



Award #: 1642336

# SI2-SSE: Enabling Chemical Accuracy in Computer Simulations: An Integrated Software Platform for Many-Body Molecular Dynamics

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## Many-body molecular dynamics: Chemical Accuracy Across Different Phases

$$E_N = \sum_i^N V^{1B}(i) \quad \text{1-body}$$

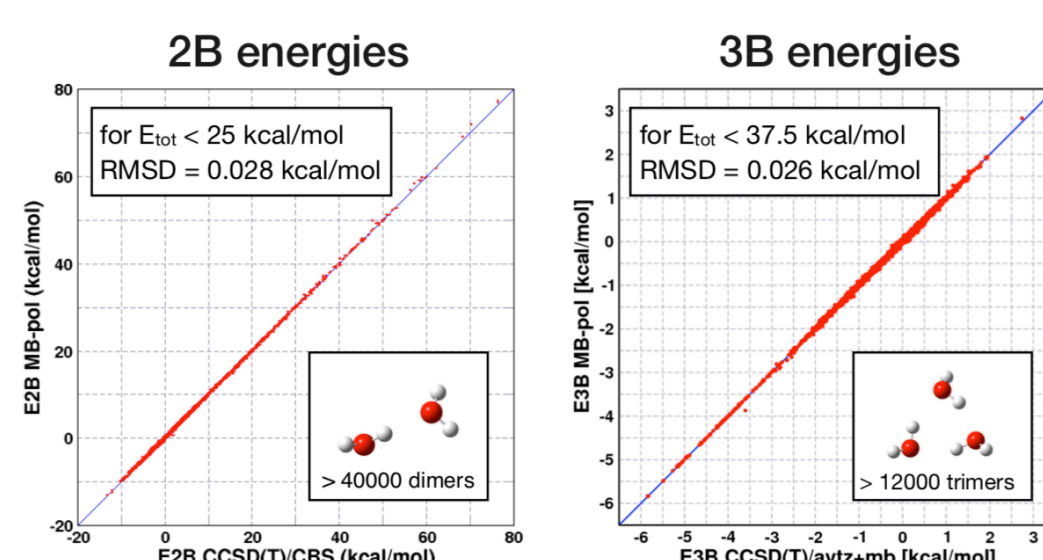
$$+ \sum_{i<j}^N V^{2B}(i,j) \quad \text{2-body}$$

$$+ \sum_{i<j<k}^N V^{3B}(i,j,k) \quad \text{3-body}$$

$$+ \dots$$

$$+ V^{NB}(1, \dots, N) \quad \text{N-body}$$

explicit 1B term  
+  
N-body induction  
+  
explicit 2B and 3B terms  
from CCSD(T)/CBS



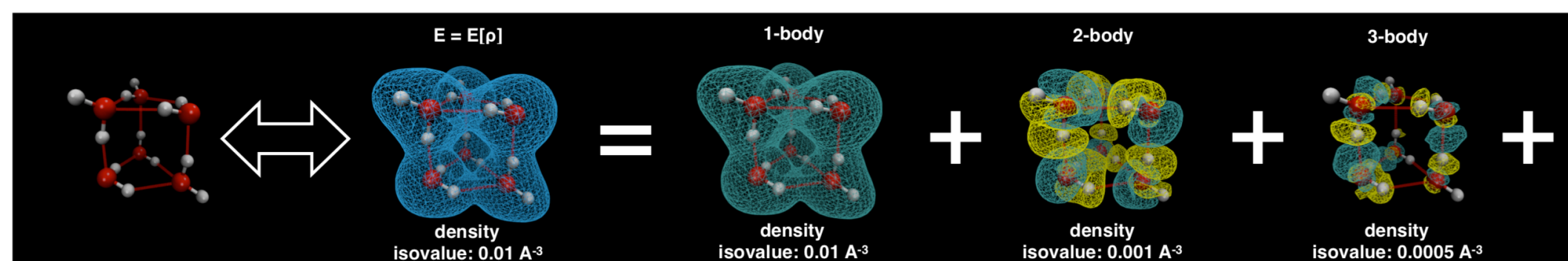
### Key features

- No recourse to empirical parameters
- Potential energy surface entirely derived from highly accurate correlated electronic structure data
- Enables predictive molecular simulations from gas phase to the condensed phase

