



Beyond the Black Box: The potential and problems of equivariant electron densities

Swiss Equivariant Learning Workshop

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Sandia National Laboratories



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Who am I?



TH
A



Nation

ship

3

Who am I?



Physics and Political
Philosophy



High School Physics and
Chemistry



Biophysics Ph.D.



Truman Fellowship



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The Grand Challenge of Molecular Modeling

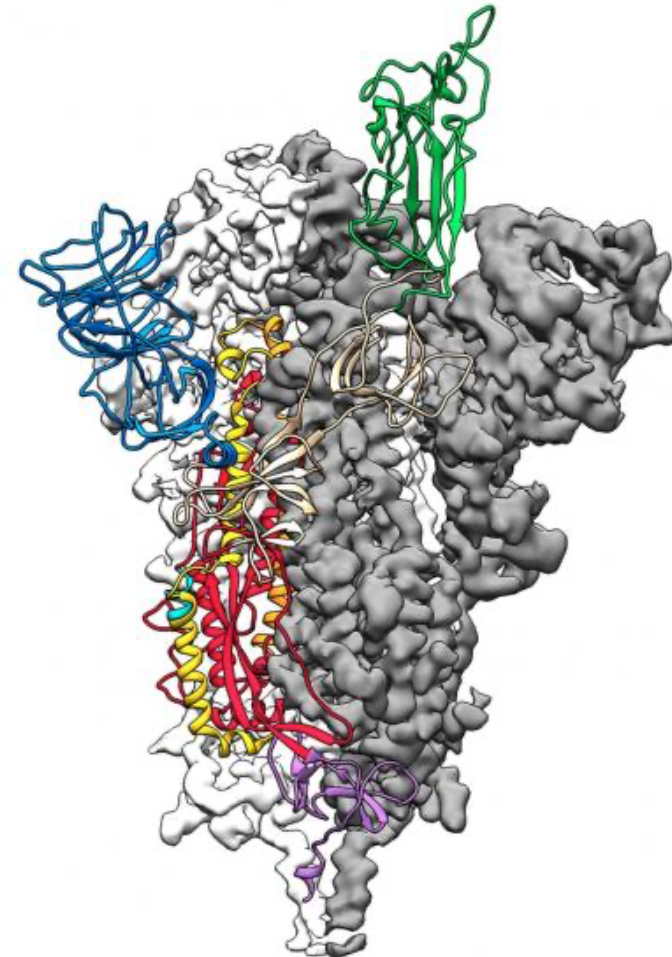
Can we **predict** the behavior of molecules at the **atomic scale**?



$$F = ma$$



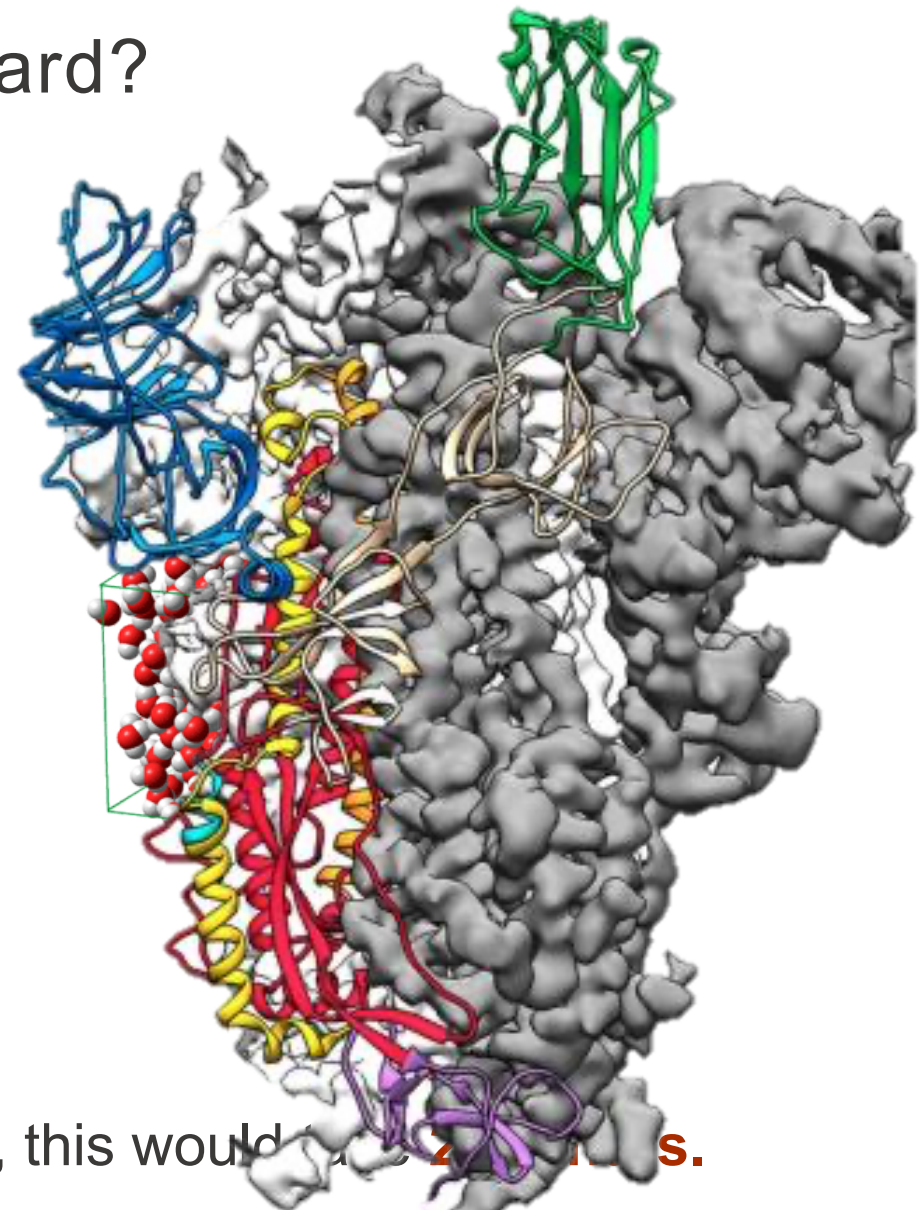
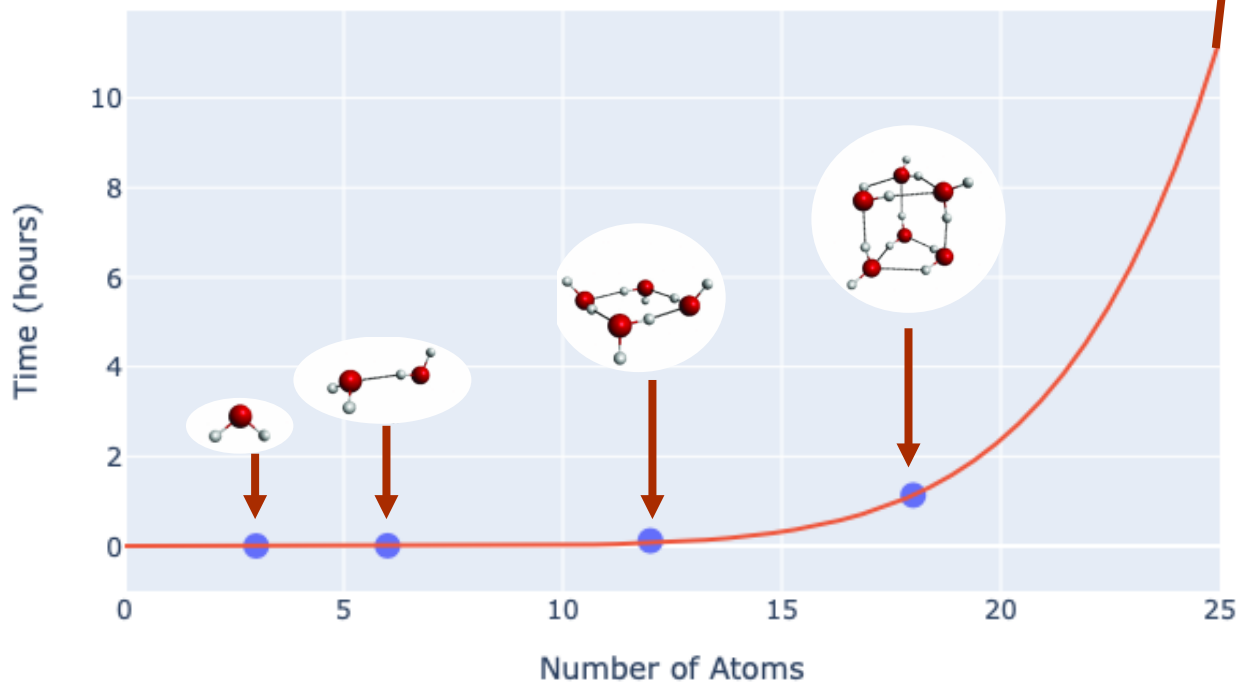
$$H\Psi = E\Psi$$



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Why is doing Quantum Mechanics hard?

Solving $H\Psi = E\Psi$ scales horrendously!



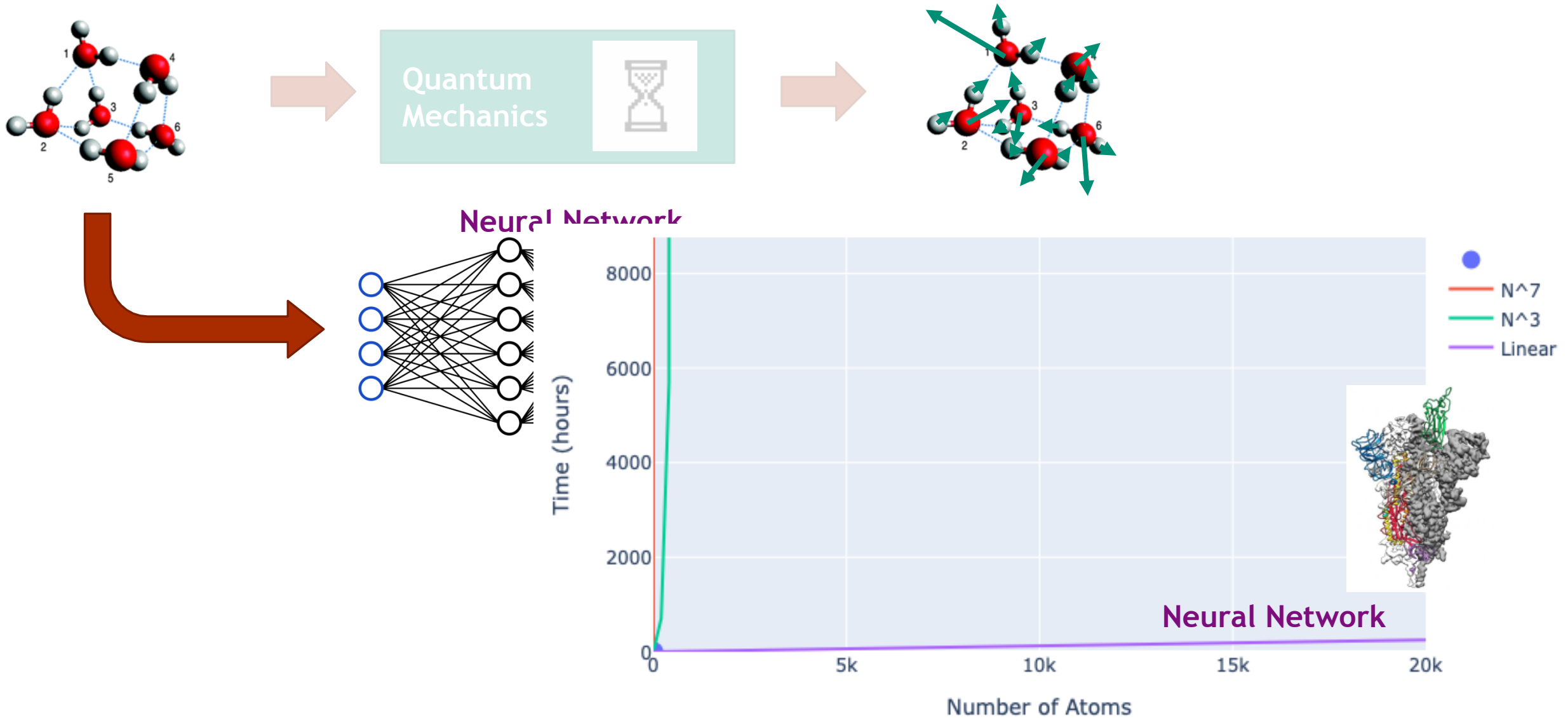
For **50 atoms**, this would take **2000 hours**.

For **100 atoms**, this would take **20 years!!!**



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Let's use Machine Learning!

