

# Going beyond Data Limitation with Physical Laws

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# Computational Methods for Molecular Science

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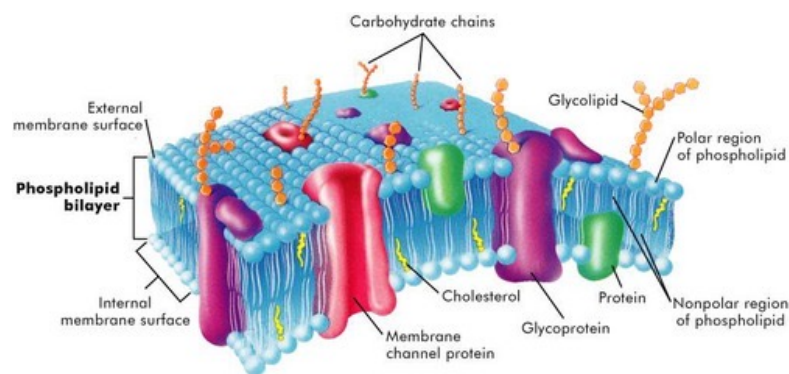
## Molecular science tasks

biomolecule understanding

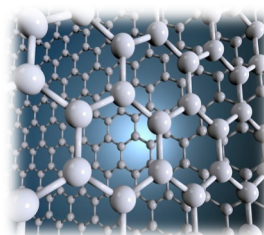
drug design

material/catalyst discovery

...

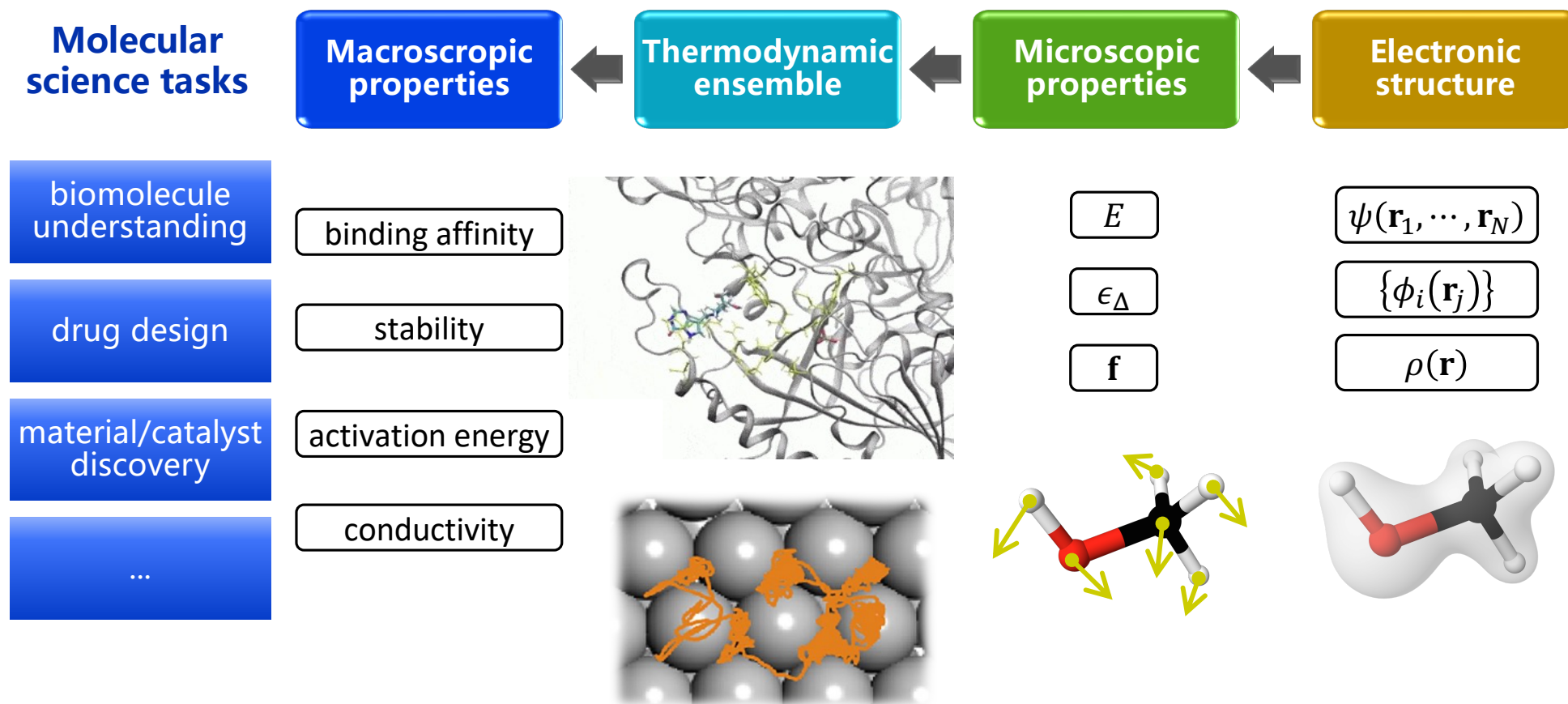


Modified from Thibodeau GA, Patton KT: Anatomy & physiology, ed 6. St Louis, 2007, Mosby.



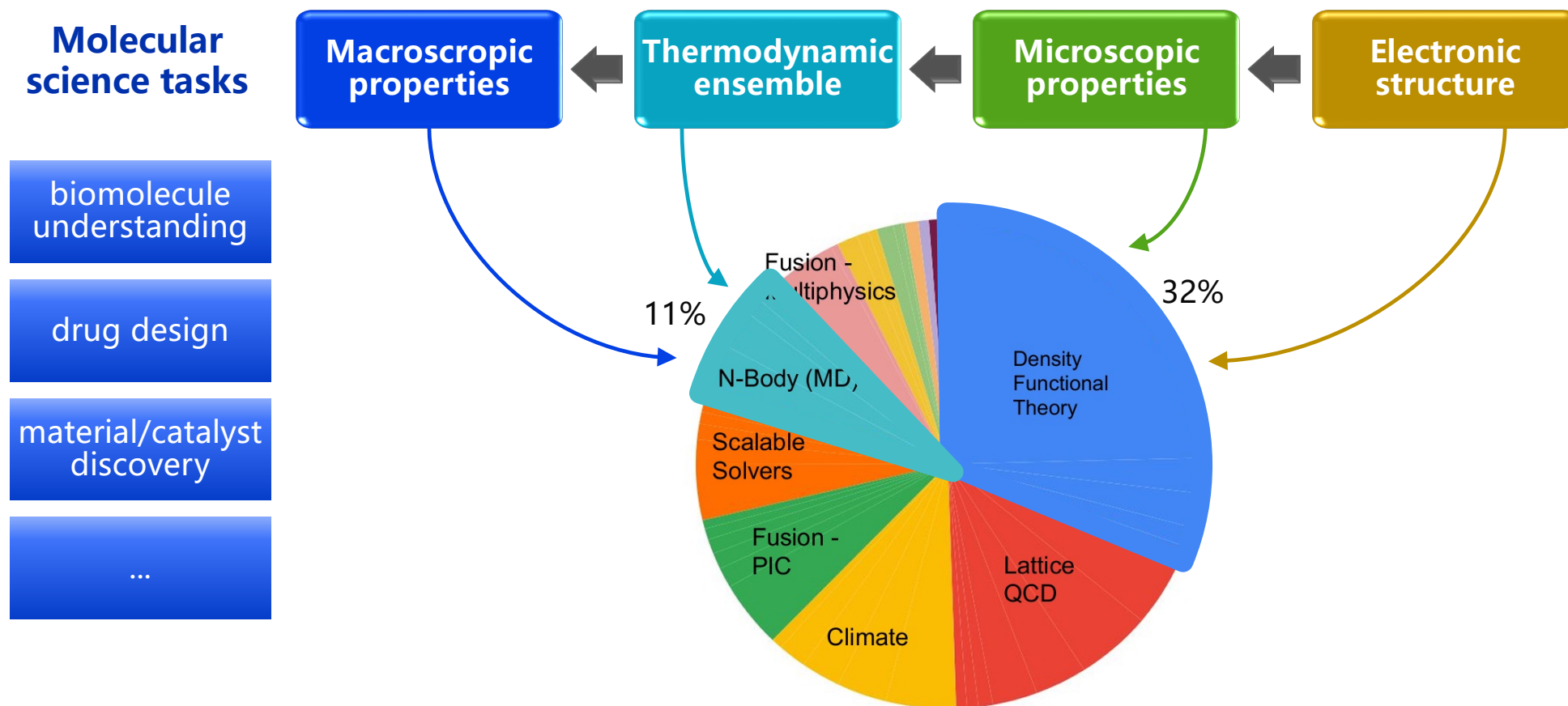
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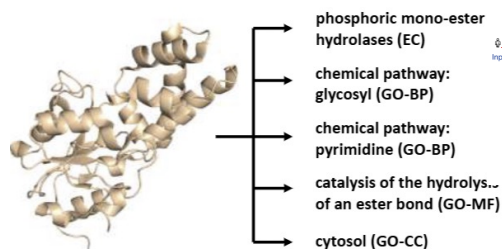


