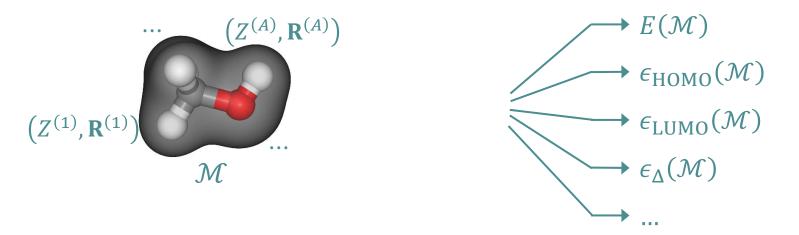
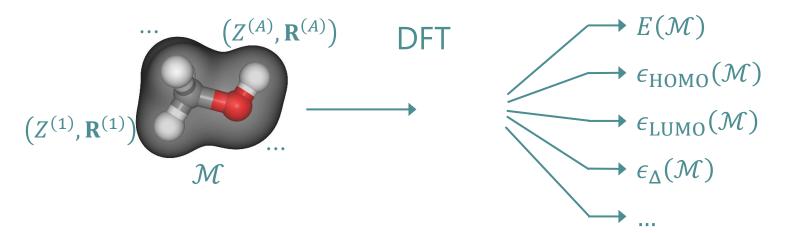


# Self-Consistency Training for Density-Functional-Theory Hamiltonian Prediction

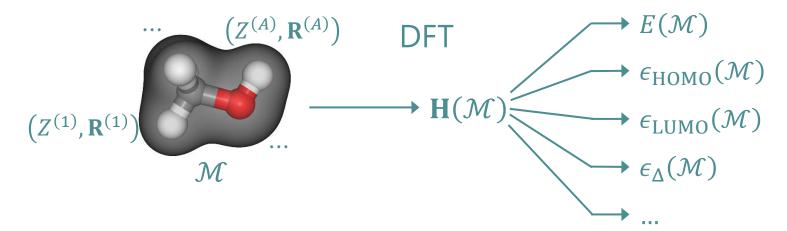
Chang Liu on behalf of the team changliu@microsoft.com



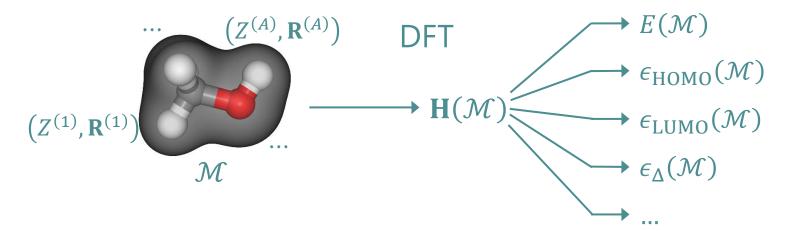
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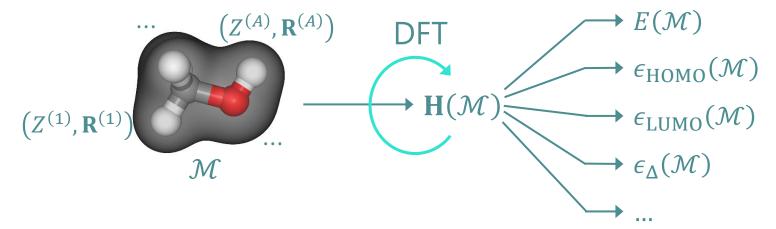


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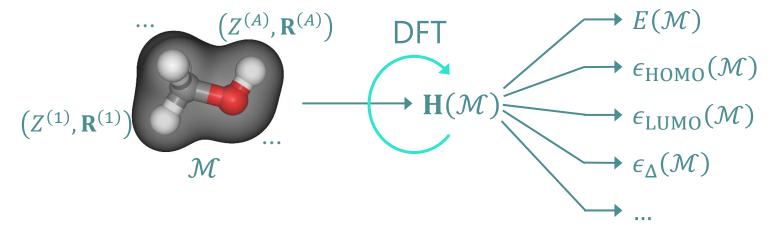


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- Hamiltonian prediction:

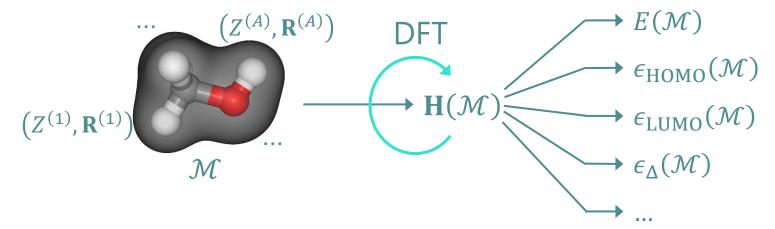
"root property" prediction, provide all properties that DFT can.