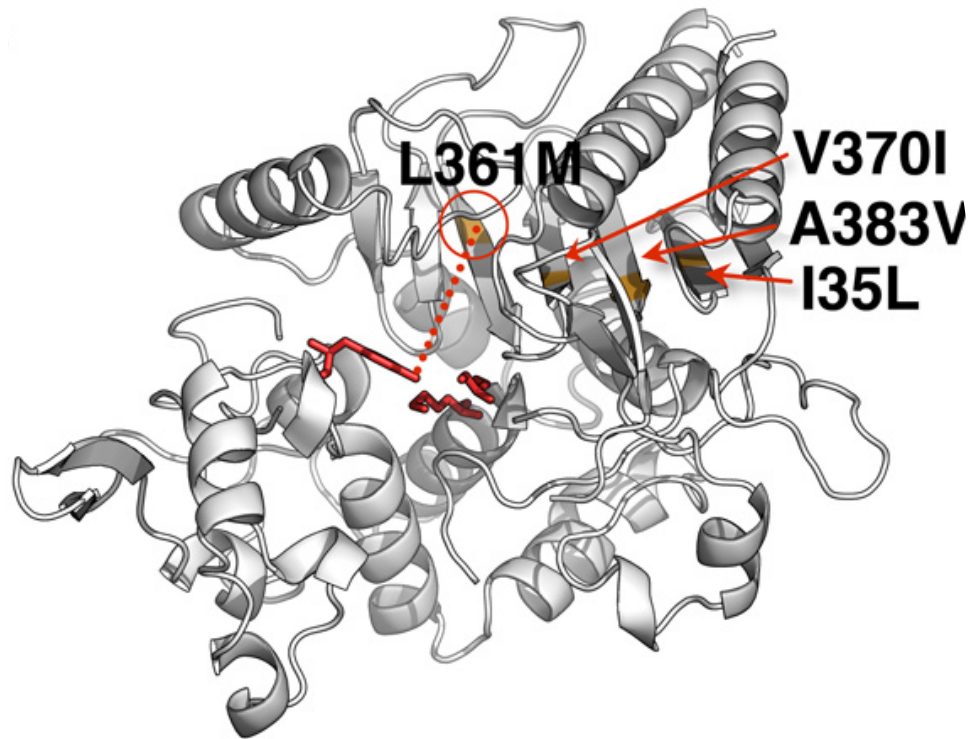




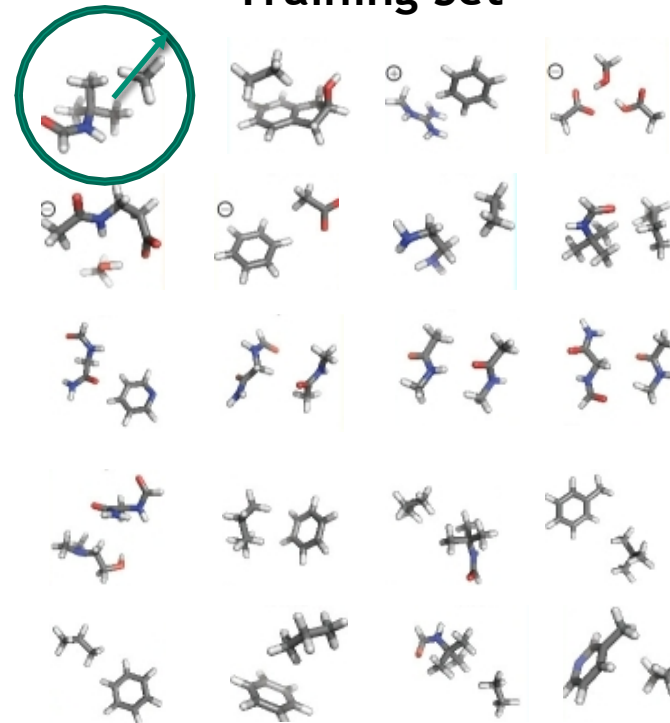
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What's the Problem? – Part 1: Training Data

We want to use ML on large molecules, but we can't train on things that big!



Training Set



Effective Radius
10 Å



What's the Problem? – Part 1: Training Data

CHEMICAL REVIEWS

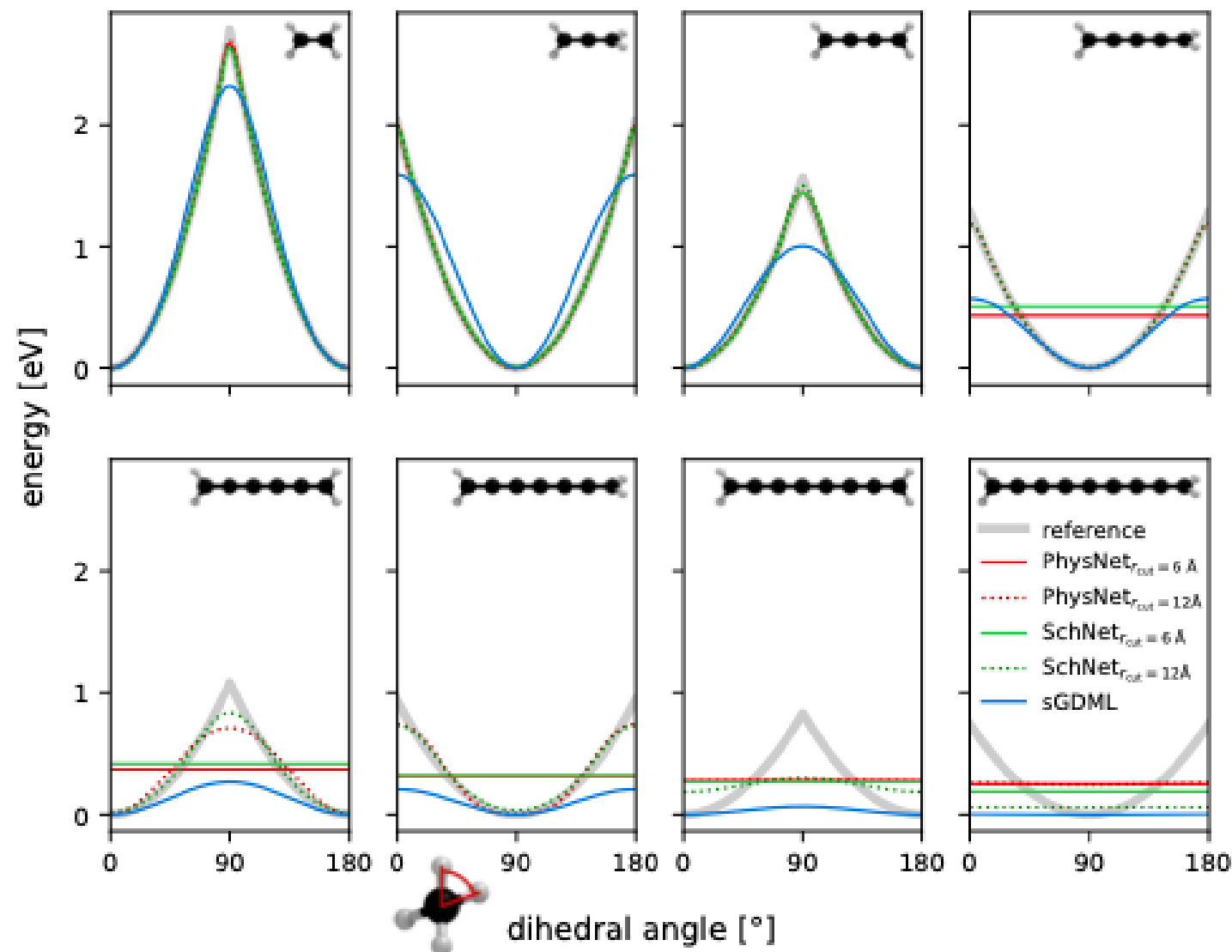
pubs.acs.org/CR



Review

Machine Learning Force Fields

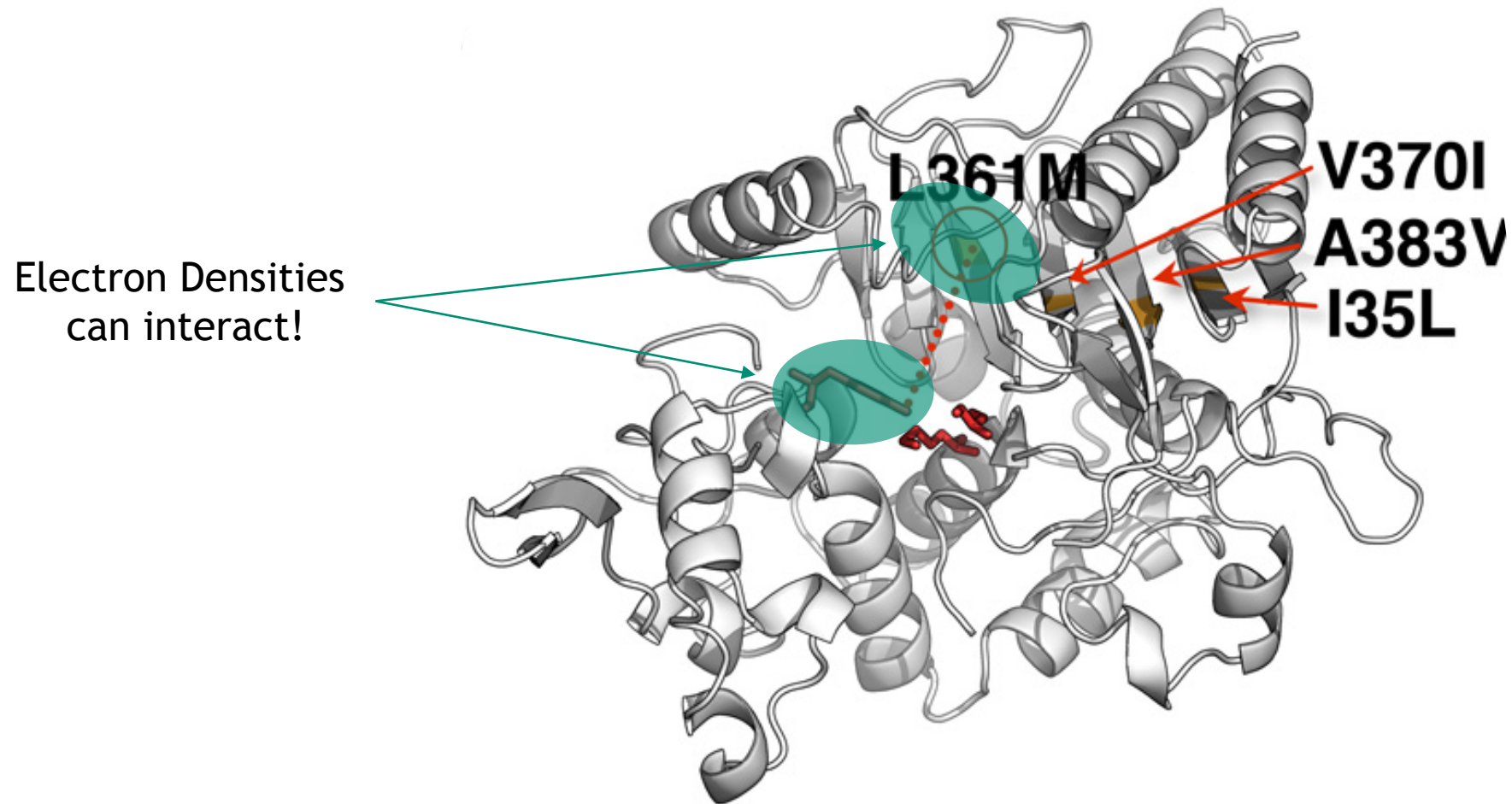
Oliver T. Unke,[△] Stefan Chmiela,[△] Huziel E. Sauceda, Michael Gastegger, Igor Poltavsky, Kristof T. Schütt, Alexandre Tkatchenko,^{*} and Klaus-Robert Müller^{*}





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Solution: Learn the **electron density**

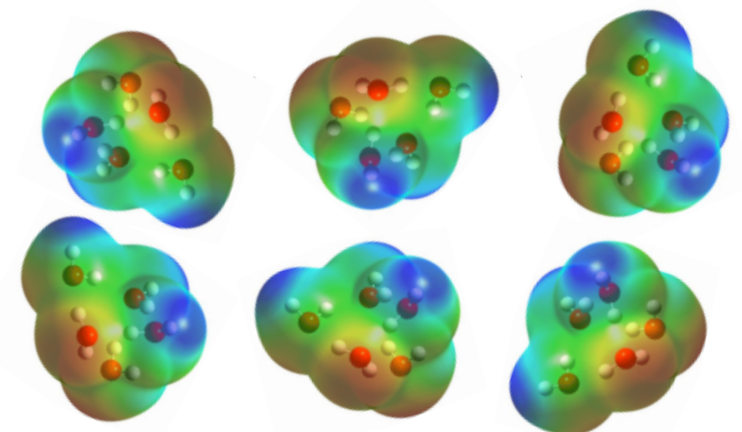
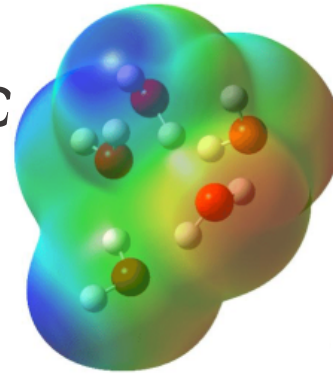




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What's the Problem? – Part 2: Learning in 3D Space

The electron density is a 3D object



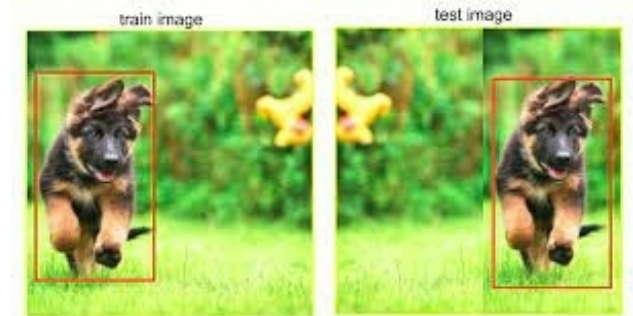
Convolution
changed
everything.



