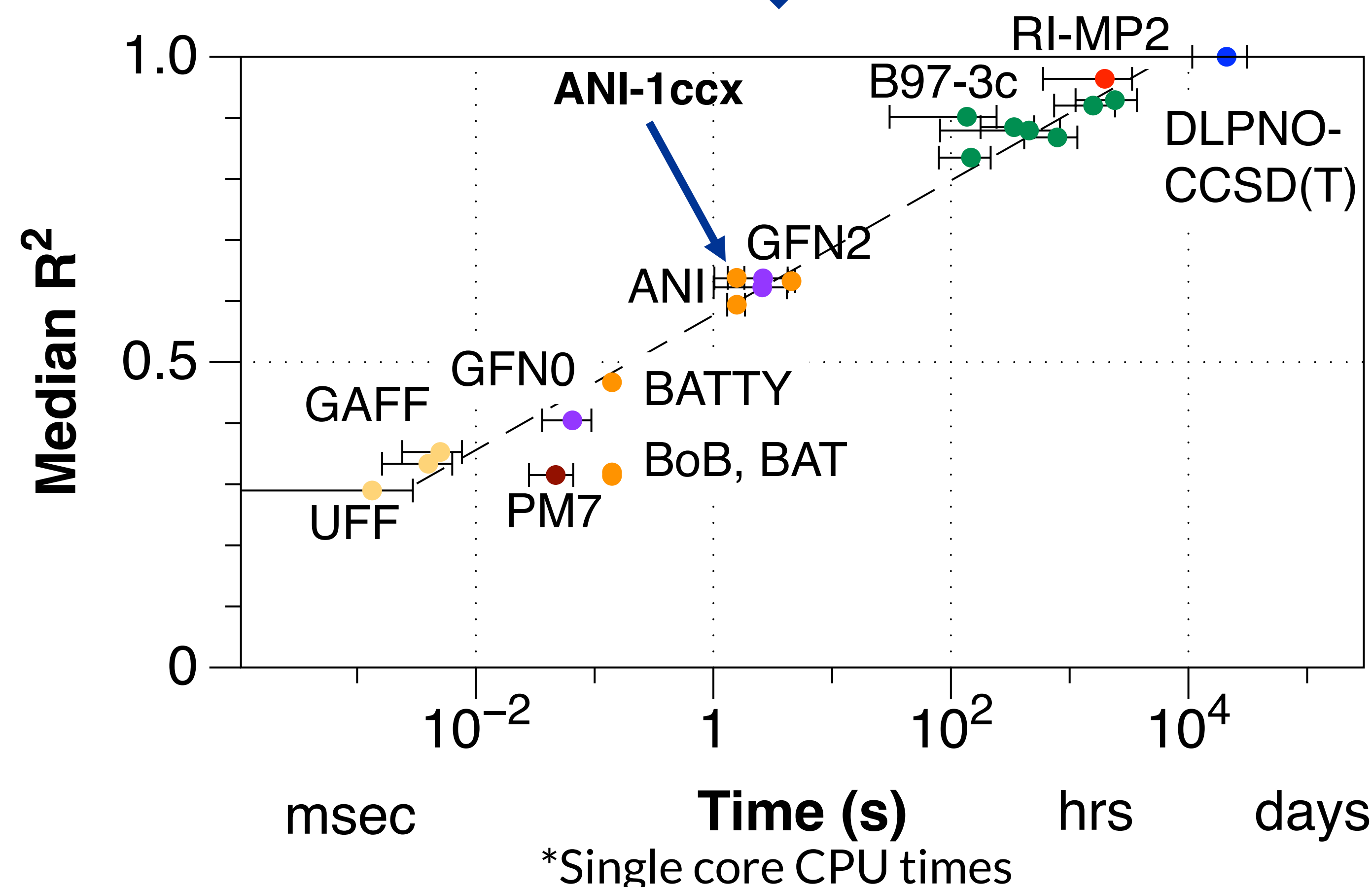
Across 10 Conformers



Across 700 Molecules



Across 39 Methods



Take a picture to  
download the full paper