### Goal of this project

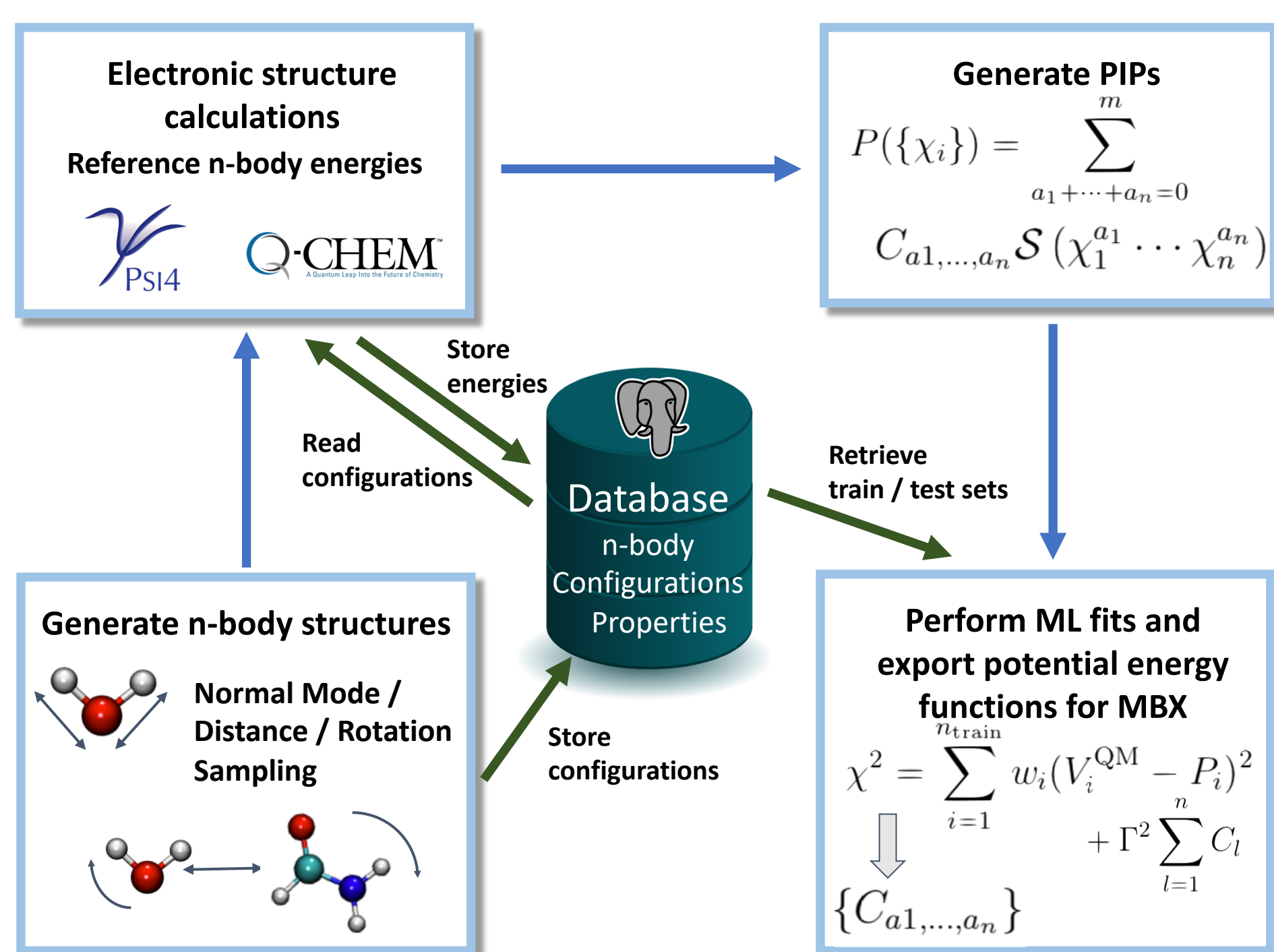
- Make MB-MD methodology widely available to a broad scientific user community