# AI for Scientific Computation

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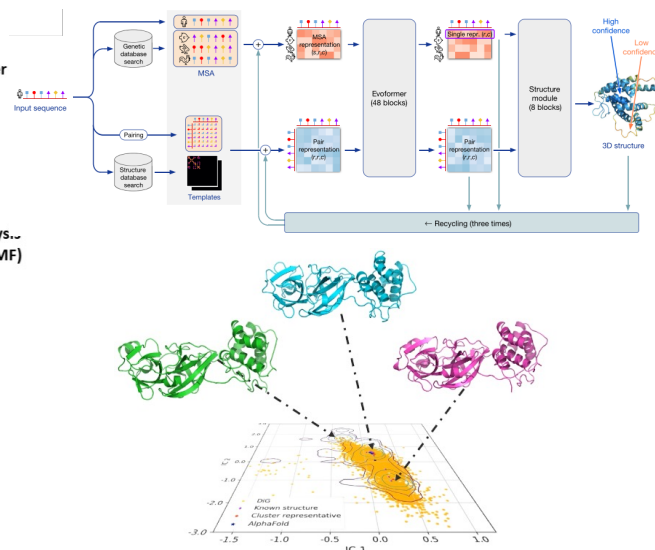
## Macroscopic properties

Macroscopic property prediction:  
scalability



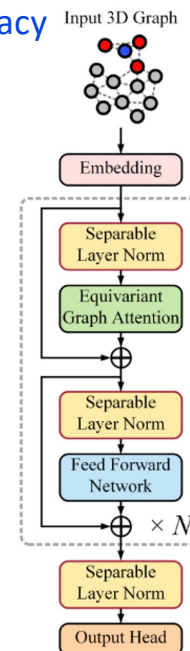
## Thermodynamic ensemble

Structure prediction and sampling:  
accuracy & scalability



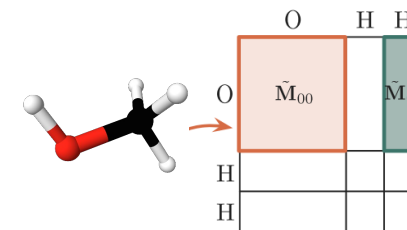
## Microscopic properties

ML force field:  
accuracy

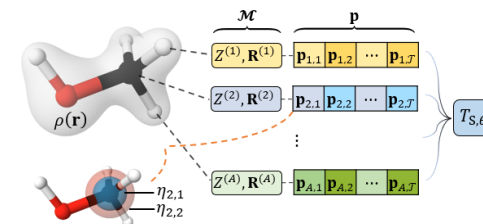


## Electronic structure

Hamiltonian prediction:  
scalability

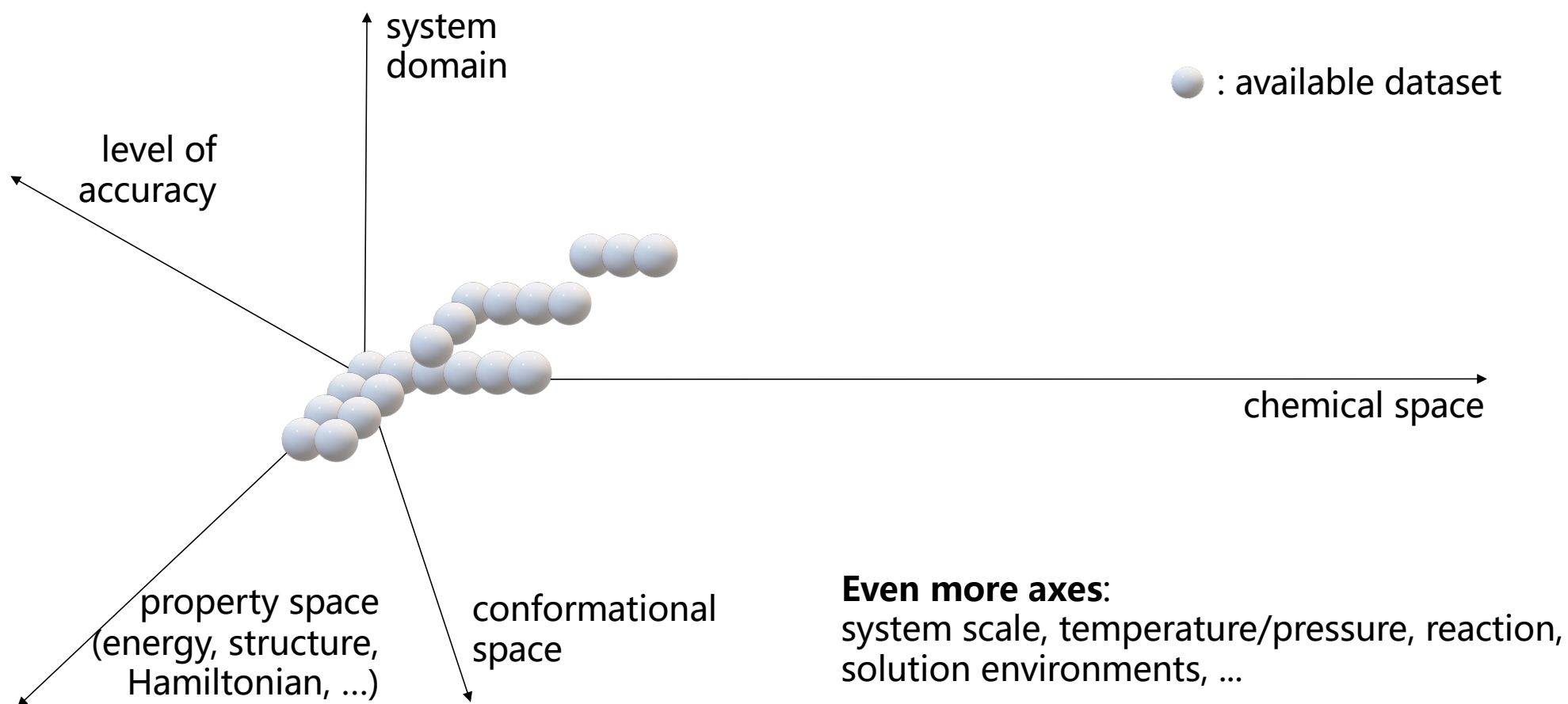


Learned functional:  
accuracy



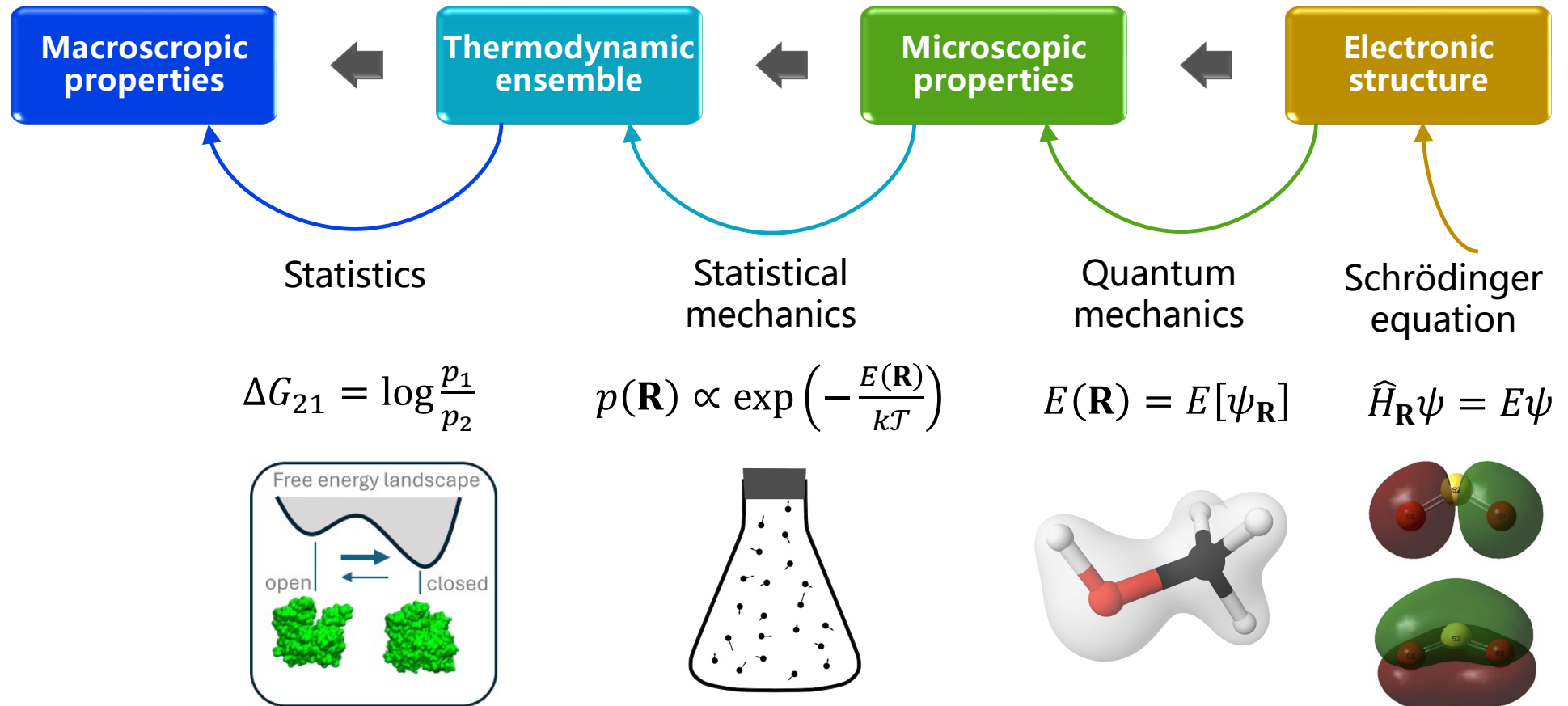