1 x1	1 x0	1 x1	0	0
0 x0	1 x1	1 x0	1	0
0 x1	0 x0	1 x1	1	1
0	0	1	1	0
0	1	1	0	0

Image

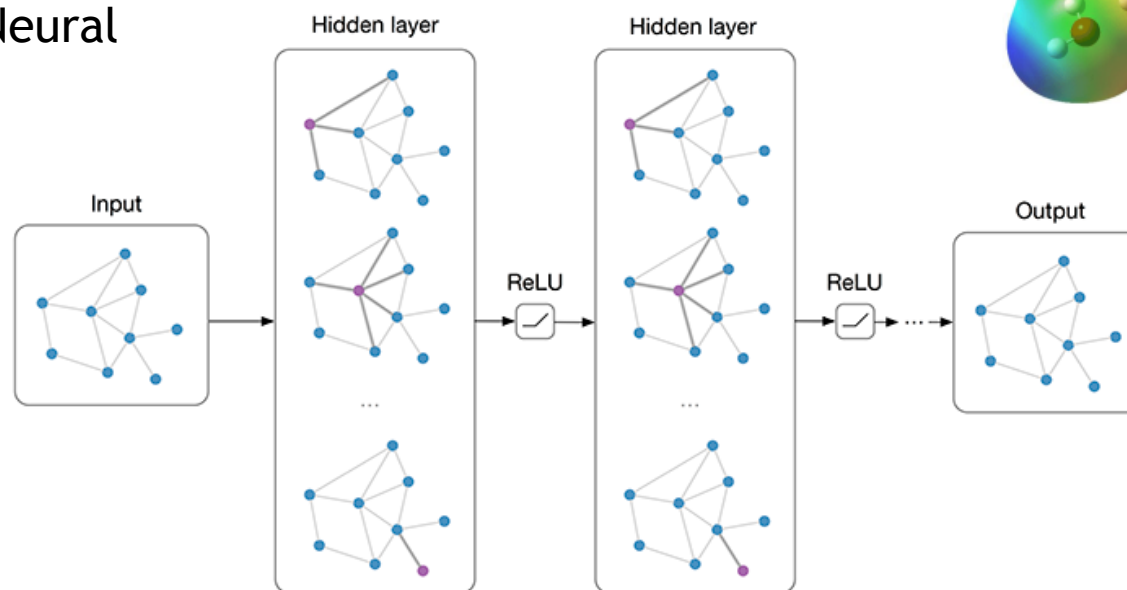
4		

Convolved
Feature

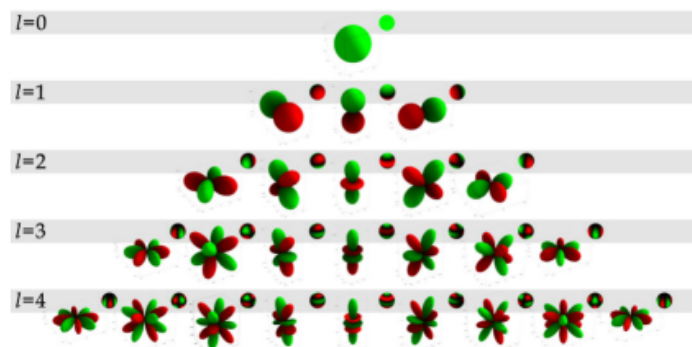
11

Solution: Euclidean Neural Networks

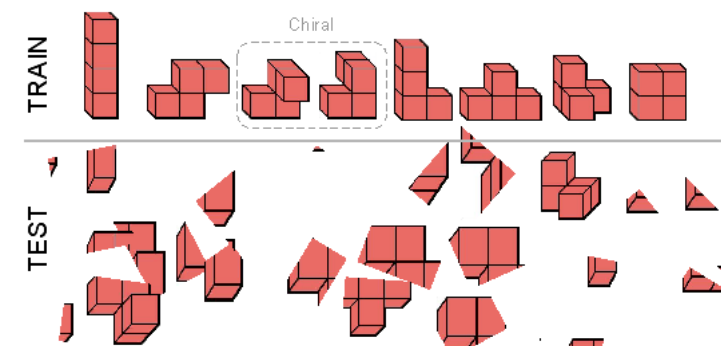
Graph Convolutional Neural Network:



e3nn features are combinations of spherical harmonics



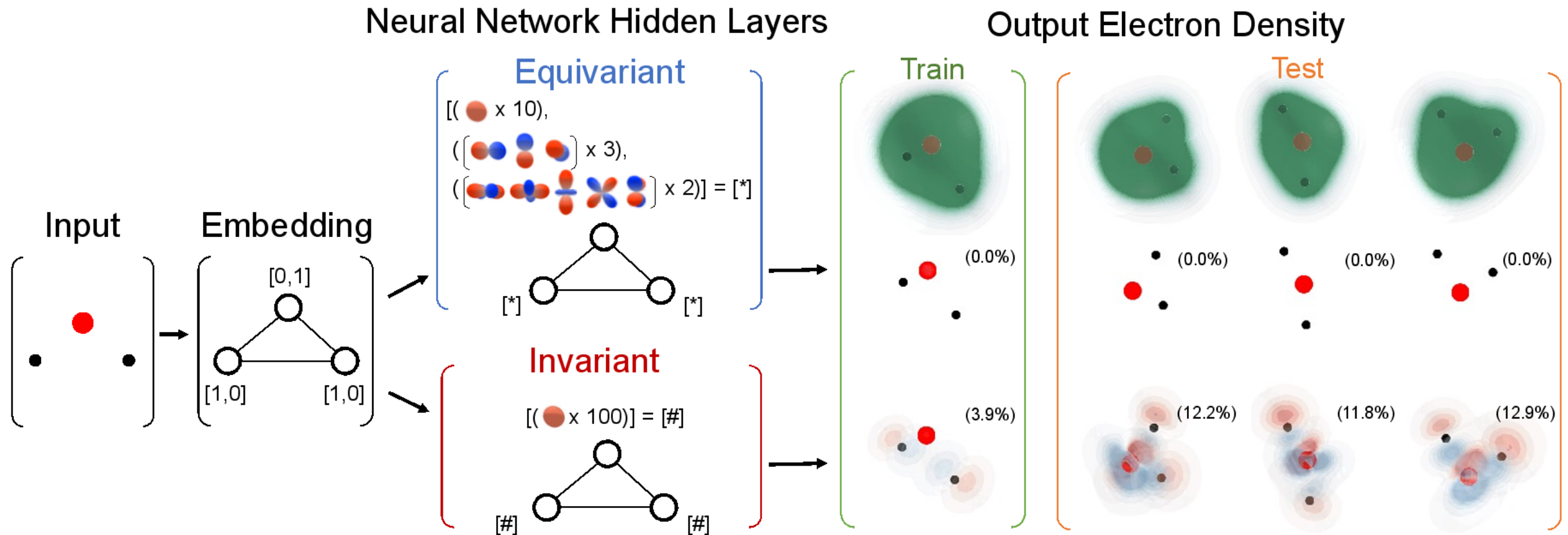
e3nn is equivariant





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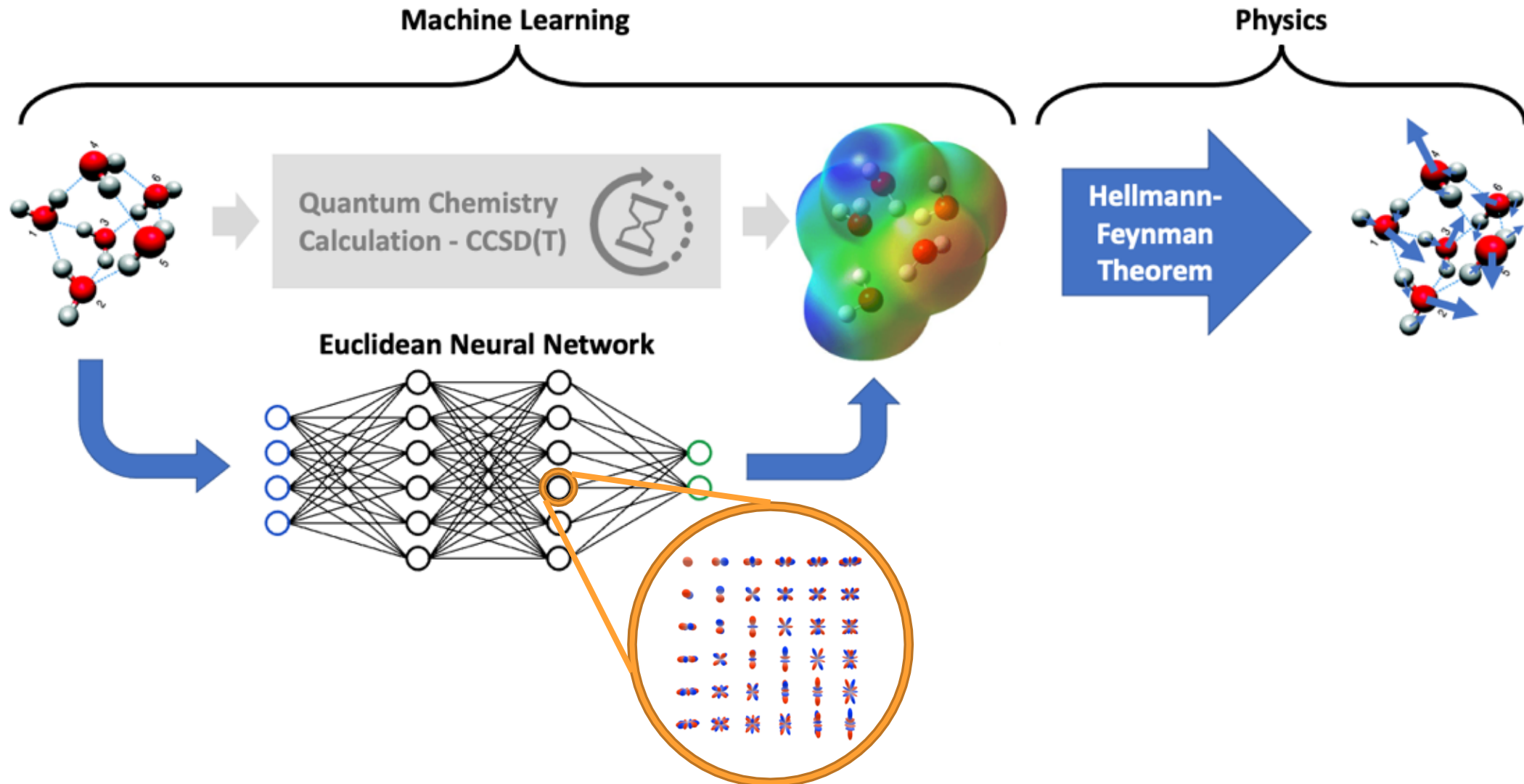
Solution: Euclidean Neural Networks





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The Big Idea: Learn the electron density with Euclidean Neural Networks





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Experiment 1: Learning the e^- density

Train on model on 500 samples of 8 water molecule clusters (DFT) (For reference, ImageNet has 14 million images.)