MB-MD in OpenMM:

[http://paesanigroup.ucsd.edu/software/mbpol\\_openmm.html](http://paesanigroup.ucsd.edu/software/mbpol_openmm.html)


- J. Phys. Chem. Lett.* **3**, 3765 (2012)
- J. Chem. Theory Comput.* **9**, 1103 (2013)
- J. Chem. Theory Comput.* **9**, 4844 (2013)
- J. Chem. Theory Comput.* **9**, 5395 (2013)
- J. Chem. Theory Comput.* **10**, 1599 (2014)
- J. Chem. Theory Comput.* **10**, 2906 (2014)
- J. Chem. Theory Comput.* **11**, 1145 (2015)
- J. Chem. Phys.* **145**, 194504 (2016)

## MB-fit: Workflow system for machine learning of many-body potential energy functions



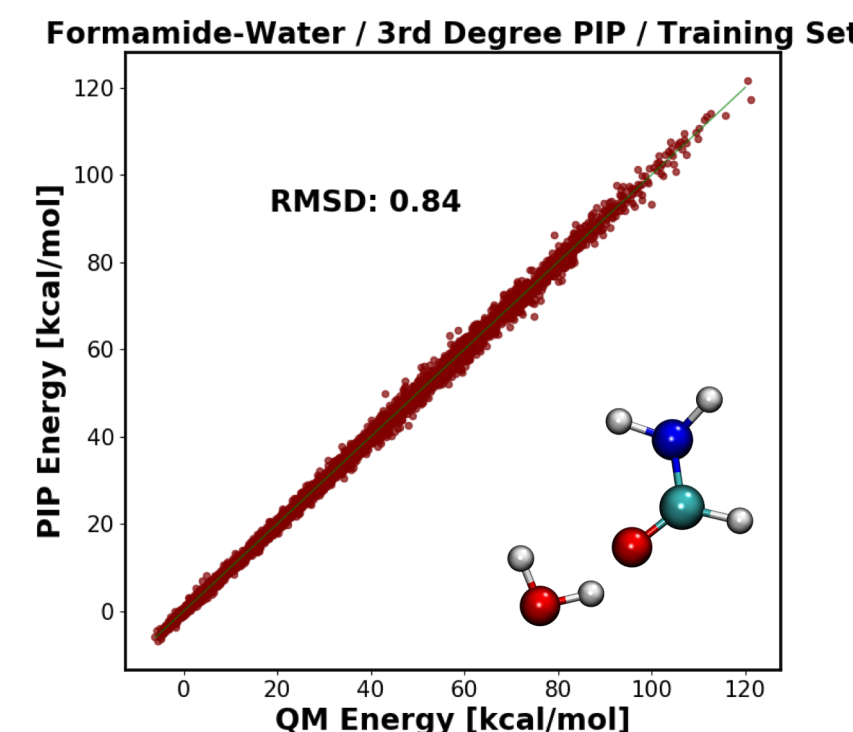
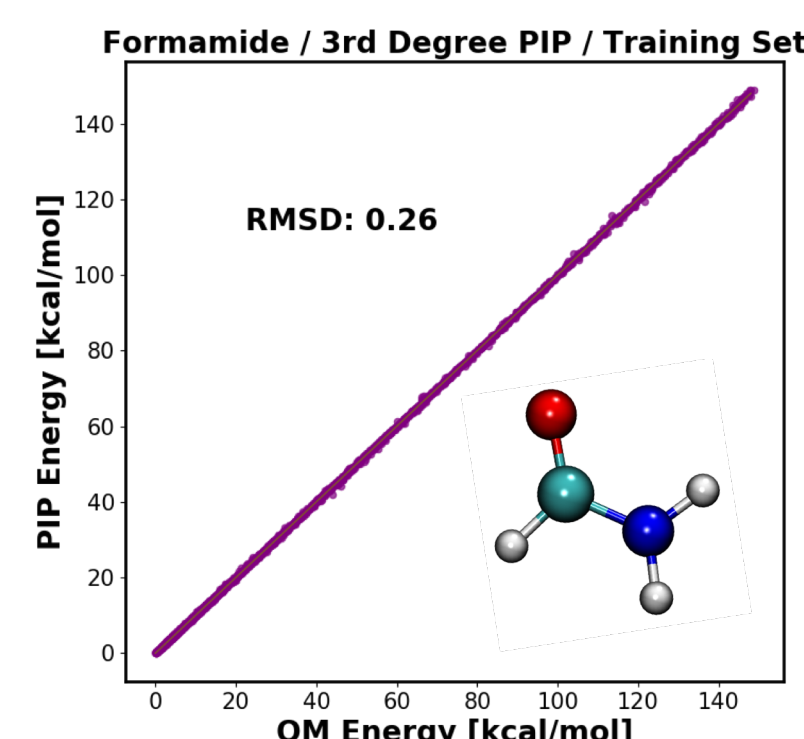
### Features