# The Curse: Data Scarcity

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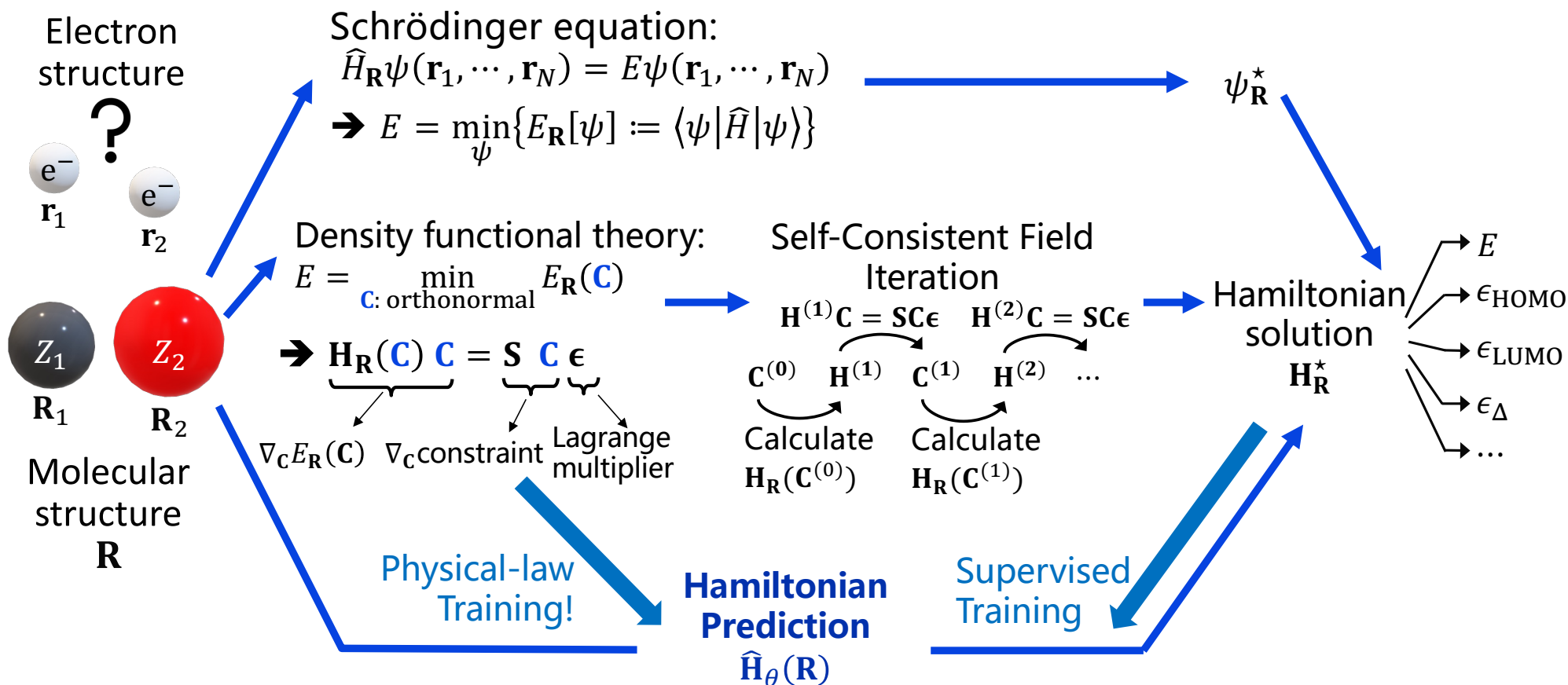
# The Blessing: Physical Laws!

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# Electronic Structure





- Self-consistency training for Hamiltonian prediction:

$$L_{\text{self\_con}}(\theta) = \left\| \hat{\mathbf{H}}_{\theta}(\mathbf{R}) - \mathbf{H}_{\mathbf{R}}\left(\mathbf{C}_{\mathbf{R}}\left(\hat{\mathbf{H}}_{\theta}(\mathbf{R})\right)\right) \right\|_{\text{F}}^2.$$

- **Label-free**: distinction from predicting other properties.
- Not just a regularization: it fully **determines** the solution.

