

# MolSets: Molecular graph deep sets learning for mixture property modeling

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# Introduction



(Generated by DALL·E)

Fascination of chemistry:  
mix things up and see what happens

Molecular mixture: a broad search space  
for materials discovery



coolant



fuel



battery  
electrolyte

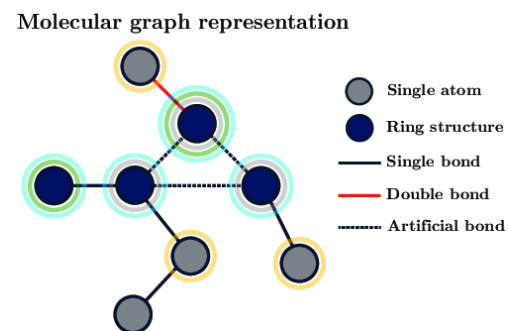
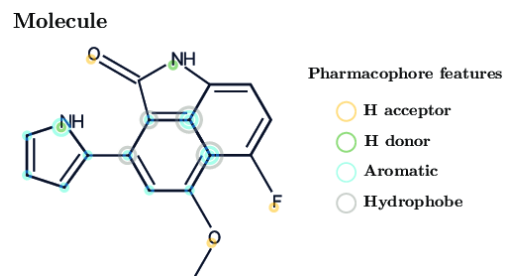
Large, combinatorial space  
Machine learning (ML) expedites the search

Challenge: multi-level complexity

- Diverse constituents × configurations

# Addressing local and global complexities

Locally, molecules have diverse chemistry and geometries.