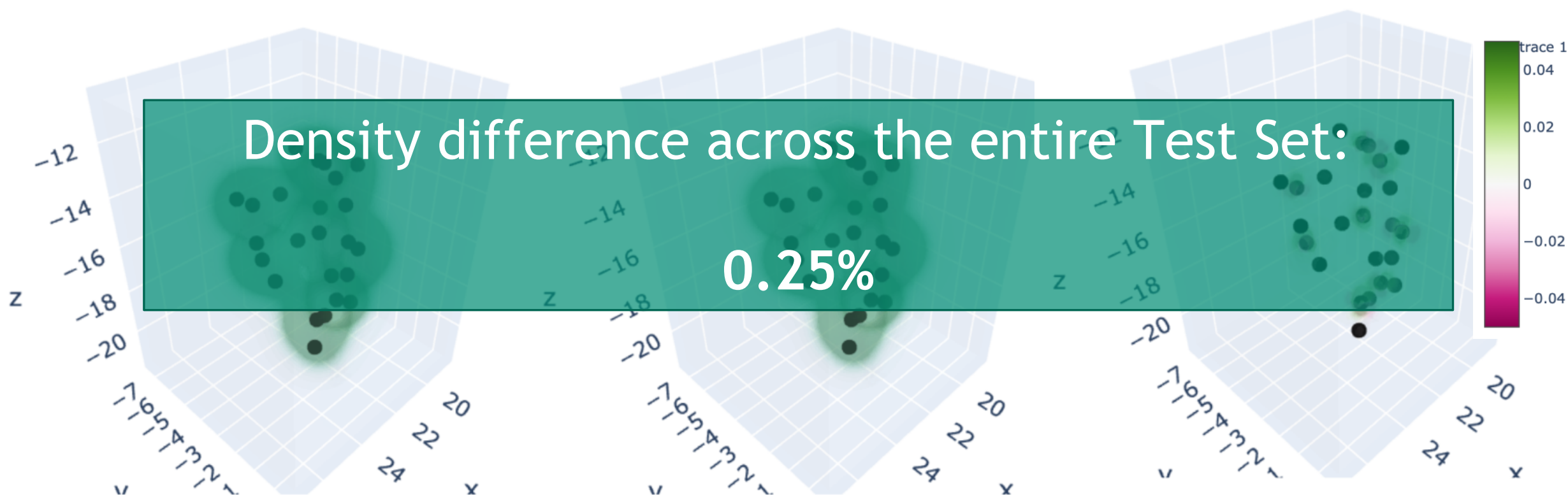
ML Density

Target Density

Difference

Density difference across the entire Test Set:

0.25%

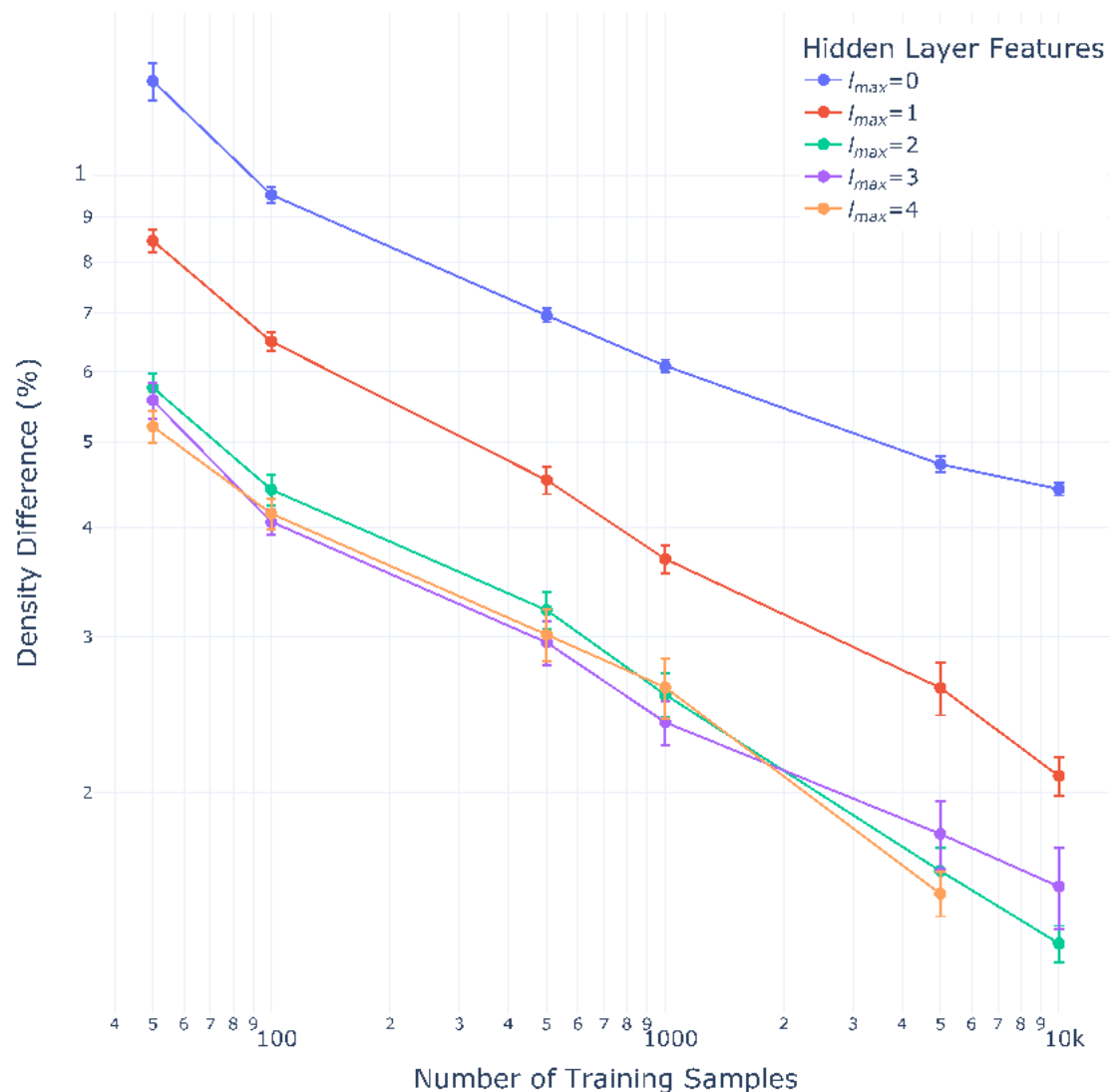




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An aside: The importance of equivariance

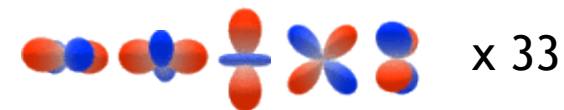
Electron Density Learning Curves



$l_{max}=0$: x 500

$l_{max}=1$: x 250 + x 83

$l_{max}=2$: x 167 + x 56 +



⋮

(Total number of 500 features in every model)

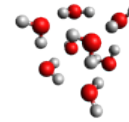


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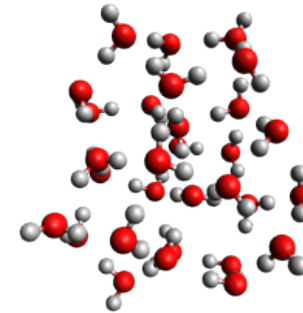
Experiment 2: Extrapolation

Train on 8 water clusters. Test on 30 water clusters.

Train



Test



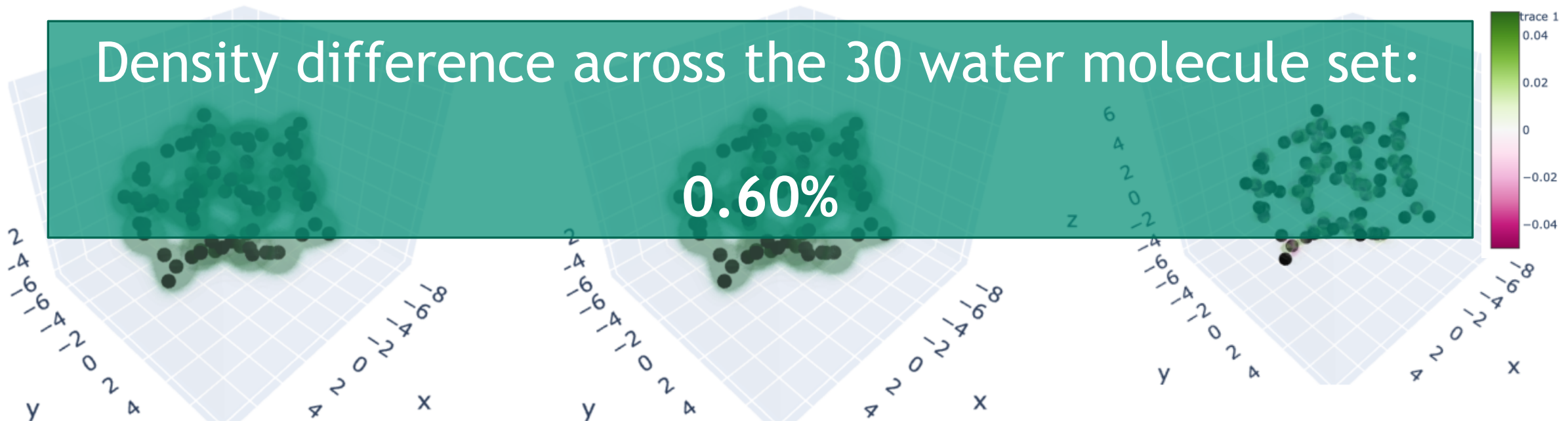
ML Density

Target Density

Difference

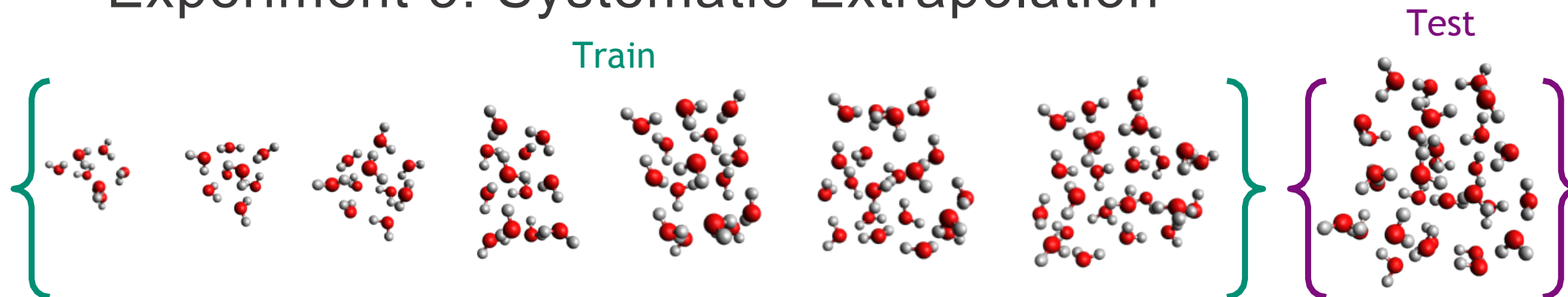
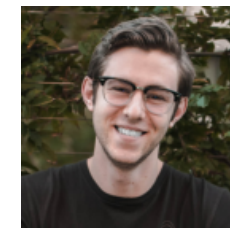
Density difference across the 30 water molecule set:

0.60%

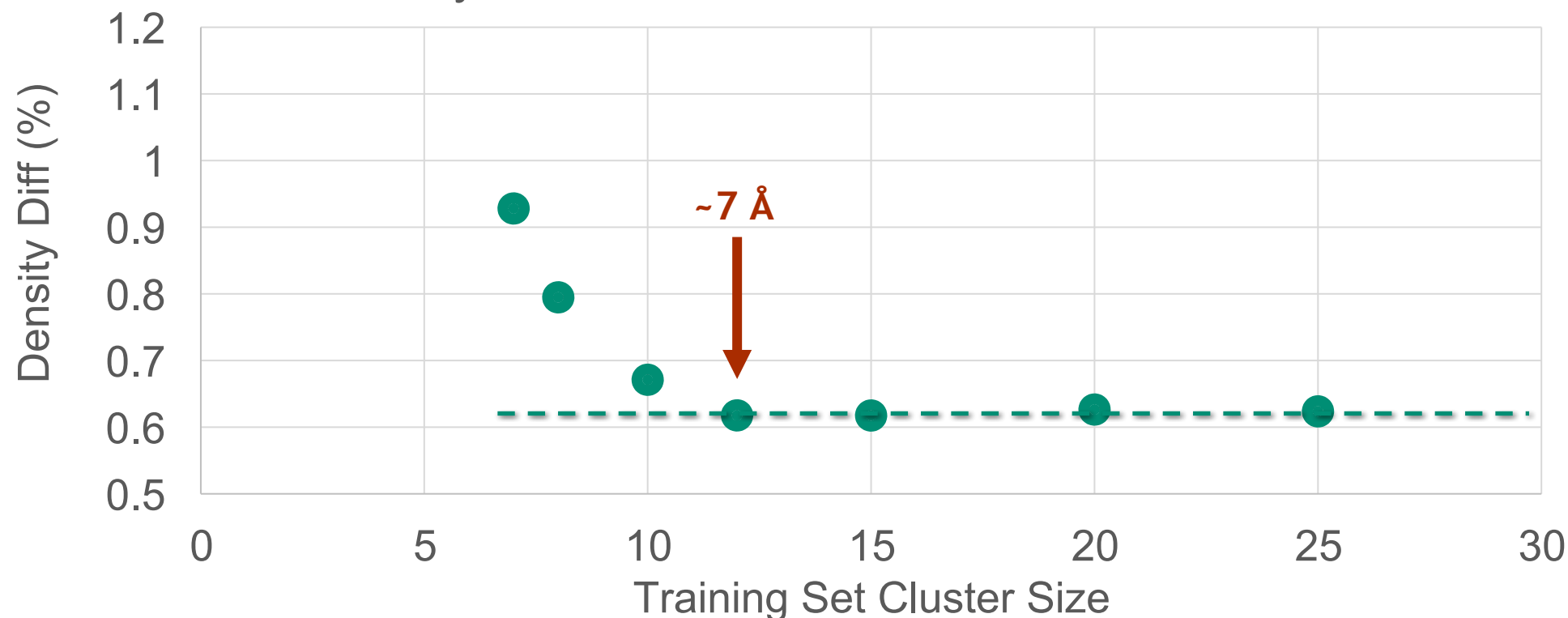


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Experiment 3: Systematic Extrapolation