# Physical-Law Training for Electronic Structure

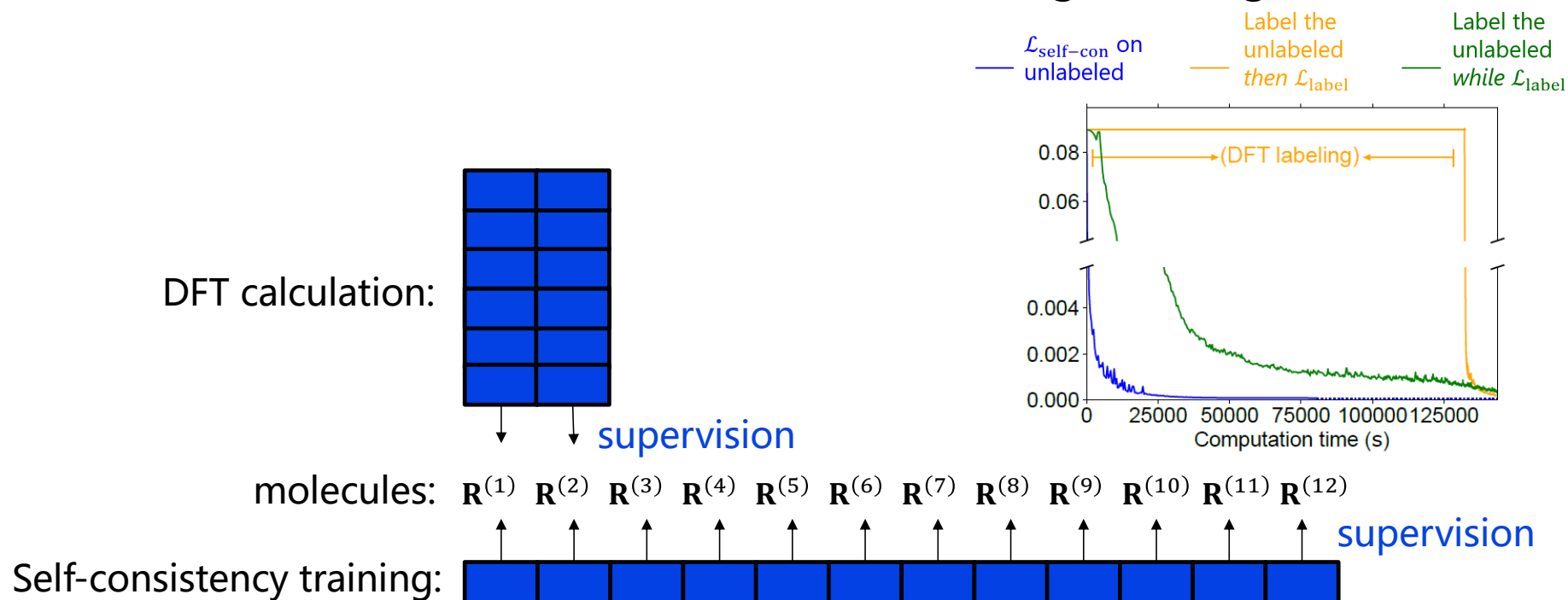
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- Generalization beyond available data

Largest labeled dataset ( $\leq 31$  atoms) + **unlabeled molecules**  $\rightarrow$  test on larger molecules

		Hamiltonian		Derived properties			
Molecule	Setting	$\mathbf{H}$ [ $\mu E_h$ ] $\downarrow$	$\epsilon$ [ $\mu E_h$ ] $\downarrow$	$\mathbf{C}$ [%] $\uparrow$	$\epsilon_{\text{HOMO}}$ [ $\mu E_h$ ] $\downarrow$	$\epsilon_{\text{LUMO}}$ [ $\mu E_h$ ] $\downarrow$	$\epsilon_{\Delta}$ [ $\mu E_h$ ] $\downarrow$
ALA3 (42 atoms)	zero-shot	237.71	$6.54 \times 10^3$	52.24	$6.90 \times 10^3$	$9.51 \times 10^4$	$9.79 \times 10^4$
	self-con	<b>52.49</b>	<b><math>1.22 \times 10^3</math></b>	<b>94.46</b>	<b><math>2.07 \times 10^3</math></b>	<b><math>3.76 \times 10^3</math></b>	<b><math>2.69 \times 10^3</math></b>
DHA (56 atoms)	zero-shot	397.87	$1.84 \times 10^4$	20.15	$1.11 \times 10^4$	$1.90 \times 10^5$	$1.85 \times 10^5$
	self-con	<b>56.12</b>	<b><math>1.81 \times 10^3</math></b>	<b>83.51</b>	<b><math>1.99 \times 10^3</math></b>	<b><math>4.01 \times 10^3</math></b>	<b><math>2.34 \times 10^3</math></b>

- Amortization effect: more efficient than running DFT to generate labels.



# Physical Law for Microscopic Properties

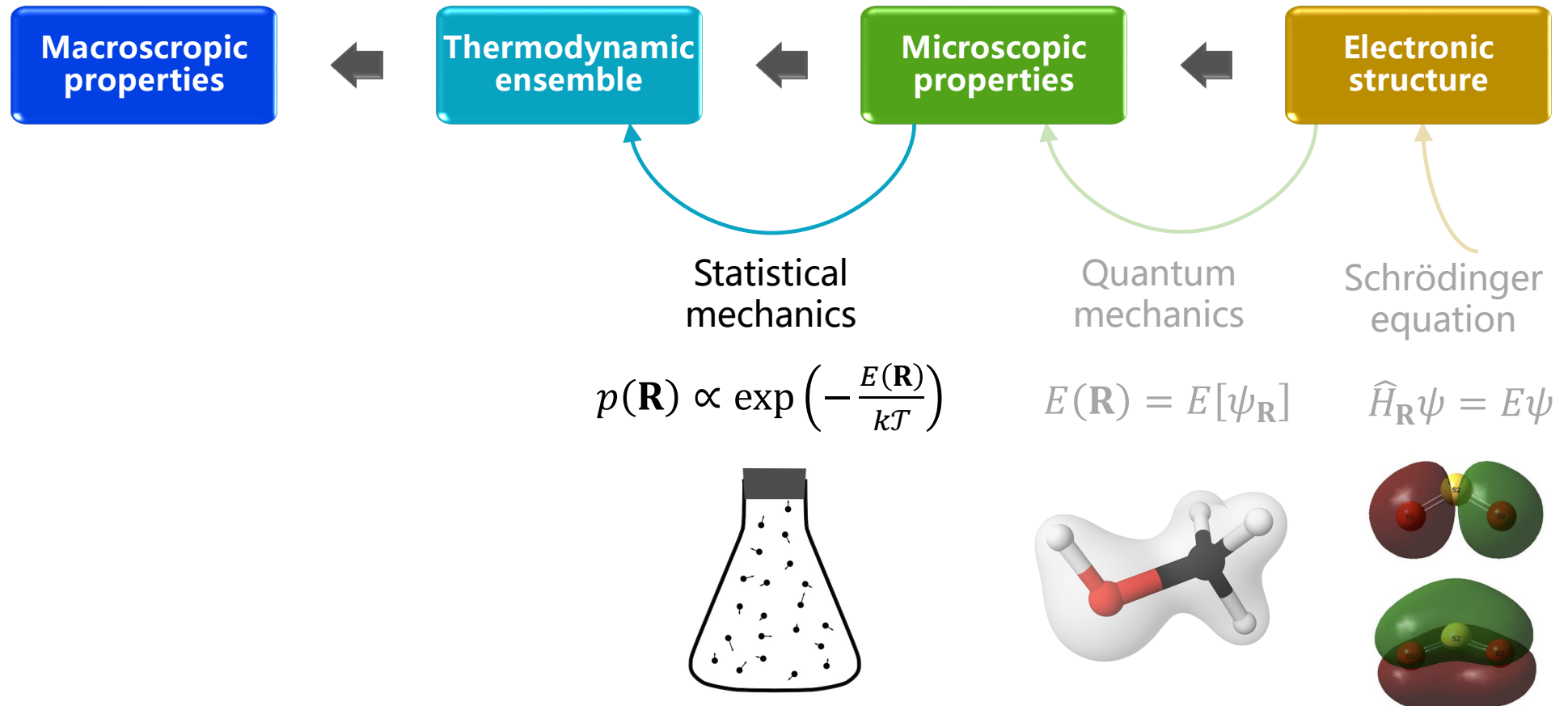
- Passing physical-law information to microscopic properties

labeled dataset ( $\leq 31$  atoms) + **unlabeled molecules**  $\rightarrow$  test on larger molecules