- Python based workflow system
- Jupyter notebooks for user documentation
- Centralized data storage via PostgreSQL
- Reference n-body structure generators
- Drivers for electronic structure calculations
- Efficient generator for permutationally invariant polynomials (PIPs)

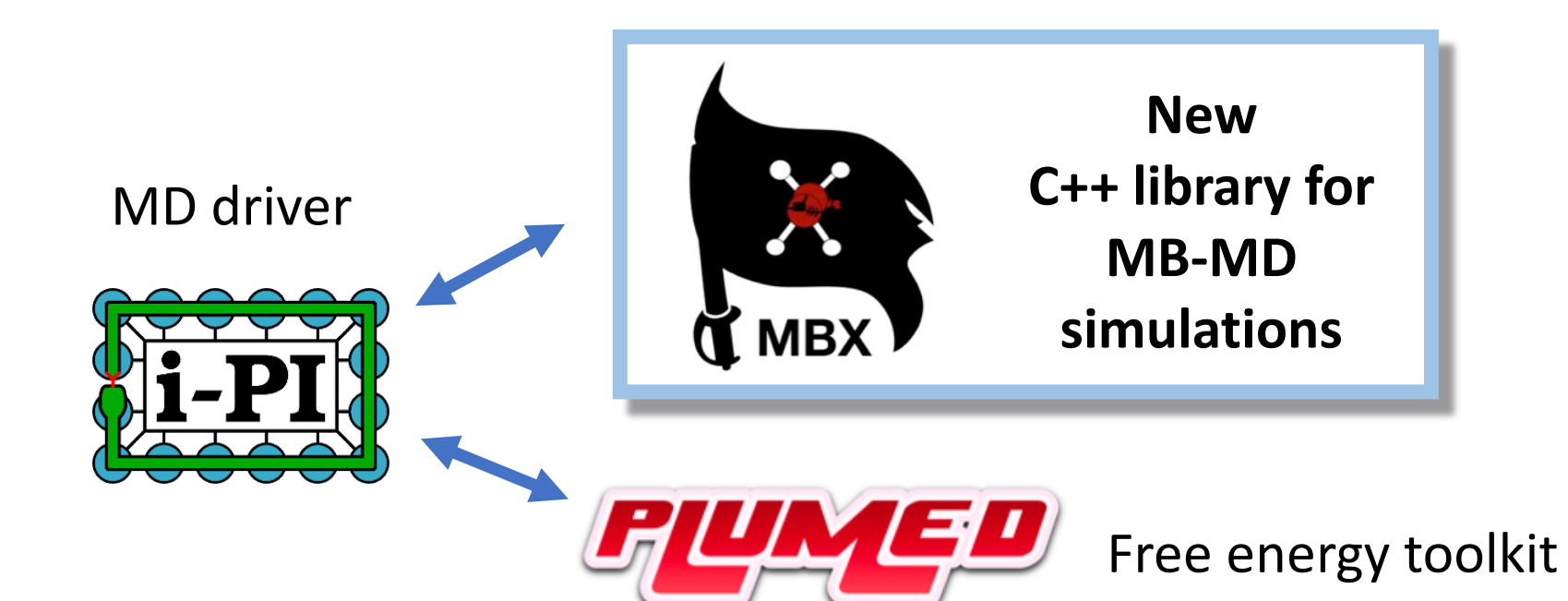
Permutationally invariant polynomials (PIPs)