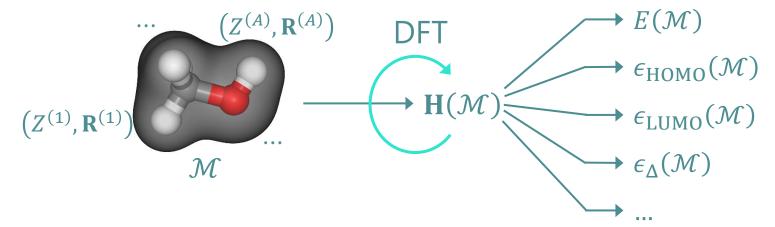
- · Hamiltonian prediction has a self-consistency principle: Training without label!
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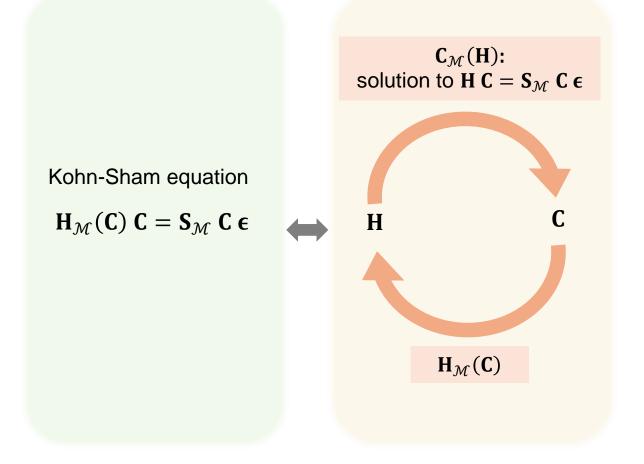
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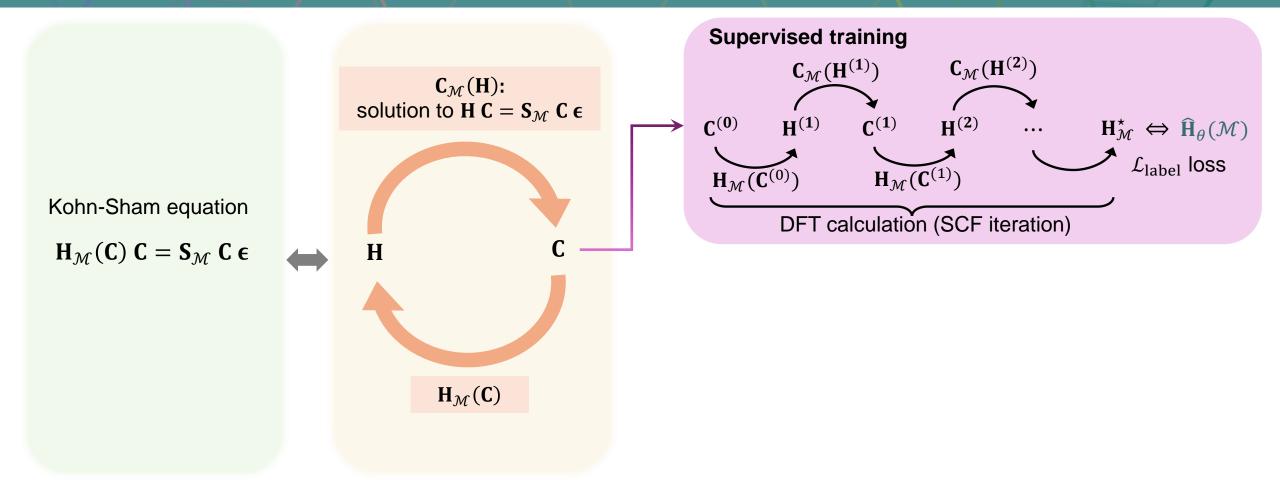
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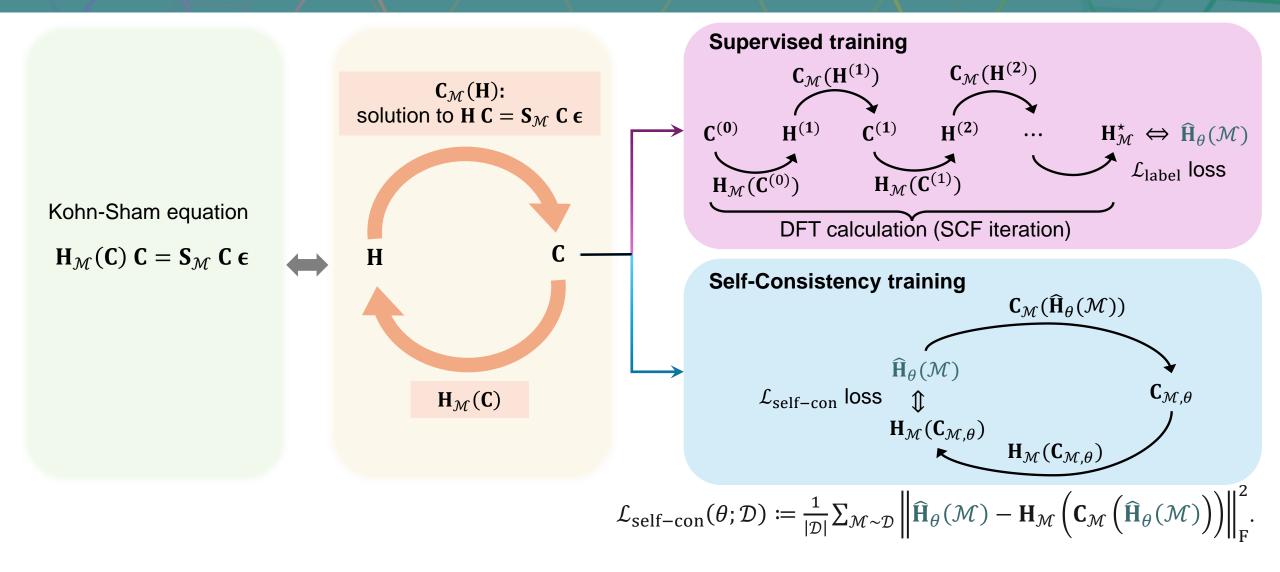
$$\underbrace{H_{\mathcal{M}}(\mathbf{C})}_{:=\nabla E_{\mathcal{M}}(\cdot)|_{\mathbf{C}\mathbf{C}^{\mathsf{T}}}} \mathbf{C} = \mathbf{S}_{\mathcal{M}} \mathbf{C} \boldsymbol{\epsilon}.$$
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# Self-Consistency Training