Molecule	Setting	$\epsilon_{\text{HOMO}} [\mu E_h] \downarrow$	$\epsilon_{\text{LUMO}} [\mu E_h] \downarrow$	$\epsilon_{\Delta} [\mu E_h] \downarrow$
ALA3 (42 atoms)	self-con	<b><math>2.07 \times 10^3</math></b>	<b><math>3.76 \times 10^3</math></b>	<b><math>2.69 \times 10^3</math></b>
	e2e (ET)	$1.74 \times 10^5$	$7.72 \times 10^3$	$2.38 \times 10^5$
	e2e (Equiformer)	$2.38 \times 10^5$	$1.16 \times 10^4$	$2.27 \times 10^5$
DHA (56 atoms)	self-con	<b><math>1.99 \times 10^3</math></b>	<b><math>4.01 \times 10^3</math></b>	<b><math>2.34 \times 10^3</math></b>
	e2e (ET)	$2.92 \times 10^5$	$2.58 \times 10^4$	$3.39 \times 10^5$
	e2e (Equiformer)	$3.76 \times 10^5$	$2.31 \times 10^4$	$4.17 \times 10^5$

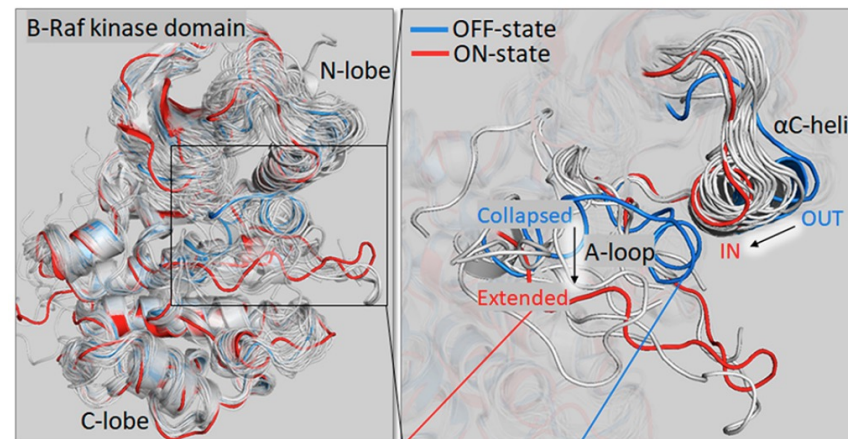
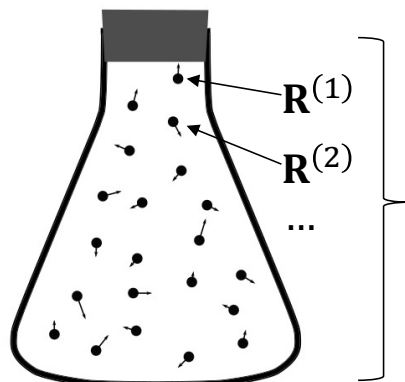
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# Thermodynamic Ensemble

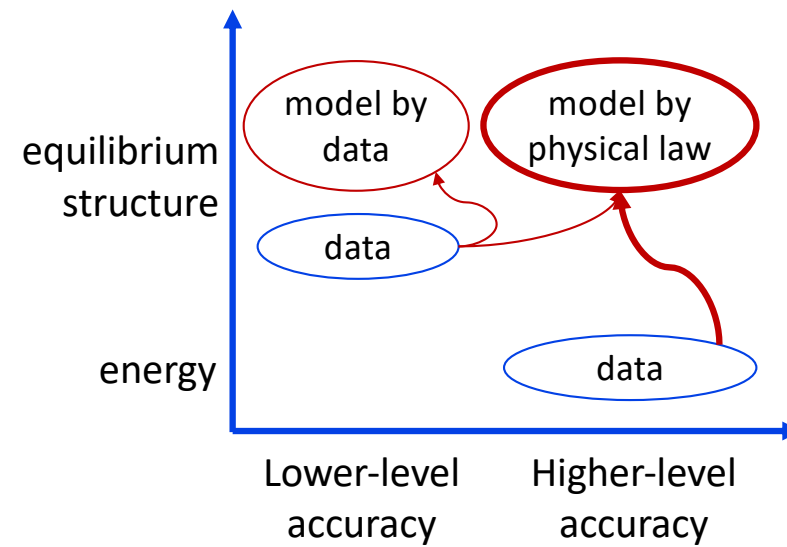
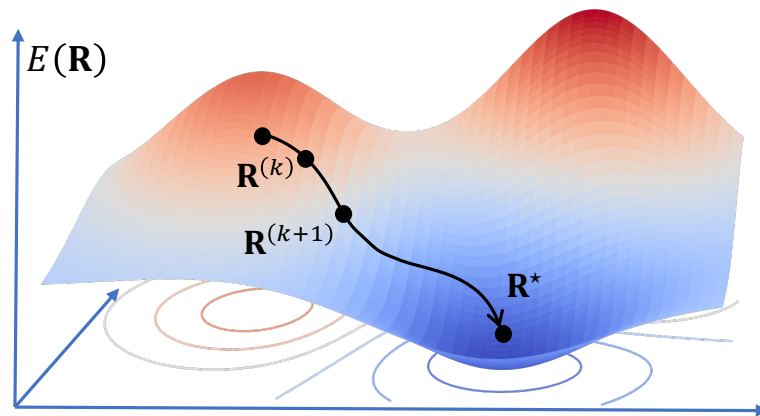
- A molecule exists in real world with structures  $\mathbf{R}$  following a distribution.
- More **detailed** description/knowledge



[https://en.wikipedia.org/wiki/Ensemble\\_\(mathematical\\_physics\)](https://en.wikipedia.org/wiki/Ensemble_(mathematical_physics))

# Data Heterogeneity: Energy and Equilibrium Structure

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# Connection between Structure and Energy

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- Optimality consistency:

$$\mathbf{R}^* = \operatorname{argmin}_{\mathbf{R}} E(\mathbf{R})$$

$$\rightarrow \min_{\theta} \mathbb{E} \max\{0, E(\mathbf{R}_{\theta}^*) - E(\mathbf{R}_{\theta}^* + \boldsymbol{\eta})\}.$$

$\mathbf{D}_{\theta}(\boldsymbol{\epsilon}, t)$  for large  $t \approx T$ .