Density Difference: 30 water molecule test set



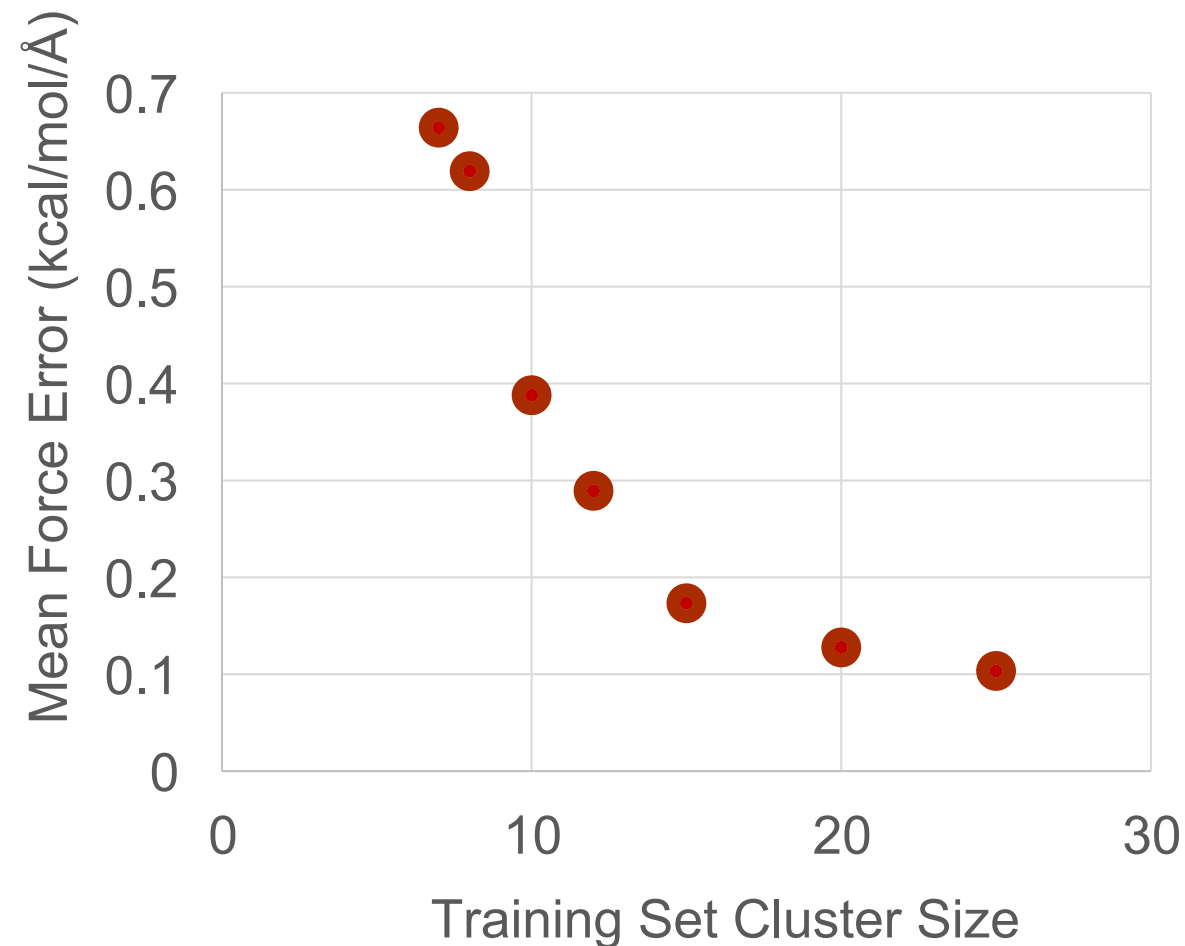


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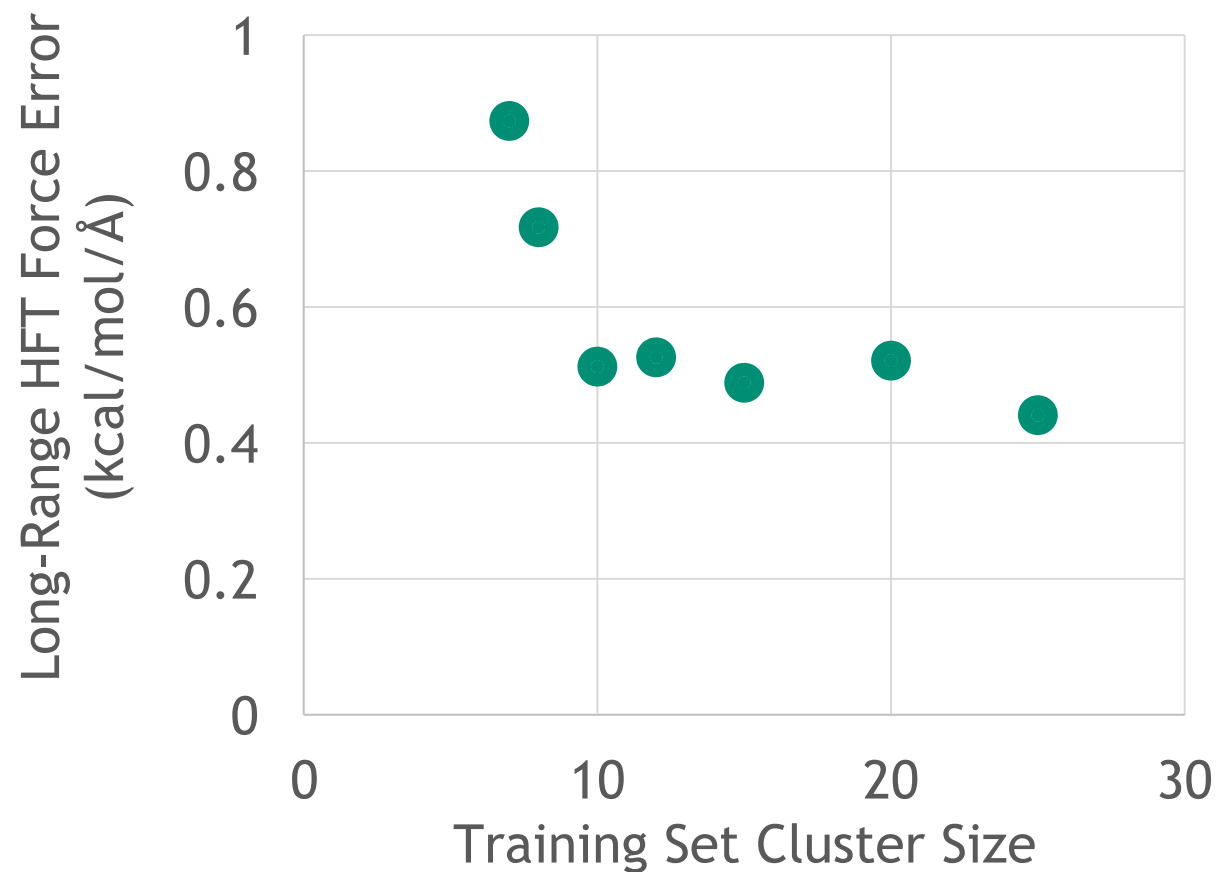
Experiment 4: Comparison with forces

(ML models from previous slide)

Force Difference: 30 waters



Long-Range Hellmann-Feynman
Force Difference: 30 waters





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Long-range interactions affect **forces** more than **density**

Energy and Force

Electron Density

13 Å+

~7 Å

By learning the electron density, we can
extrapolate much more effectively



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Long-range interactions affect **forces** more than **density**

This is consistent with our understanding of **perturbation theory**.

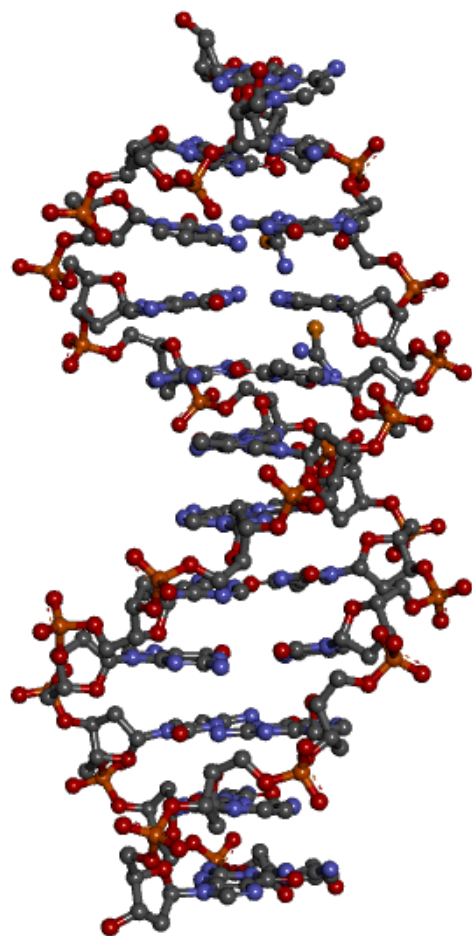


$$E_{PT} = E^{(0)} + \underbrace{E_{elst}^{(1)}}_{\text{Electrostatics}} + \underbrace{E_{exch}^{(1)} + E^{(2)} + \dots}_{\text{Polarization Exchange-Repulsion Dispersion}}$$

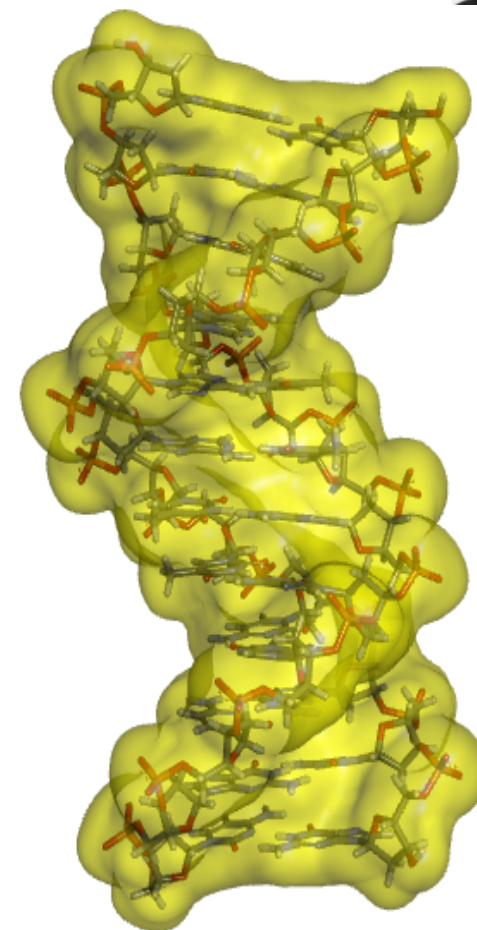
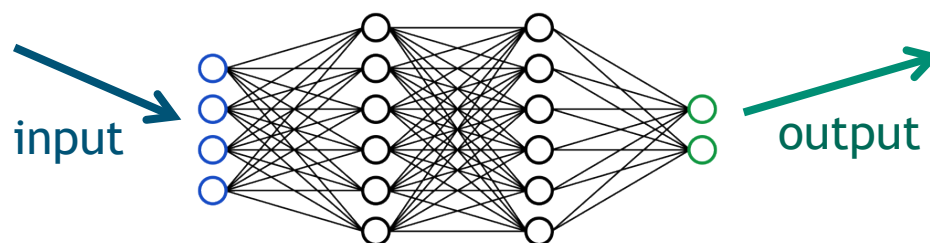
Electrostatics

Polarization
Exchange-Repulsion
Dispersion

DNA – ML electron densities at the biomolecular scale



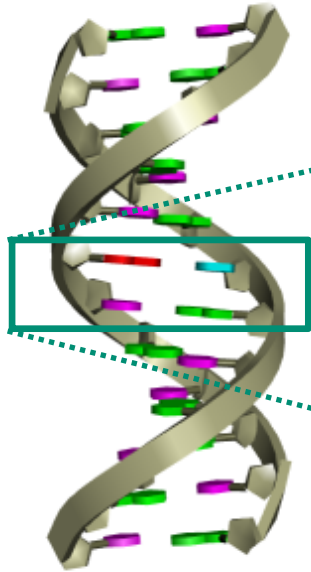
$$\hat{H}\psi = E\psi$$



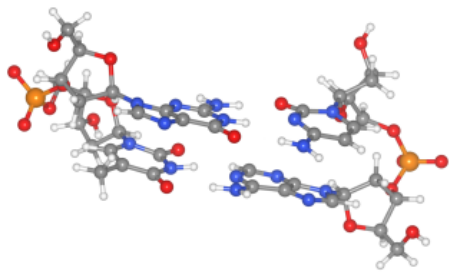
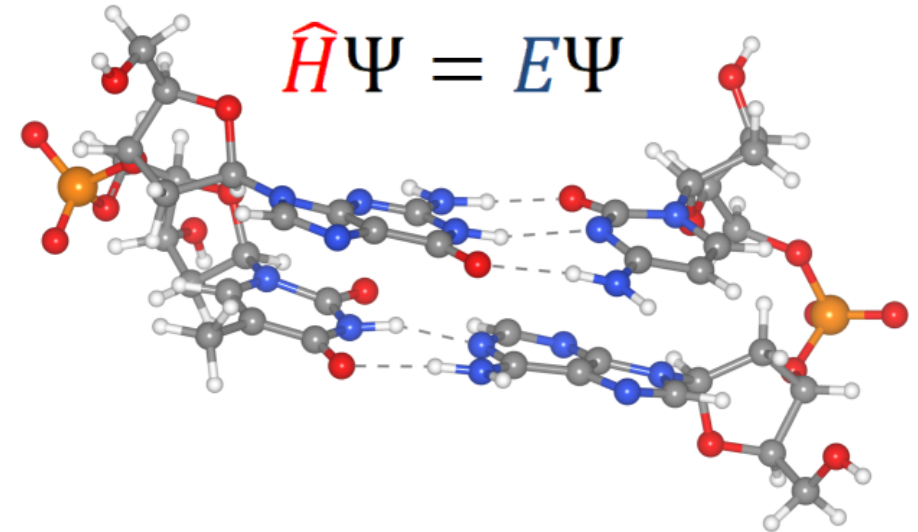
DNA – Learning the electron density