• Self-consistency loss:

$$\mathcal{L}_{\text{self-con}}(\theta; \mathcal{D}) \coloneqq \frac{1}{|\mathcal{D}|} \sum_{\mathcal{M} \sim \mathcal{D}} \left\| \widehat{\mathbf{H}}_{\theta}(\mathcal{M}) - \mathbf{H}_{\mathcal{M}}\left( \mathbf{C}_{\mathcal{M}}\left( \widehat{\mathbf{H}}_{\theta}(\mathcal{M}) \right) \right) \right\|_{F}^{2}.$$

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- · Numerically stable implementation of differentiation through eigensolver.
- GPU implementation of Hamiltonian construction  $H_{\mathcal{M}}(\mathbf{C})$ .

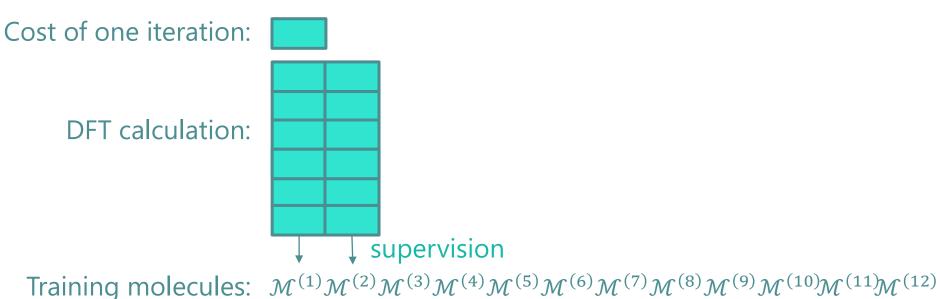
1. Generalization beyond labeled data:  $\mathcal{L}_{label}(\theta; \overline{\mathcal{D}^{(1)}}) + \lambda \mathcal{L}_{self-con}(\theta; \underline{\mathcal{D}^{(2)}})$ . limited labeled dataset unlimited unlabeled dataset

