



Sparse graph attention networks as efficient ionic liquid potentials

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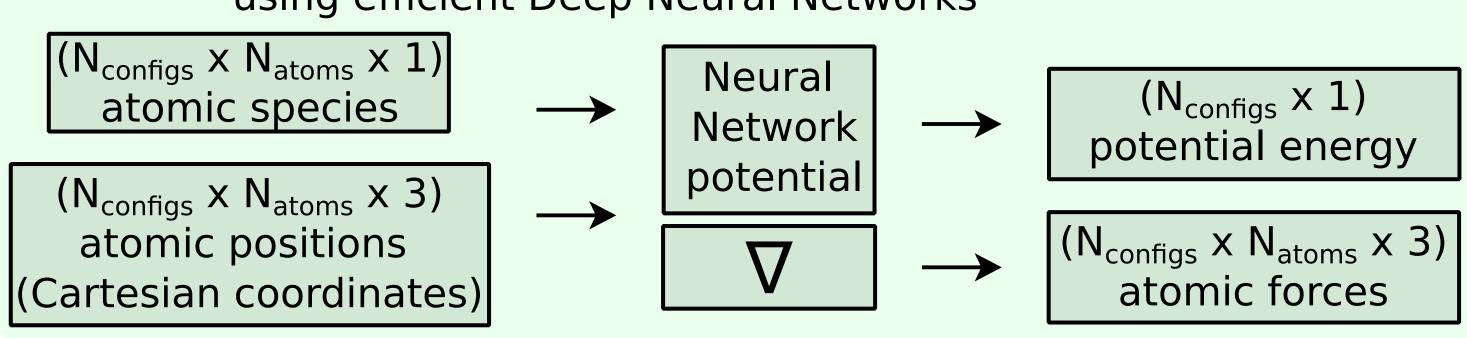
Problem context: What's a potential?

Domain: Computational (molecular) chemistry Predict the energy of molecules Goal: Learn the potential energy surface Task: (atomic positions → energy mapping)

Problem: "Exponential barrier" of electronic contribution, high-dimensional task

(3Natoms-6 degrees of freedom)

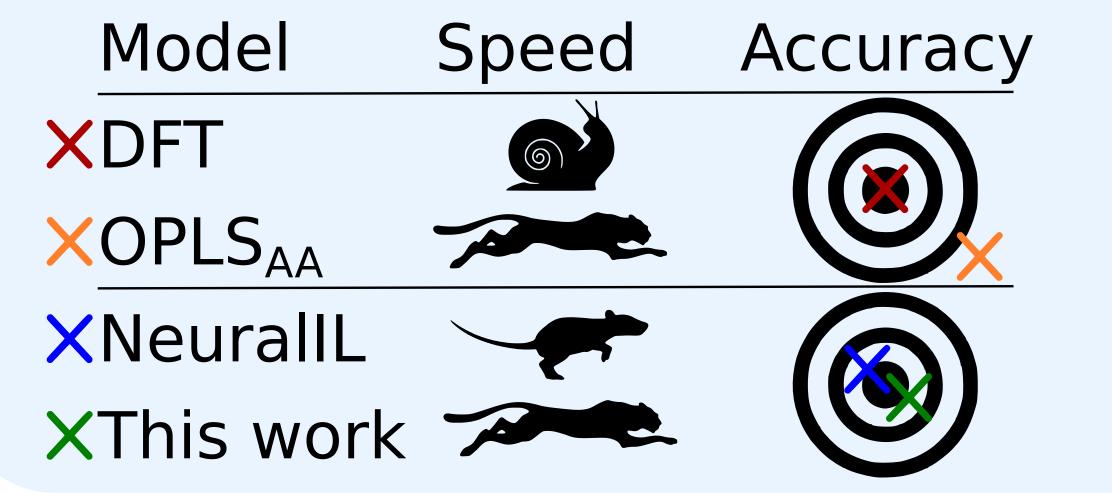
Solution: Approximate the Schrödinger Equation using efficient Deep Neural Networks