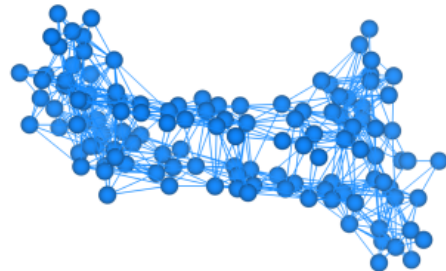
Molecular dynamics



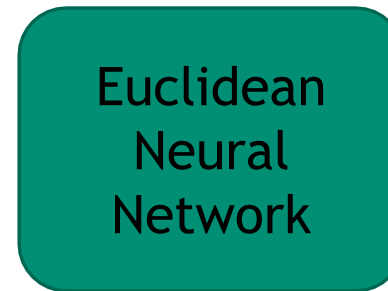
Quantum



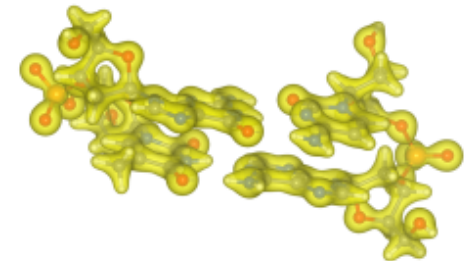
Atomic coordinates



Input layer



Hidden layers



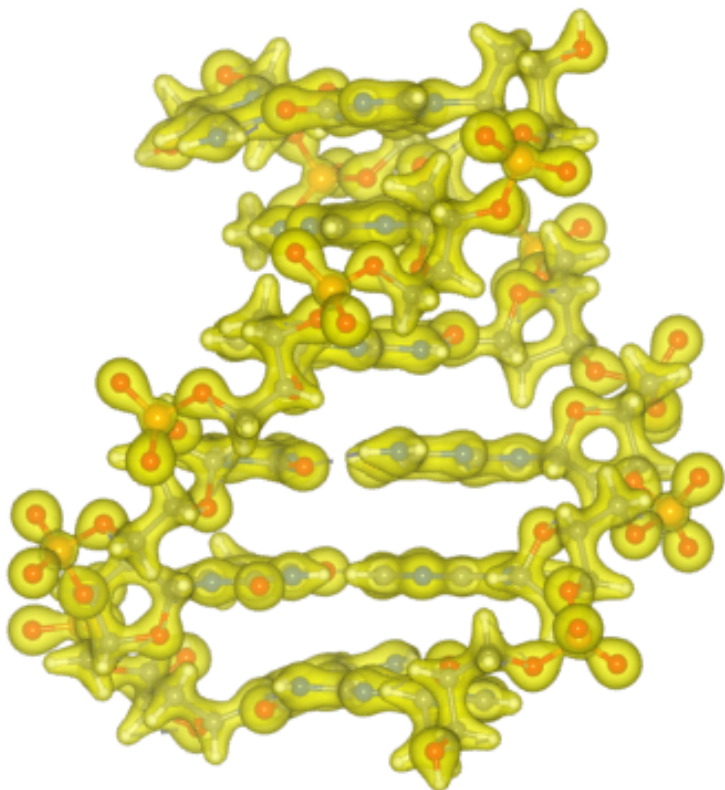
Output layer



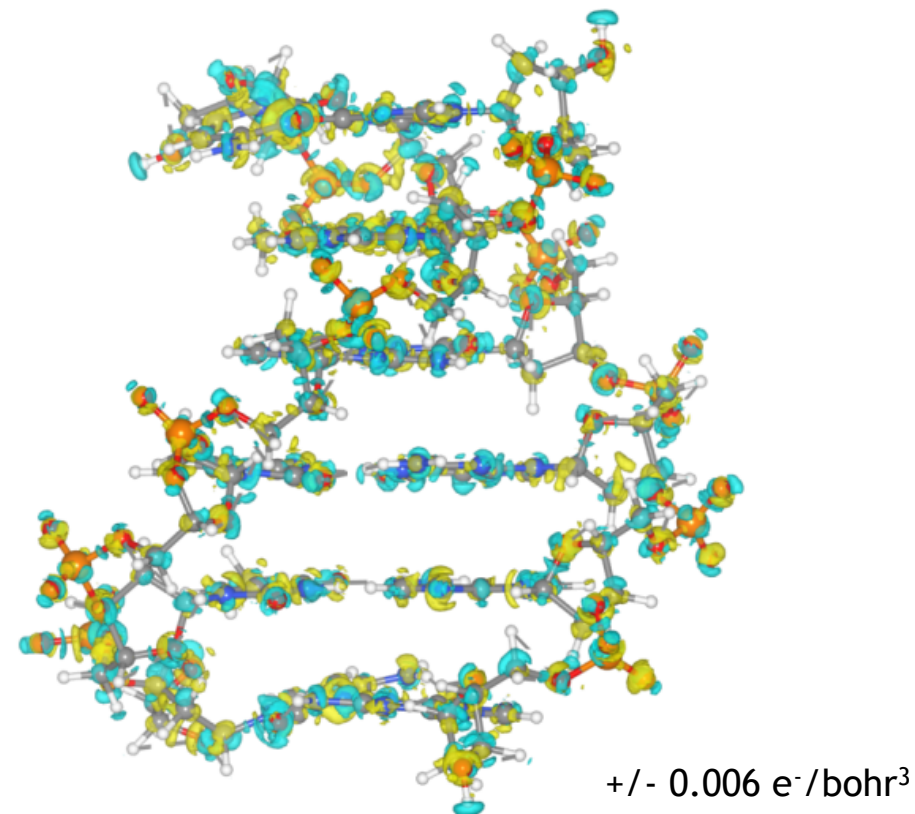
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A real, experimental structure! (PDB code 251d)

ML prediction



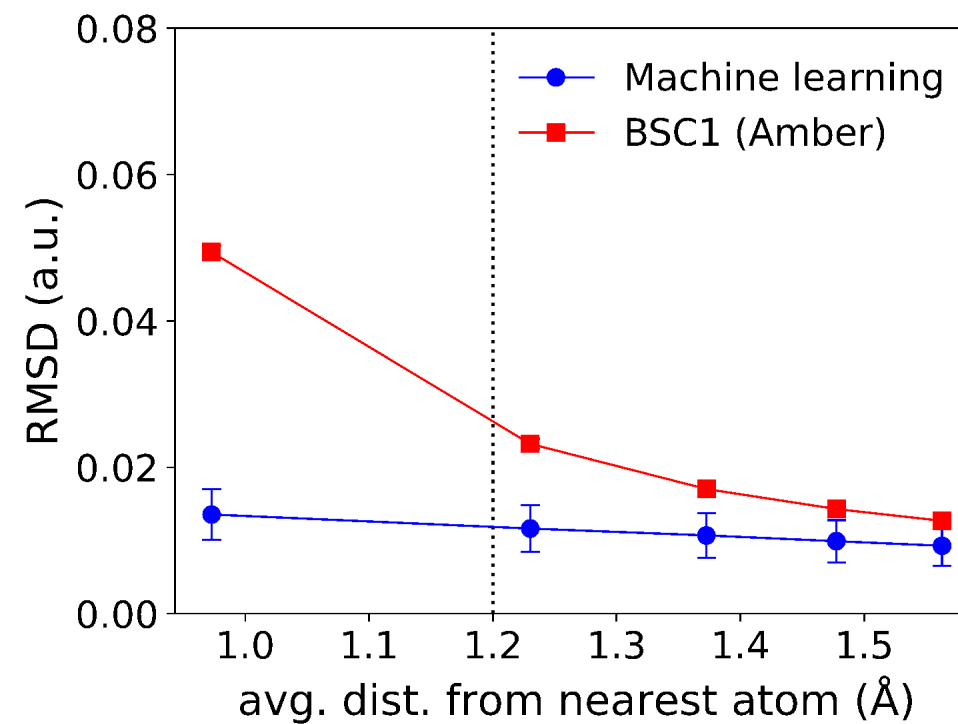
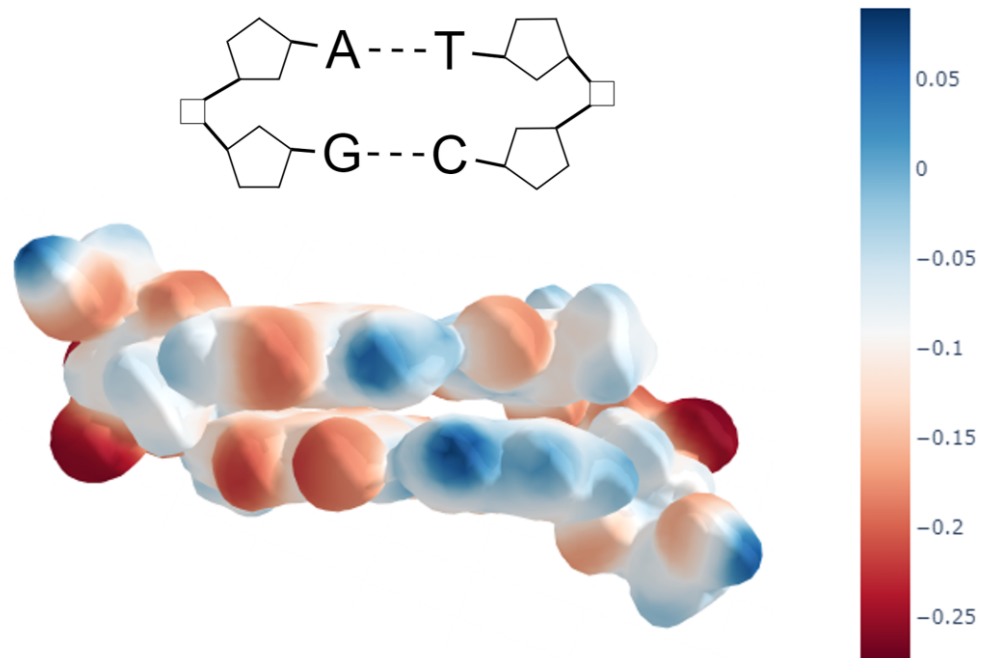
Density difference: ML - target