- Score consistency:

$$\mathbf{R}^* \sim \exp\left(-\frac{E(\mathbf{R})}{k\mathcal{T}}\right) \text{ for small } \mathcal{T} \rightarrow$$

$$\min_{\theta} \mathbb{E}_{\mathbf{R}} \left\| \nabla \log p_{\theta}(\mathbf{R}) + \frac{\nabla E(\mathbf{R})}{k\mathcal{T}} \right\|^2.$$

$\frac{\alpha_t \mathbf{D}_{\theta}(\mathbf{R}, t) - \mathbf{R}}{\sigma_t^2}$  for small  $t \approx 0$

# Accuracy beyond Training Data

**Evaluation:**  
error w.r.t high-accuracy structure

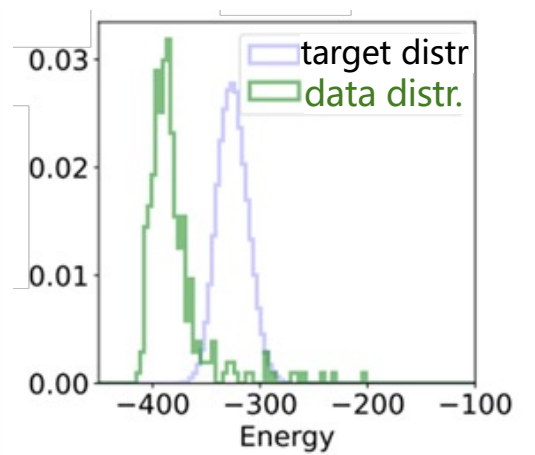
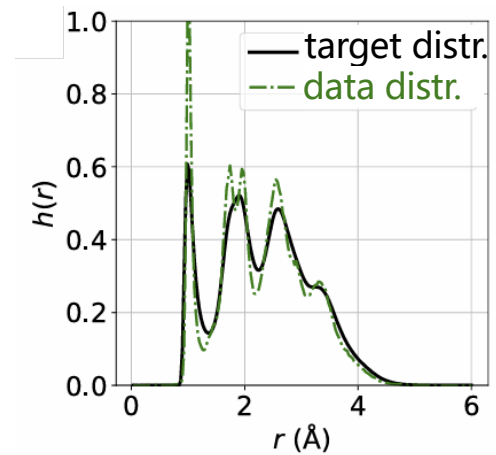
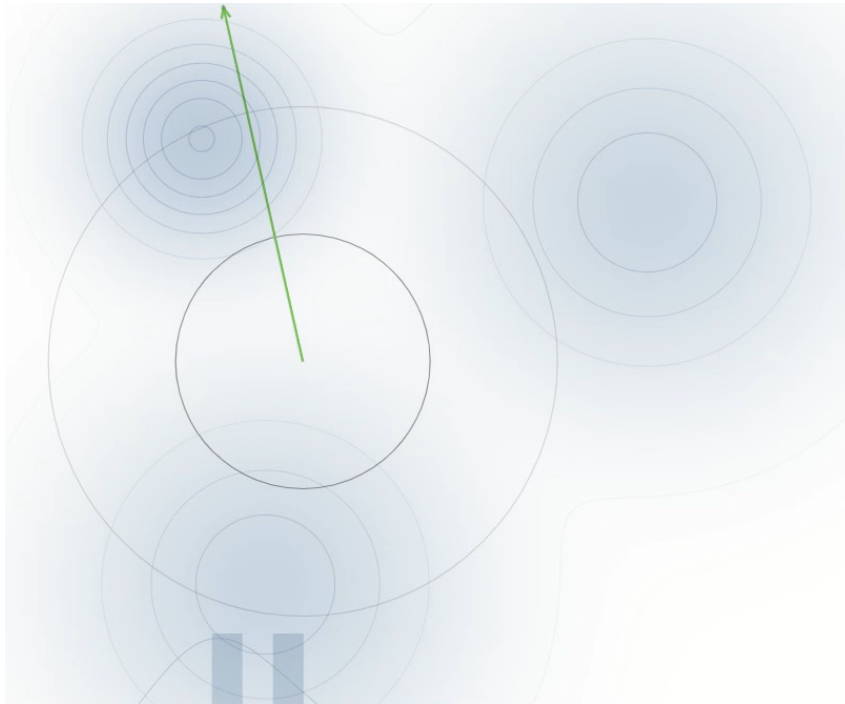
Test Set	PCQ		QM9	
	Mean	Min	Mean	Min
Struct. Stat.				
low-accuracy structure data	1.189	0.655	0.928	0.545
+ high-accuracy energy data	<b>1.158</b>	<b>0.645</b>	<b>0.848</b>	<b>0.490</b>

Ren et al. Physical Consistency Bridges Heterogeneous Data in Molecular Multi-Task Learning. *NeurIPS*, 2024.

# Problem with Learning Thermodynamic Ensemble

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- Data are biased from finite-length simulation



# Connection between Distribution and Energy

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- Energy function defines the target:  $p_0(\mathbf{R}_0) \propto \exp\left(-\frac{E(\mathbf{R}_0)}{kT}\right)$ .
- Diffusion-model learning target:

$$\mathbf{s}_\theta(\mathbf{R}_t, t) \rightarrow \operatorname{argmin}_\theta \mathbb{E}_{\underbrace{p_0(\mathbf{R}_0)}_{\approx q(\mathbf{R}_0)} p(\mathbf{R}_t|\mathbf{R}_0)} \left\| \mathbf{s}_\theta(\mathbf{R}_t, t) - \frac{1}{\alpha_t} \underbrace{\nabla \log p_0(\mathbf{R}_0)}_{-\frac{\nabla E(\mathbf{R}_0)}{kT}} \right\|^2.$$

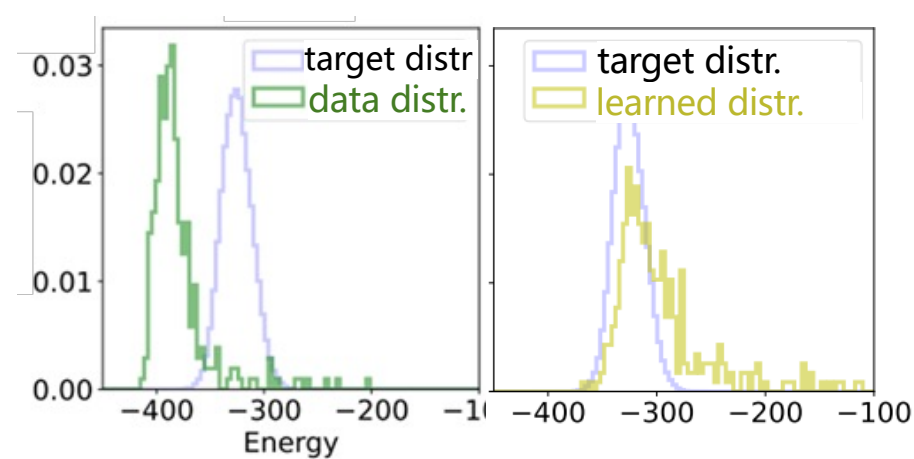
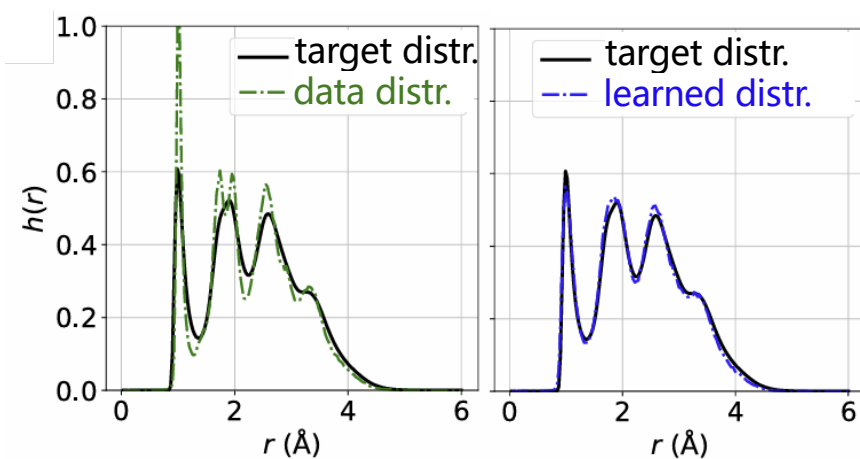
$$\mathbf{s}_{\theta_{\text{debiased}}}(\mathbf{R}_t, t) \rightarrow \underbrace{\approx q(\mathbf{R}_0)}_{\text{data distribution}}$$

- Energy function corrects data bias:

$$\left\| \mathbf{s}_{\theta_{\text{debiased}}}(\mathbf{R}_t, t) - \nabla \log p_t(\mathbf{R}_t) \right\|^2 \leq \left\| \mathbf{s}_{\theta_{\text{data}}}(\mathbf{R}_t, t) - \nabla \log p_t(\mathbf{R}_t) \right\|^2.$$