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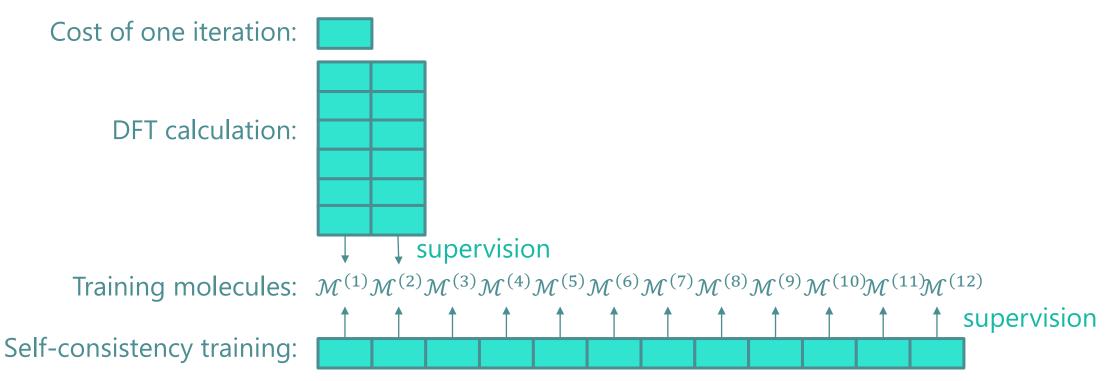
Cost of one iteration:

Training molecules:  $\mathcal{M}^{(1)}\mathcal{M}^{(2)}\mathcal{M}^{(3)}\mathcal{M}^{(4)}\mathcal{M}^{(5)}\mathcal{M}^{(6)}\mathcal{M}^{(7)}\mathcal{M}^{(8)}\mathcal{M}^{(9)}\mathcal{M}^{(10)}\mathcal{M}^{(11)}\mathcal{M}^{(12)}$ 

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# Generalization beyond Labeled Data

· Data-scarce scenario (MD17): 100 labeled + 24900 unlabeled  $\rightarrow$  test.

	Direct prediction			Derived molecular properties				As DFT init.
			[		l			
Molecule	Setting	$\mathbf{H}\left[\mu E_{\mathbf{h}}\right]\downarrow$	$\epsilon \left[\mu E_{\rm h}\right] \downarrow$	$\mathbf{C}\left[\% ight]\uparrow$	$\epsilon_{\mathrm{HOMO}} \left[ \mu E_{\mathrm{h}} \right] \downarrow$	$\epsilon_{ m LUMO} \left[ \mu E_{ m h} \right] \downarrow$	$\epsilon_{\Delta} \left[ \mu E_{\rm h} \right] \downarrow$	SCF Accel. [%] $\downarrow$
Ethanol	label	160.36	712.54	99.44	911.64	6800.84	6643.11	68.3
	label + self-con	<b>75.65</b>	<b>285.49</b>	<b>99.94</b>	<b>336.97</b>	<b>1203.60</b>	<b>1224.86</b>	<b>61.5</b>
Malondi-	label	101.19	456.75	99.09	471.92	1093.22	1115.94	69.1
aldehyde	label + self-con	<b>86.60</b>	<b>280.39</b>	<b>99.67</b>	<b>274.45</b>	279.14	<b>324.37</b>	<b>62.1</b>
Uracil	label	88.26	1079.51	95.83	1217.17	12496.1	11850.56	65.8
	label + self-con	<b>63.82</b>	<b>315.40</b>	<b>99.58</b>	<b>359.98</b>	<b>369.67</b>	<b>388.30</b>	<b>54.5</b>