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Electrostatic Potential

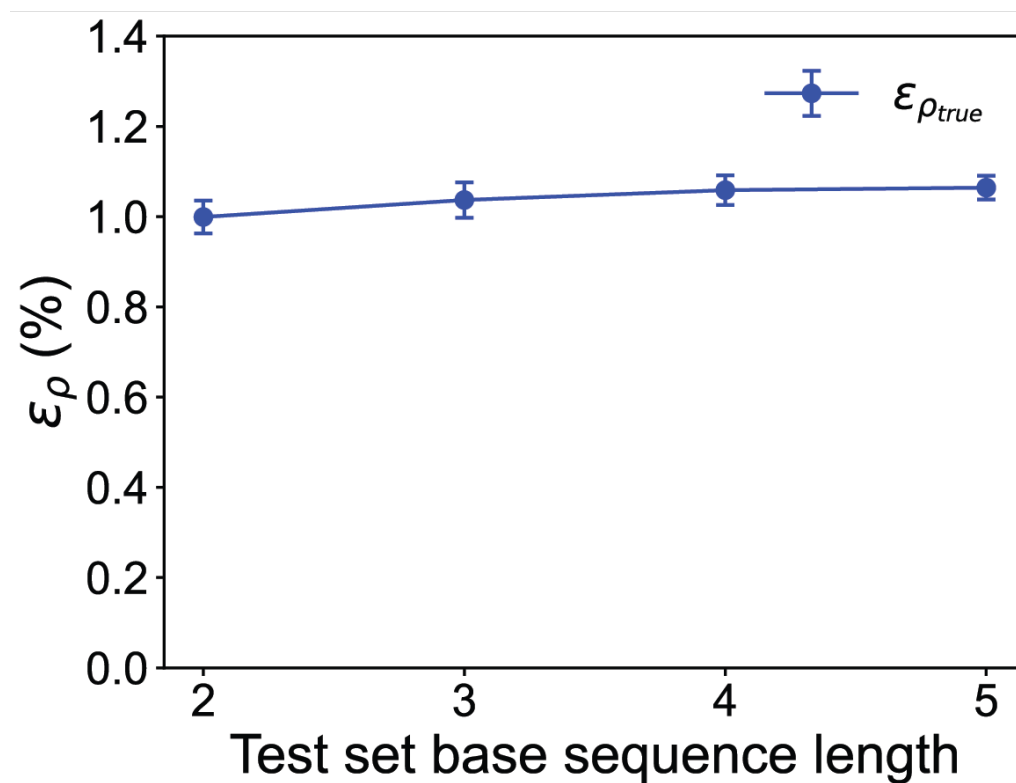




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Systematic Extrapolation

Sequence Length Size Dependence

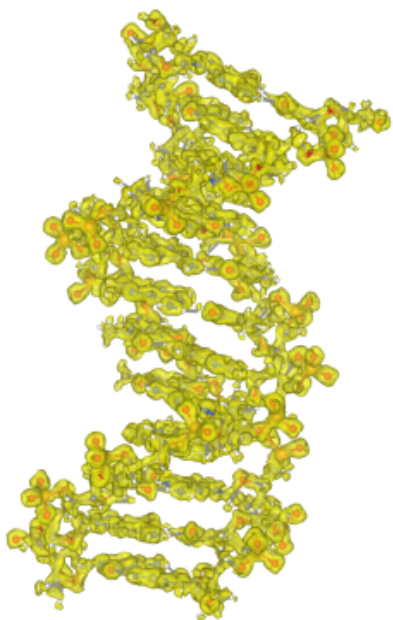




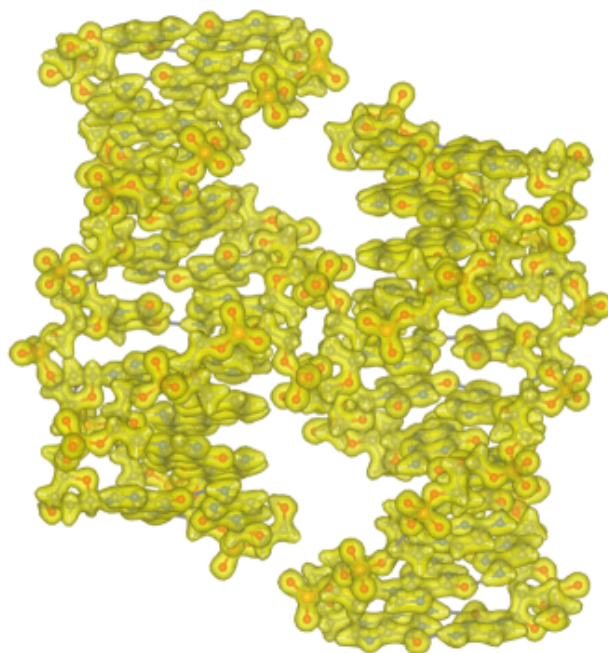
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We can predict very large structures!

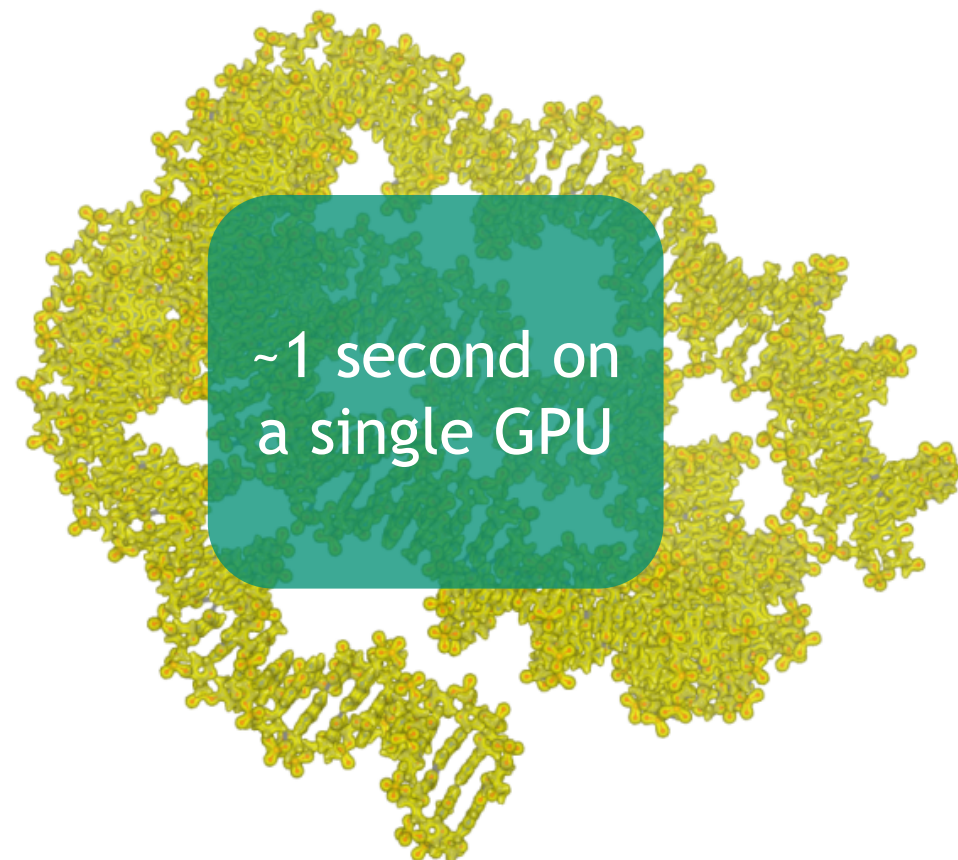
Drew-Dickerson dodecamer
758 atoms, 3780 electrons
(PDB code 4c64)



Stacked Holliday junction
1260 atoms, 6280 electrons
(PDB code 1dcw)



Nucleosome core particle
147 bps, 9346 atoms, 46980 electrons
(PDB code 1kx5)

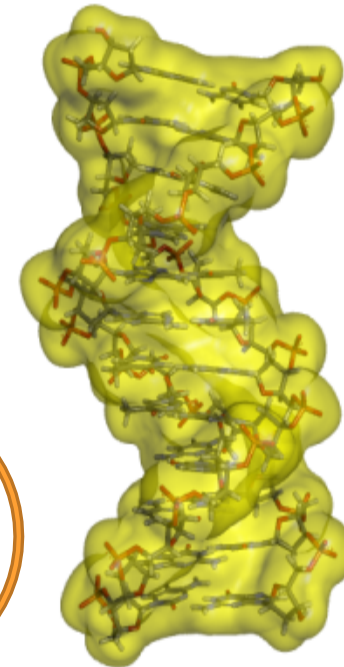
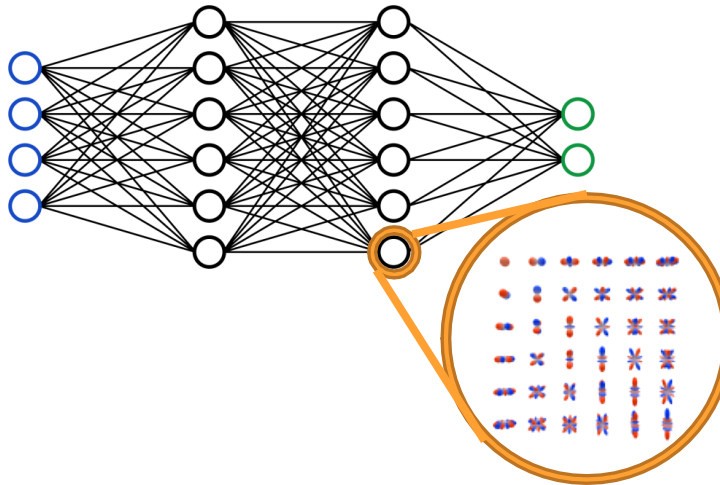
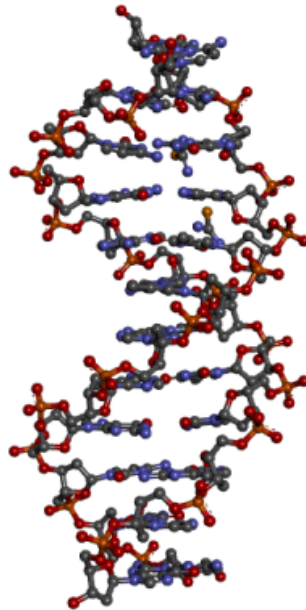


Take-Aways

A. Any serious model has to be able to "train on small, run on big".

B. Euclidean Neural Networks + physics is the tool for the job

C. This works for water and DNA!

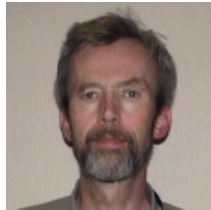




Acknowledgements



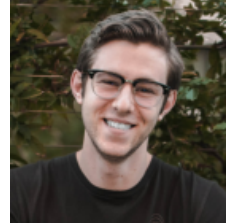
Steve Plimpton



Aidan Thompson



Susan Rempe



Lucas Tecot
(UCLA)



Tess Smidt
(MIT)



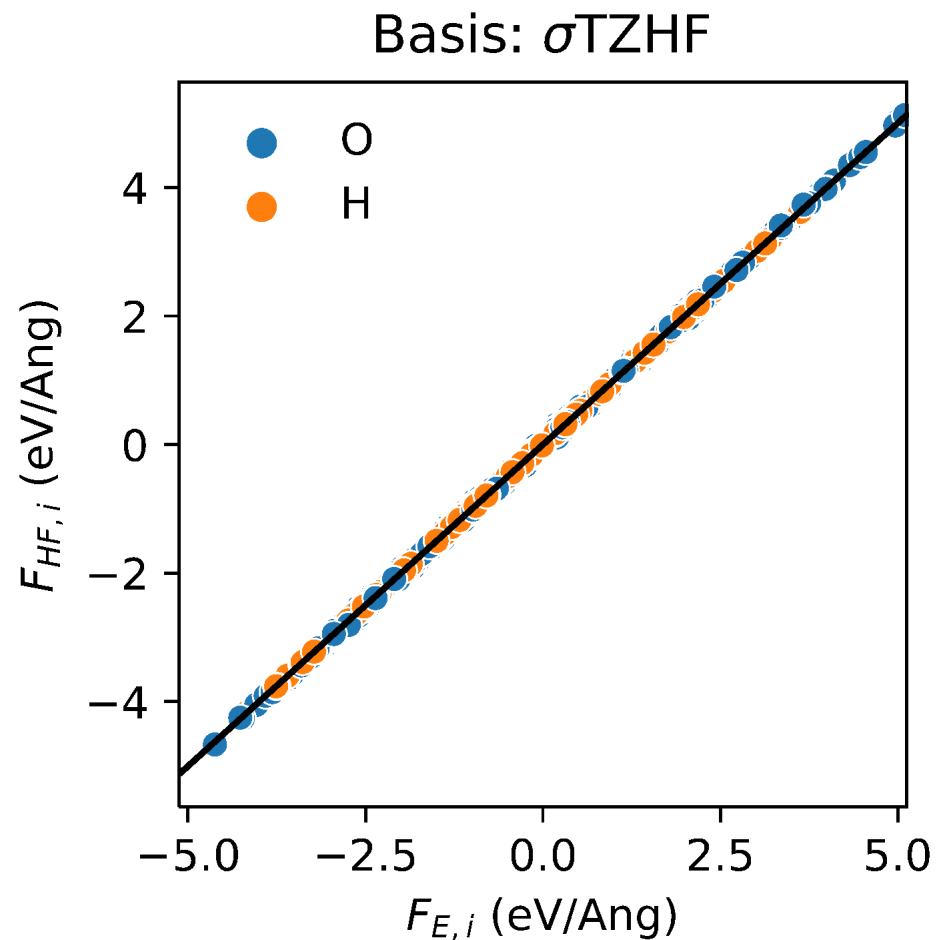
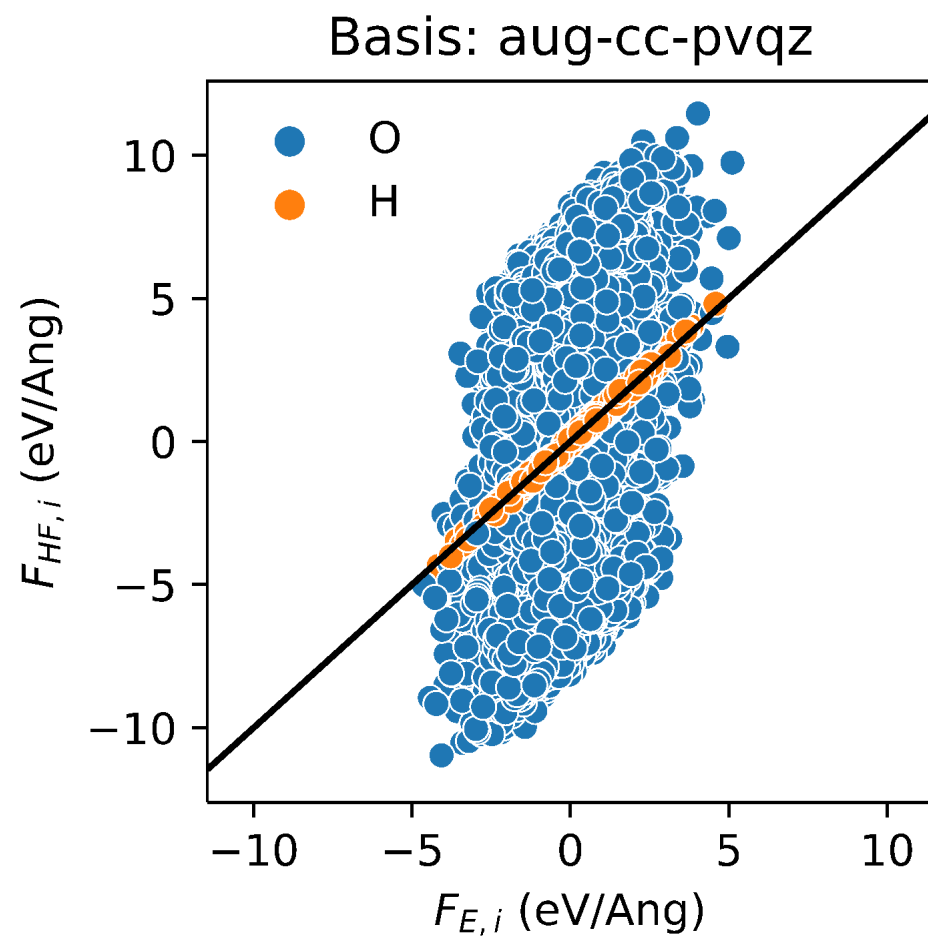
Shivesh Pathak
(Sandia)



Will Bricker and Alex
Lee
(UNM)



Hellmann-Feynman Forces



Solution: Learn the **electron density**

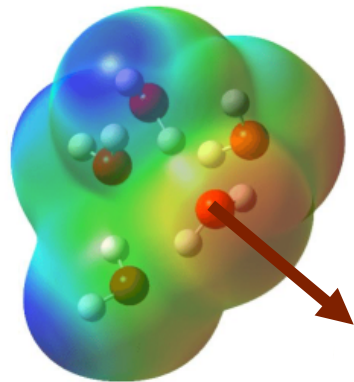
The Hellmann-Feynman Theorem:

The force on any atom can be exactly calculated from the surrounding **electron density** **using only classical electrostatics**.