# Accuracy beyond Training Data

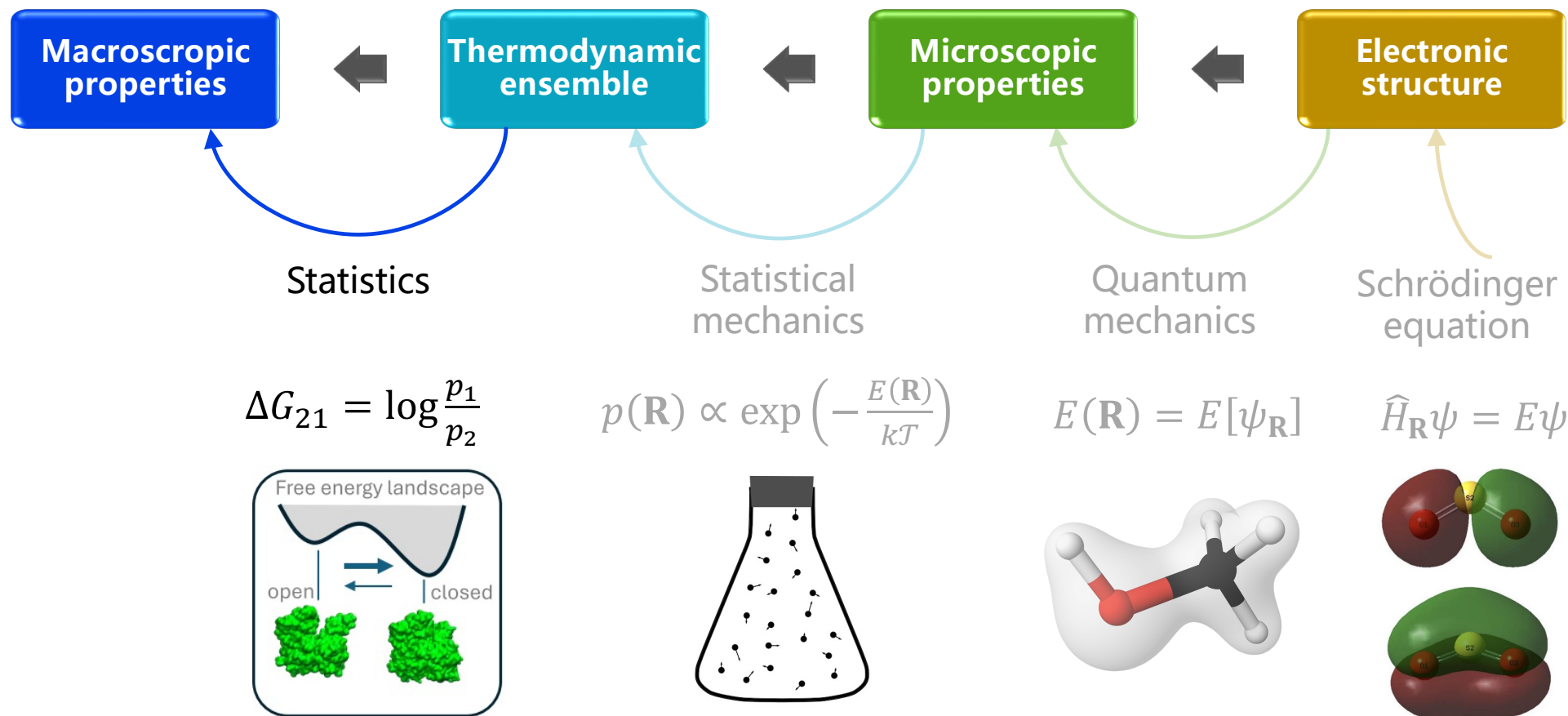
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Guo et al. Potential Score Matching: Debiasing Molecular Structure Sampling with Potential Energy Guidance. *TMLR*, 2025.

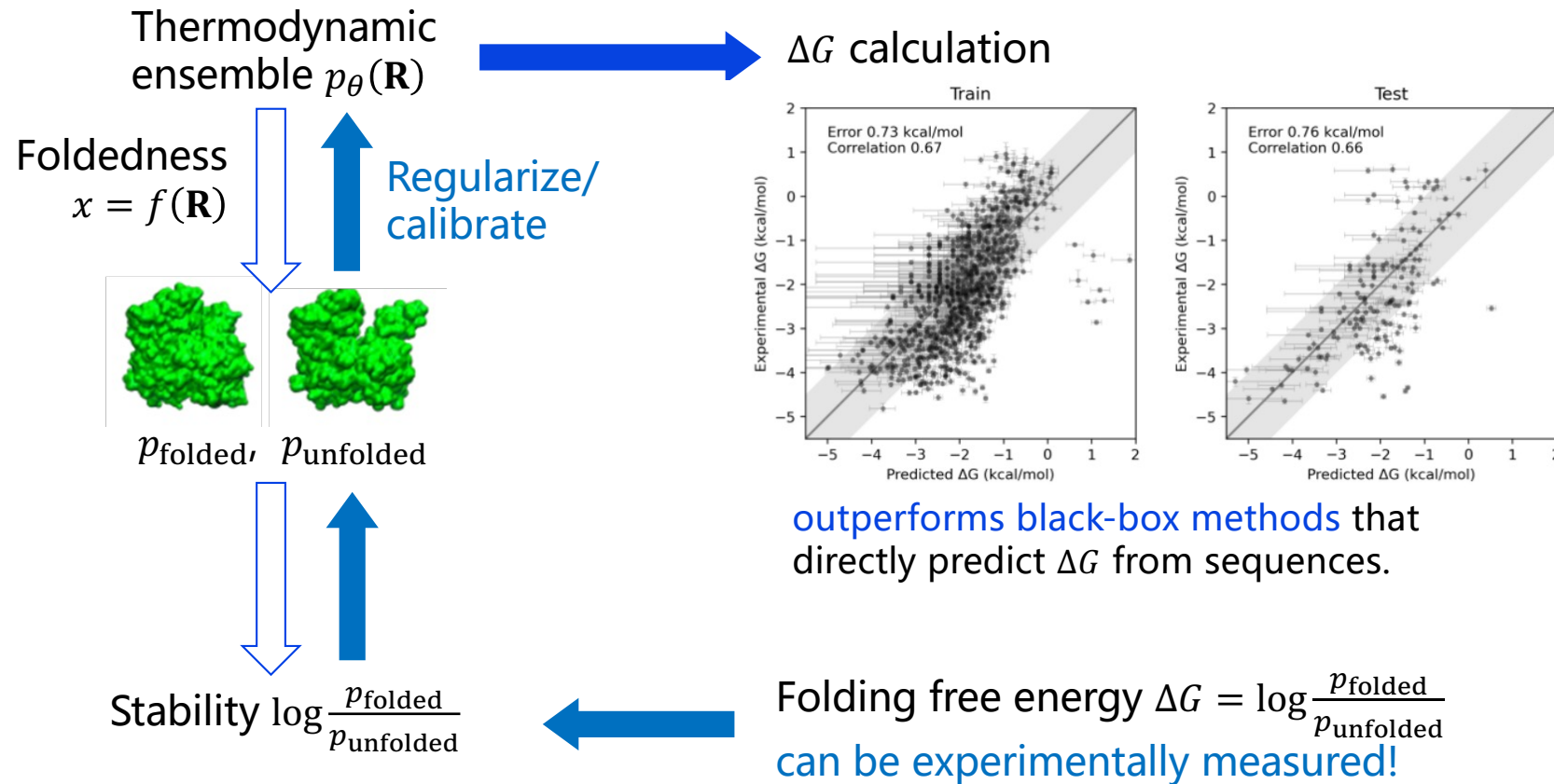
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# Stability Calculation from Ensemble Model