$$\chi_i = \exp(-k_i d_i) \quad \text{or} \quad \chi_i = \exp(-k_i d_i) / d_i$$

$$P(\{\chi_i\}) = \sum_{a_1+\dots+a_n=0}^m C_{a_1,\dots,a_n} S(\chi_1^{a_1} \dots \chi_n^{a_n})$$



## MBX: High-performance C++ / OpenMP code for MB-MD simulations



### Features

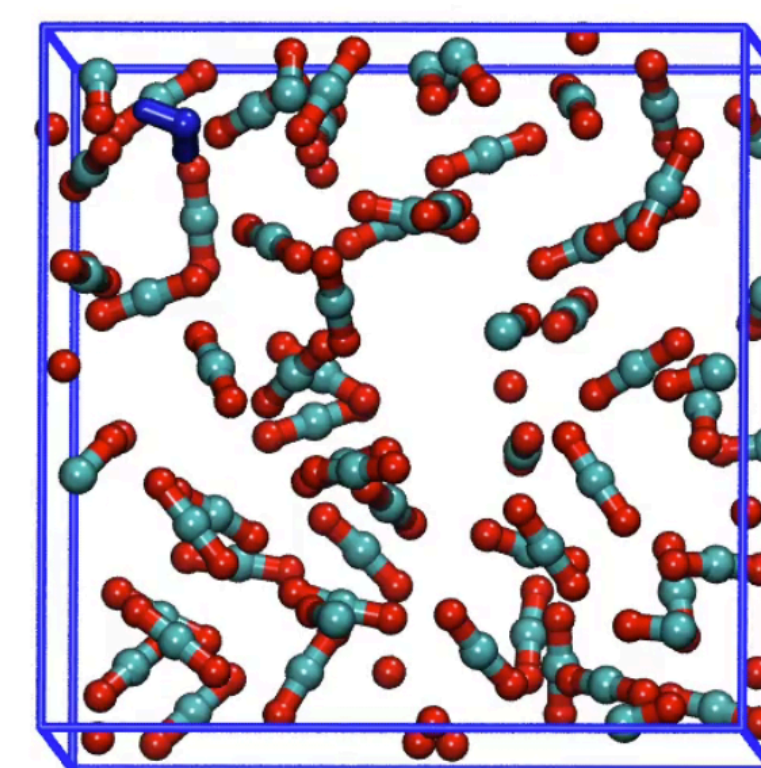
- Modern C++ implementation
- OpenMP parallel
- Periodic boundary conditions
- Long-range electrostatics and dispersion energies via helpME library (<https://github.com/andysim/helpme>)
- User-friendly API for integration with simulation codes
- MD driver via i-PI
- Free energies via PLUMED

### Software available at

<http://paesanigroup.ucsd.edu/software/mbx.html>

### Application example:

MBX powered condensed phase simulation of dilute H<sub>2</sub>O in CO<sub>2</sub> with coupled cluster accuracy (chemrxiv.11288465.v1)



OpenMP  
Enabling HPC since 1997

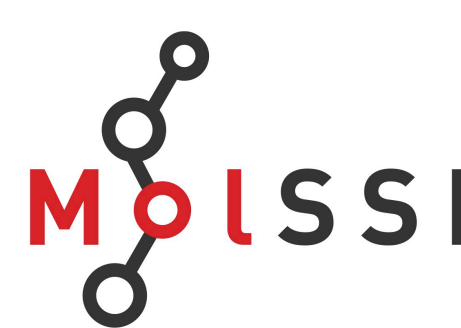
- OpenMP parallel for multi-core CPUs
- MPI support for multi-node clusters in progress

## Broader Impacts

- Training of a large number of postdoctoral scholars, graduate students, undergraduate students and high-school interns in computational research and software engineering
- Open source software tools for predictive molecular simulations for the broader scientific community

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