Machine Learning

$$H\Psi = E\Psi$$

$$\rho = \Psi^*\Psi$$

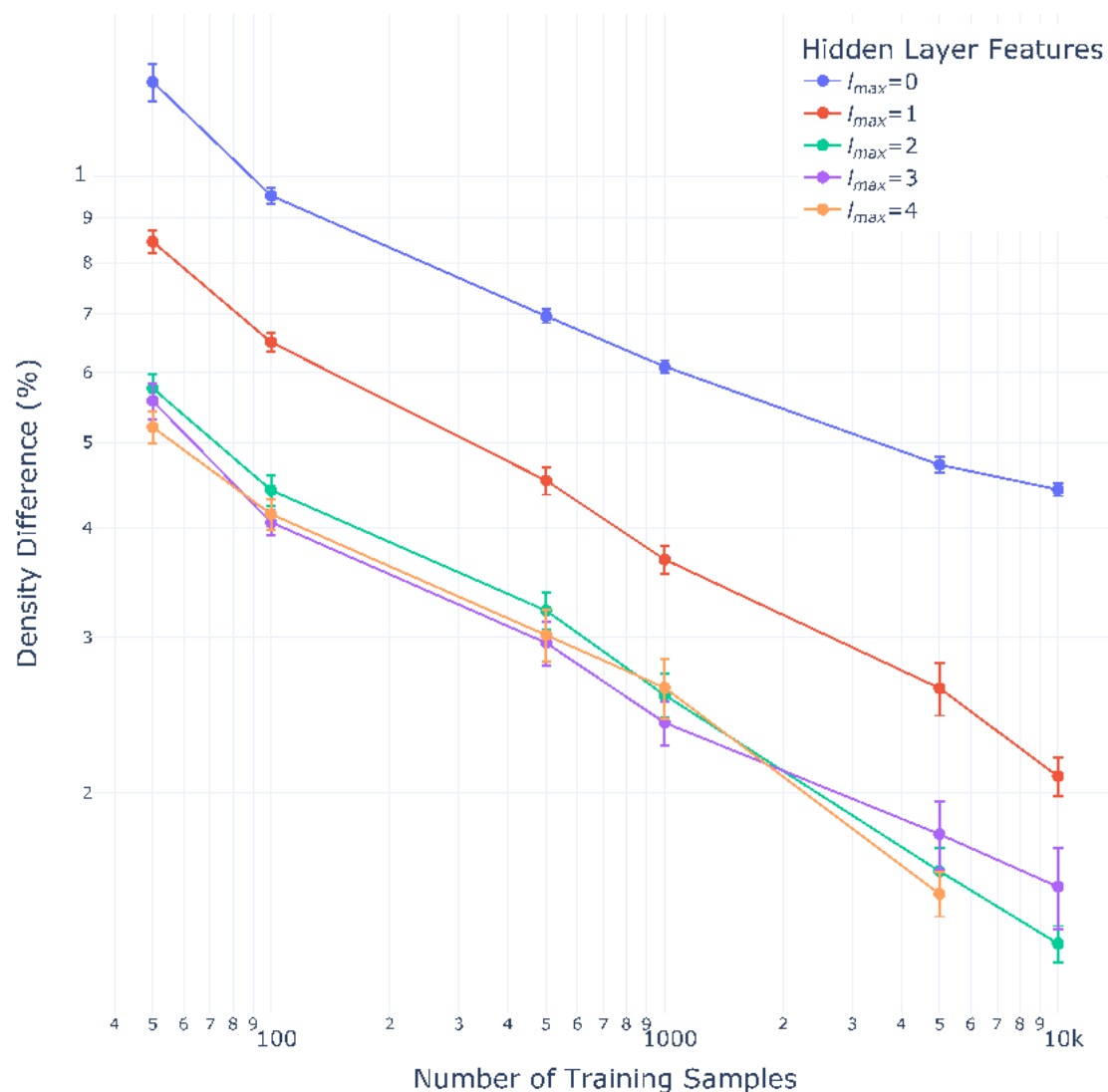


Physics

$$\vec{F}_i = \sum_j^{N_{\text{nuclei}}} \frac{Z_i Z_j}{R_{ij}^3} - Z_i \int \frac{\rho}{r^3} dv$$

Experiment 2: The importance of higher-order features

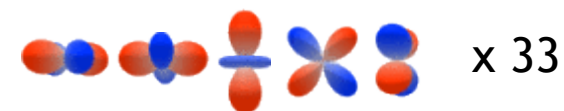
Electron Density Learning Curves



$l_{max}=0$: x 500

$l_{max}=1$: x 250 + x 83

$l_{max}=2$: x 167 + x 56 +



⋮

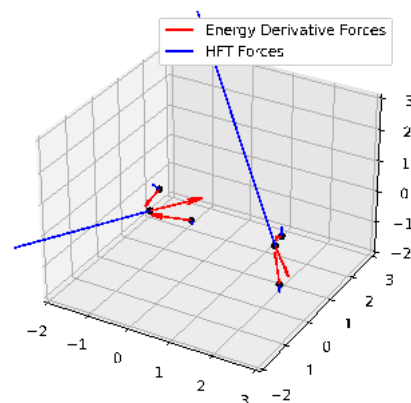
(Total number of 500 features in every model)

Hellmann-Feynman Forces



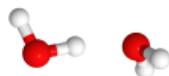
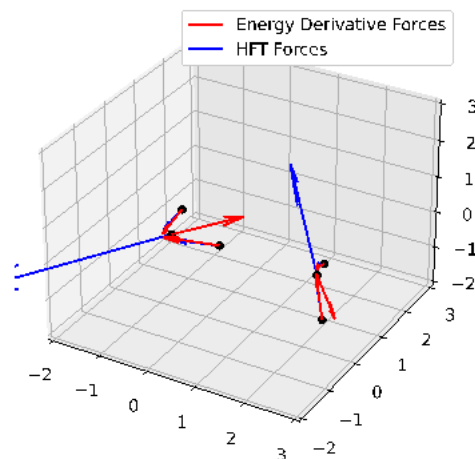
aug-cc-pvdz

82 basis functions



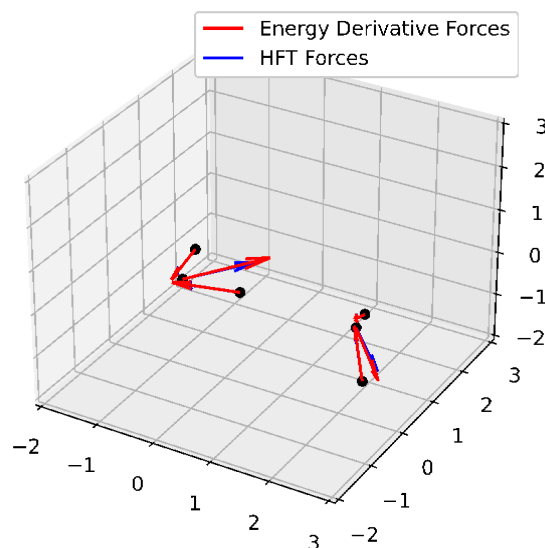
aug-cc-pvqz

344 basis functions



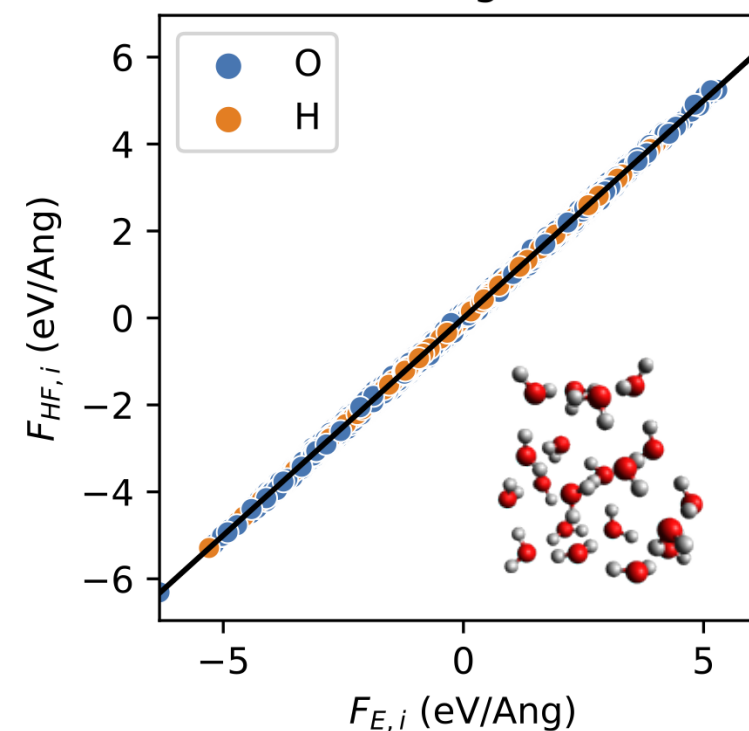
Hellmann-Feynman Basis

154 basis functions

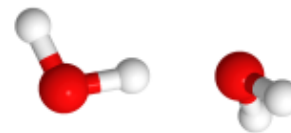


**Hellmann-Feynman Force Error:
20 water molecule clusters**

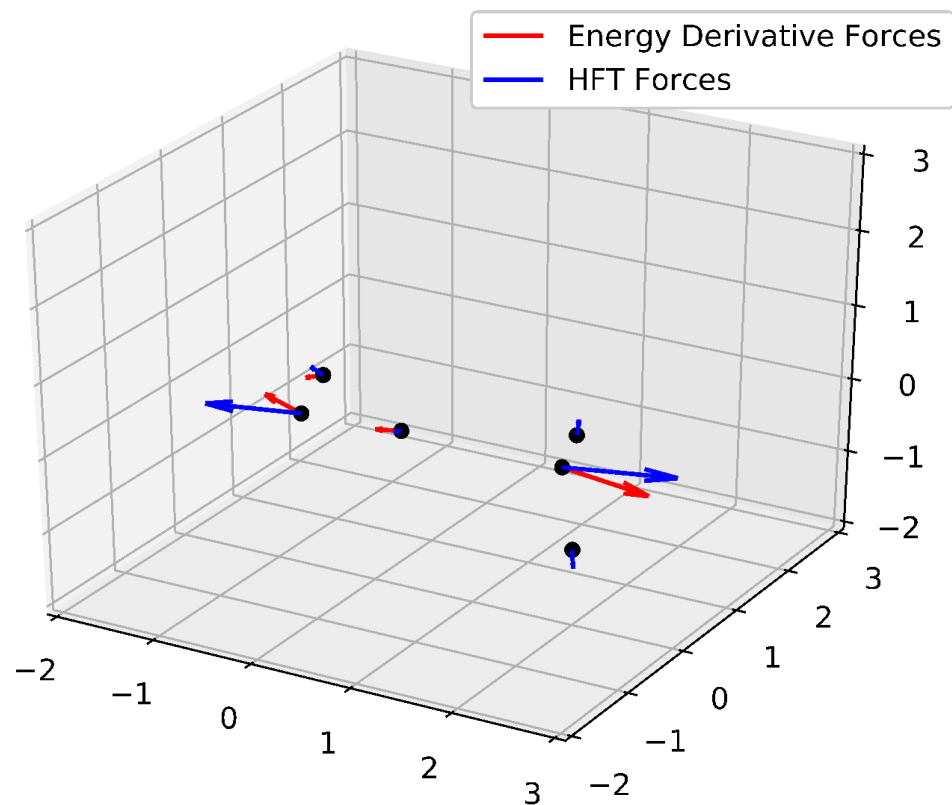
Basis: large-basis



HFT forces with CCSD(T) densities



hft-basis-MEDIUM
154 basis functions



hft-basis-LARGE
334 basis functions