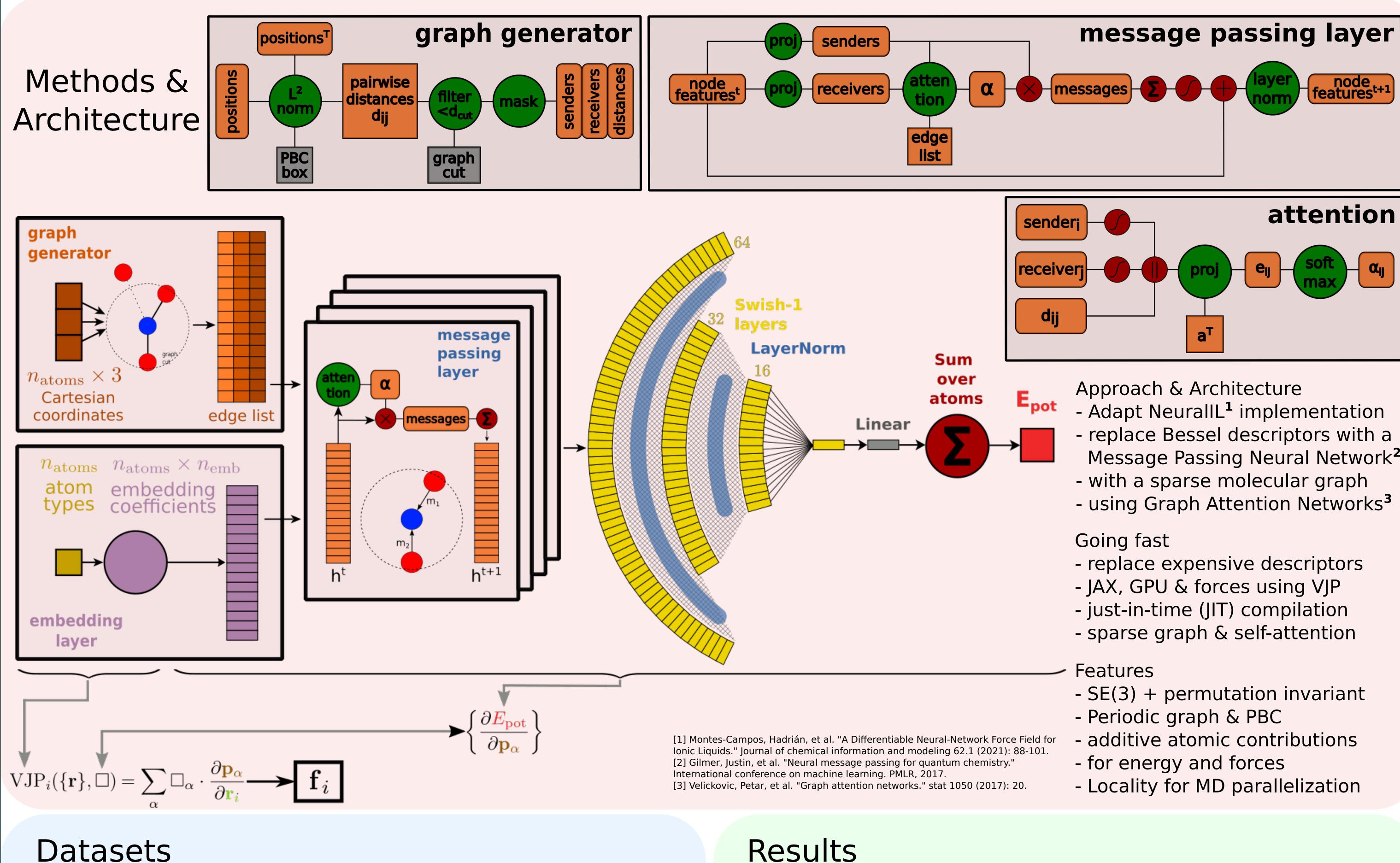


... but why? Molecular dynamics!

Obtain 3N_{atoms} forces as gradient of energy w.r.t. positions

- Use atomic forces to integrate Newton's equations of motion
- requires very small timestep $\Delta t \sim fs$, $(10^{-15} s)$
- 1M+ steps to watch something interesting happen...
- thus 1M+ model evaluations, making efficiency crucial
- model() & grad() for value (energy) and grad (forces)





Approach & Architecture

- Adapt NeurallL¹ implementation
- replace Bessel descriptors with a Message Passing Neural Network²

attention

- with a sparse molecular graph
- using Graph Attention Networks³

Going fast

- replace expensive descriptors
- JAX, GPU & forces using VJP
- just-in-time (JIT) compilation
- sparse graph & self-attention

Features

- SE(3) + permutation invariant
- Periodic graph & PBC
- additive atomic contributions
- for energy and forces
- Locality for MD parallelization

Ionic liquid EAN

- salt in liquid phase at room temperature
- 15 EAN pairs of anion (NO_3^-) & cation $(C_2H_5NH_3^+) \rightarrow 225$ atoms
- 741 configurations sampled from OPLS-AA MD trajectory
- reference energy & forces calculated using DFT
- training on 3N_{atoms} atomic forces

ANI-1

- ~57K organic molecules in displaced states
- max. 29 {C, N, O, H} atoms with 1 8 heavy (non-H) atoms
- ~21M configurations sampled from vibration normal modes
- Sample 5% 10% of subsets 1-7 to train & develop architecture
- final run: 5% of ANI-1, ~900K train, ~100K validation and test - all C₂H₆N₂ and C₄H₄N₄ isomers excluded for separate evaluation
- training only on energy, DFT reference energies

EAN

- best model uses 7 JAT layers, 1 head of dim 48
- 3K epochs in \sim 4:30h on GPU, \sim 100 configurations/sec
- forces MAE ~80-88 meV/Å, RMSE ~140-170 meV/Å
- 4x speed up against NeurallL baseline
- reaches chemical accuracy but does not outperform baseline
- predicts dissociation

ANI-1

- best model uses 5 JAT layers, 4 heads of dim 48
- 80 epochs in ~9h on GPU, ~3300 configuration/sec
- energy MAE \sim 16-30 meV/atom (validation & test set)
- reaches chemical accuracy, especially for larger molecules
- MAE ~42 meV/atom on $C_2H_6N_2$, ~18 meV/atom on $C_4H_4N_4$ - excellent results scaling to larger training sets