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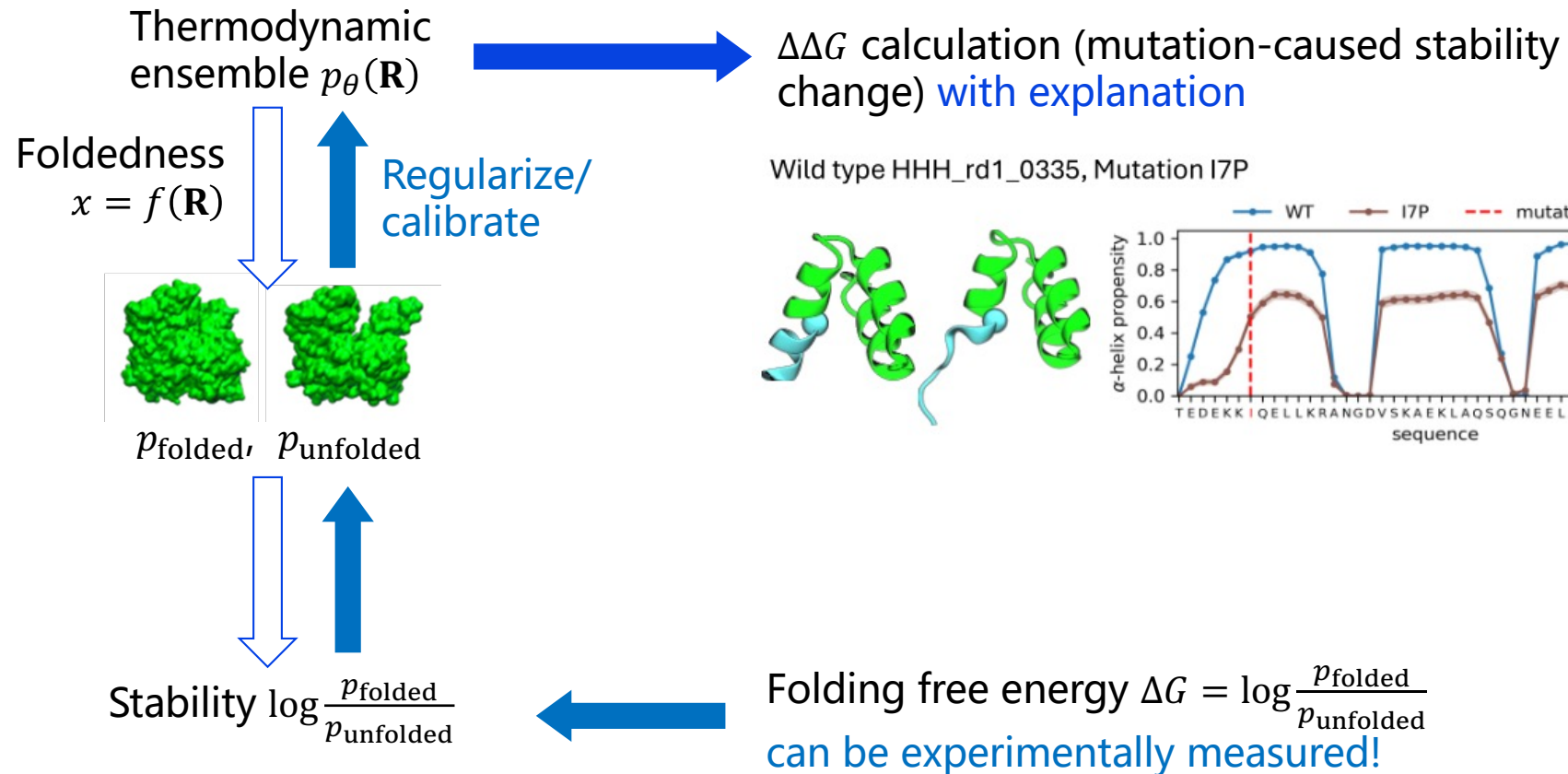


Lewis et al., Scalable emulation of protein equilibrium ensembles with generative deep learning, *Science*, 2025.



# Stability Calculation from Ensemble Model

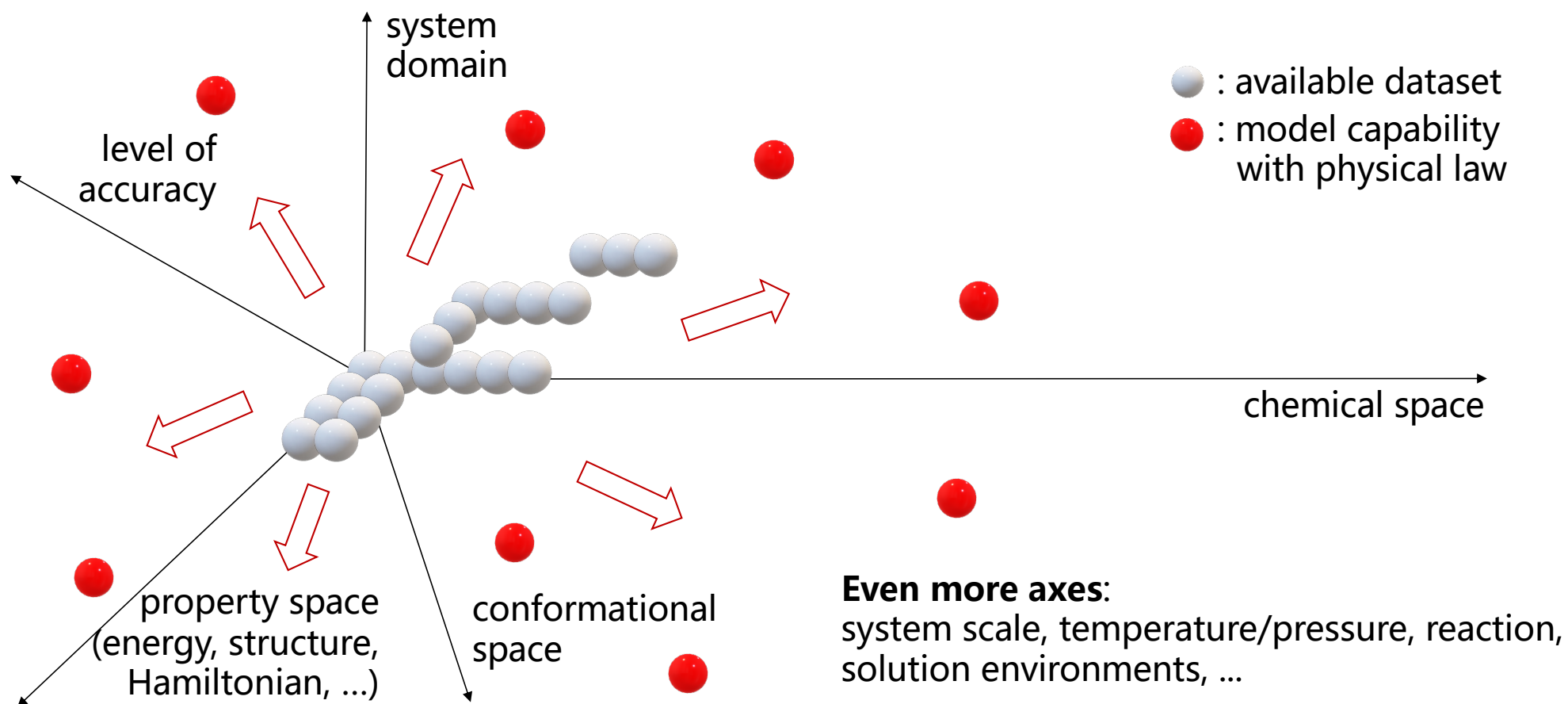
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Lewis et al., Scalable emulation of protein equilibrium ensembles with generative deep learning, *Science*, 2025.

# Physical Law: Reasoning in Scientific Computing

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Thank you