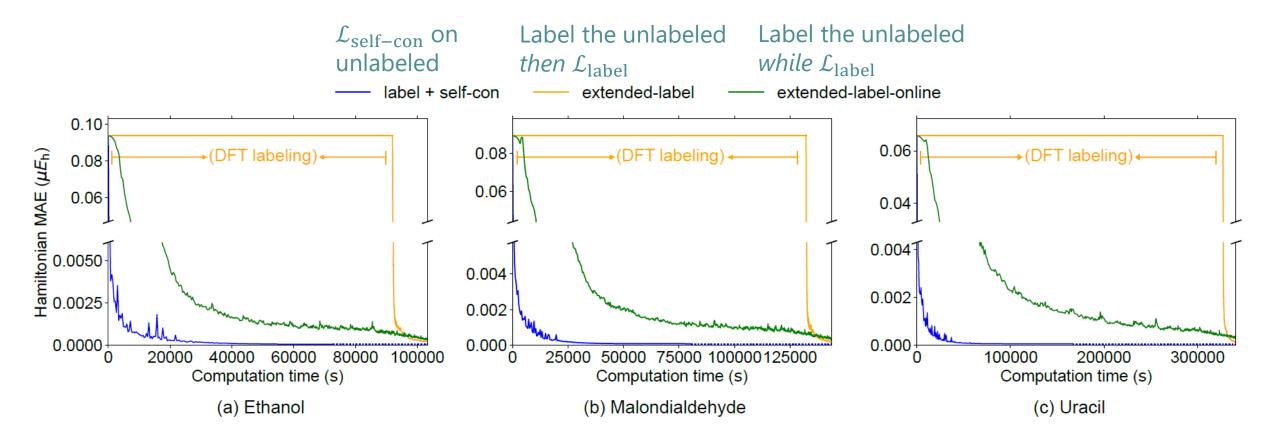
# Generalization beyond Labeled Data

#### • Out-of-distribution (OOD) scenario (QH9):

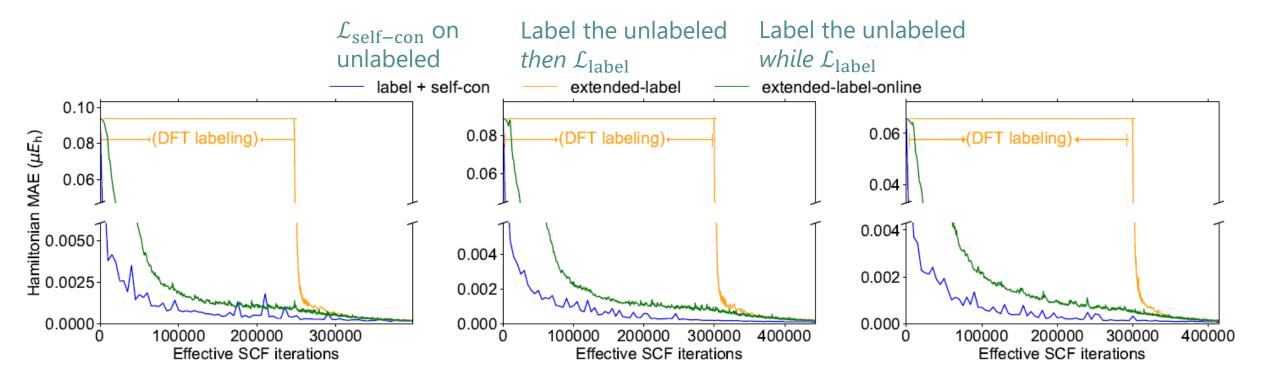
labeled small molecules + finetune on unlabeled large molecules  $\rightarrow$  test on large molecules.

Setting	$\mathbf{H}\left[ \mu E_{\mathbf{h}}\right] \downarrow$	$\mathbf{\epsilon} \left[ \mu E_{\mathbf{h}} \right] \downarrow$	$\mathbf{C}\left[\% ight]\uparrow$	$\epsilon_{\mathrm{HOMO}}\left[\mu E_{\mathrm{h}}\right]\downarrow$	$\epsilon_{\rm LUMO} \left[ \mu E_{\rm h} \right] \downarrow$	$\epsilon_{\Delta} \left[ \mu E_{\rm h} \right] \downarrow$	SCF Accel. [%] $\downarrow$
zero-shot	69.67	403.52	95.72	778.86	12230.49	12203.12	66.3
self-con (all-param)	65.74	375.31	<b>97.31</b>	565.50	<b>1130.55</b>	<b>1316.96</b>	<b>64.5</b>
self-con (adapter)	<b>64.48</b>	<b>268.83</b>	97.12	<b>449.80</b>	1220.54	1394.29	65.0

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#### · OOD scenario: labeled small molecules + finetune on unlabeled large molecules $\rightarrow$ test on large molecules. label + self-con extended-label extended-label-online (*nE*<sup>h</sup>) 0.00025 0.00020 0.00015 0.00010 →(DFT labeling)+ →(DFT labeling)+ 0.00025 0.00020 0.00015 0.00010 -0.00005 0.00005-200000 100000 300000 50000 100000 150000 200000 0 Computation time (s) Effective SCF iterations

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Setting	$\mathbf{H}\left[ \mu E_{\mathbf{h}}\right] \downarrow$	$\mathbf{\epsilon} \left[ \mu E_{\mathbf{h}} \right] \downarrow$	$\mathbf{C}\left[\% ight]\uparrow$	$\epsilon_{\mathrm{HOMO}}\left[\mu E_{\mathrm{h}}\right]\downarrow$	$\epsilon_{ m LUMO} \left[ \mu E_{ m h} \right] \downarrow$	$\epsilon_{\Delta}\left[\mu E_{\rm h}\right]\downarrow$	SCF Accel. [%] $\downarrow$
extended-label self-con	<b>59.67</b>	330.05	96.63	541.92	6372.12	6445.33	65.2
	64.48	<b>268.83</b>	97.12	<b>449.80</b>	<b>1220.54</b>	<b>1394.29</b>	<b>65.0</b>

• Direct efficiency comparison with DFT: time for solving MD17 structures under the same stopping criteria.

Molecule	criterion $[\mu E_h]$	$t_{\text{self-con}}\left[\mathbf{s}\right]$	$t_{ m DFT}[ m s]$
Ethanol	31.0	$4.50  imes 10^4$	$6.40 \times 10^{4}$
Malondialdehyd	e 88.9	$4.81  imes 10^4$	$1.05 \times 10^{5}$
Uracil	177.2	$1.23  imes 10^5$	$2.15 \times 10^{5}$

#### Extending Applicable Scale of Hamiltonian Prediction

· Labeled QH9 molecules ( $\leq$  31 atoms) + Finetune on unlabeled larger molecules  $\rightarrow$  Test on larger molecules (MD22).

Molecule	Setting	$\mathbf{H}\left[ \mu E_{\mathbf{h}}\right] \downarrow$	$\mathbf{\epsilon} \left[ \mu E_{\mathbf{h}} \right] \downarrow$	$\mathbf{C}\left[\% ight]\uparrow$	$\epsilon_{\mathrm{HOMO}}\left[\mu E_{\mathrm{h}}\right]\downarrow$	$\epsilon_{ m LUMO} \left[ \mu E_{ m h} \right] \downarrow$	$\epsilon_{\Delta}\left[\mu E_{\rm h}\right]\downarrow$	SCF Accel. [%] $\downarrow$
ALA3 42 atoms	zero-shot self-con	237.71 <b>52.49</b>	$6.54 \times 10^3$ <b>1.22</b> ×10 <sup>3</sup>	52.24 <b>94.46</b>	$6.90 \times 10^3$ 2.07 × 10 <sup>3</sup>	$9.51 \times 10^4$ 3.76×10 <sup>3</sup>	$9.79 \times 10^4$ <b>2.69</b> ×10 <sup>3</sup>	84.6 <b>64.7</b>
	e2e (ET) e2e (Equiformer)	N/A N/A	N/A N/A	N/A N/A	$1.74 \times 10^{5}$ $2.38 \times 10^{5}$	$7.72 \times 10^{3}$ $1.16 \times 10^{4}$	$2.38 \times 10^{5}$ $2.27 \times 10^{5}$	N/A N/A
DHA 56 atoms	zero-shot self-con	397.87 <b>56.12</b>	$1.84 \times 10^4$ <b>1.81</b> ×10 <sup>3</sup>	20.15 <b>83.51</b>	$1.11 \times 10^4$ <b>1.99</b> ×10 <sup>3</sup>	$1.90 \times 10^5$ 4.01×10 <sup>3</sup>	$1.85 \times 10^5$ 2.34×10 <sup>3</sup>	170.8 <b>67.0</b>
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- Labeled QH9 molecules ( $\leq$  31 atoms) + Finetune on unlabeled larger molecules  $\rightarrow$  Test on larger molecules (MD22).
- · Outperform end-to-end property predictors: merit of scientific-law supervision!

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