## Graph representation

- Atoms  $\rightarrow$  nodes
- Atomic properties
- Bonds  $\rightarrow$  edges

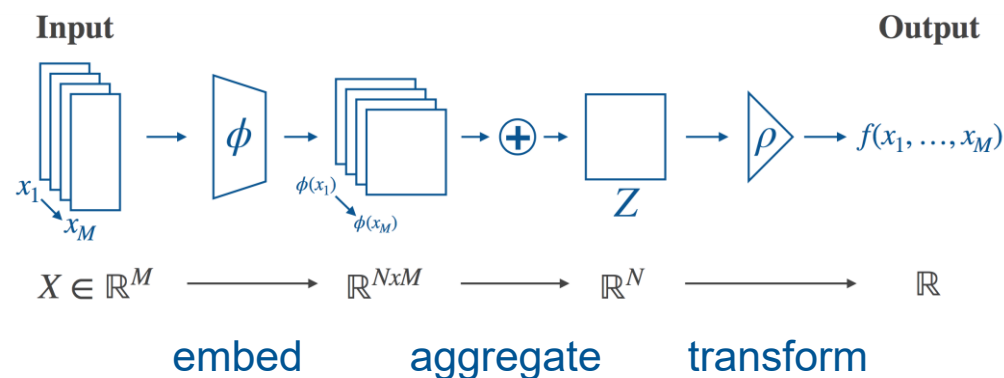
## Graph neural network

- Message passing between nodes
- Readout

Globally, mixture should be permutation invariant

$$f(\{x_1, x_2, x_3\}) = f(\{x_2, x_1, x_3\})$$

The “Deep Sets” model architecture

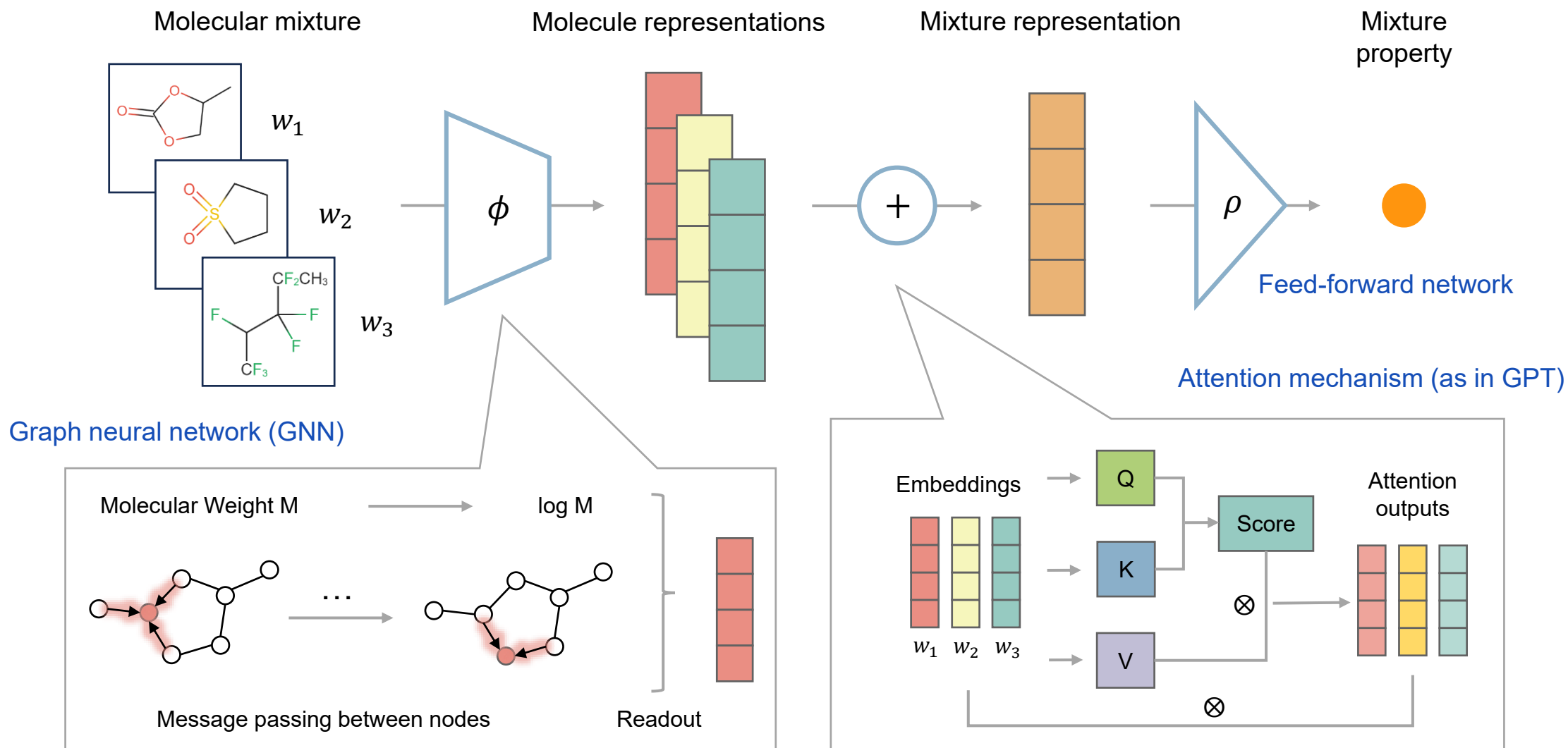


Molecular mixtures as sets:

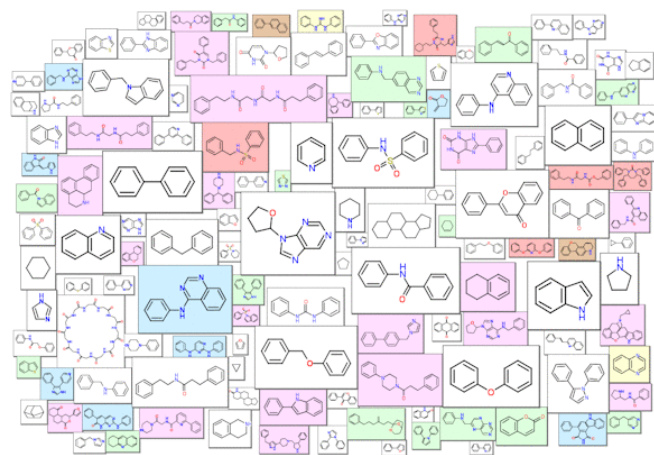
$$\{(x_1, w_1), (x_2, w_2), \dots\}$$

\*  $w$ : weight fraction,  $\neq$  “influence” fraction

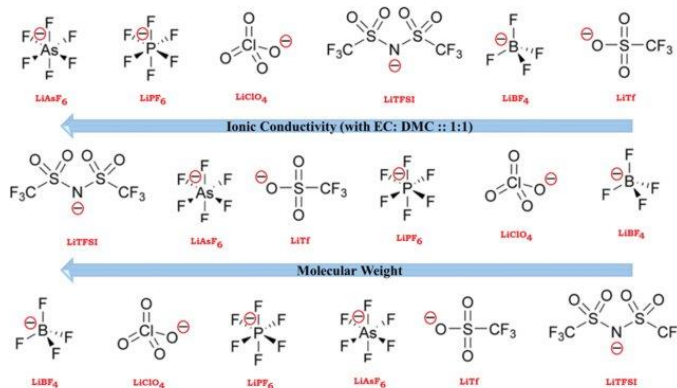
# MolSets model architecture



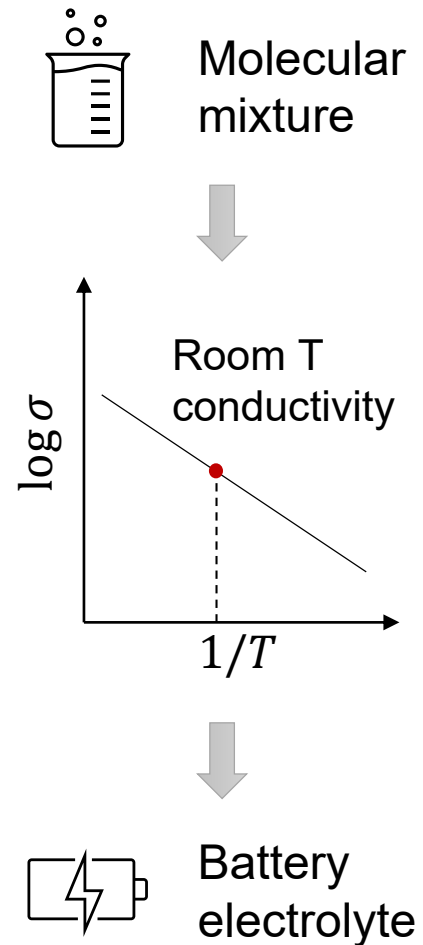
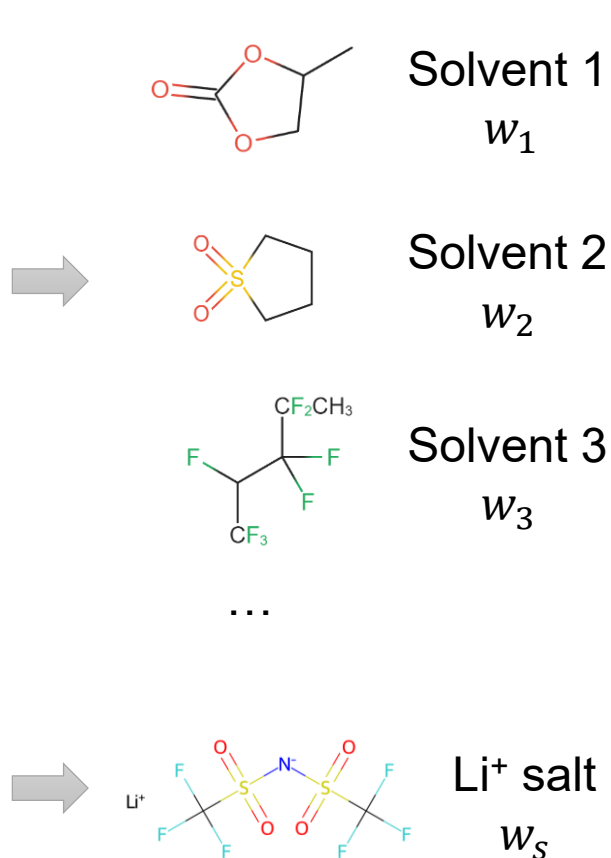
# Molecular mixture electrolytes



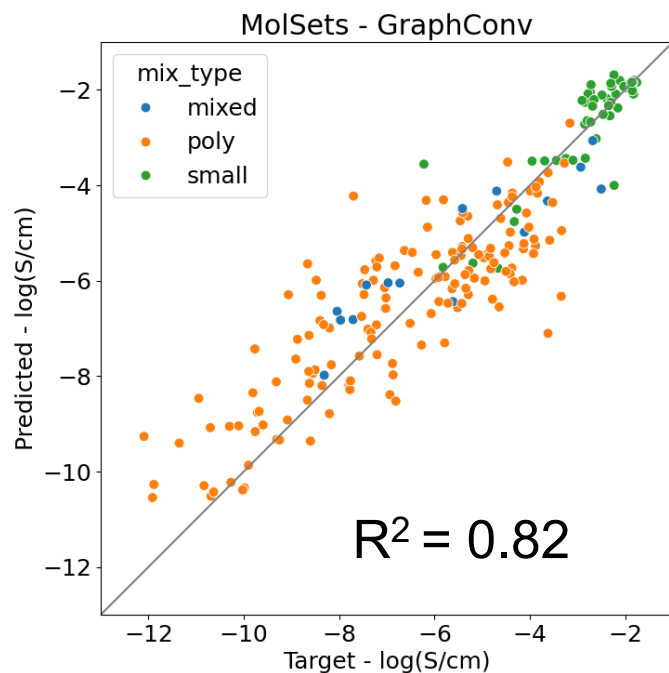
~200 molecules (incl. polymers)



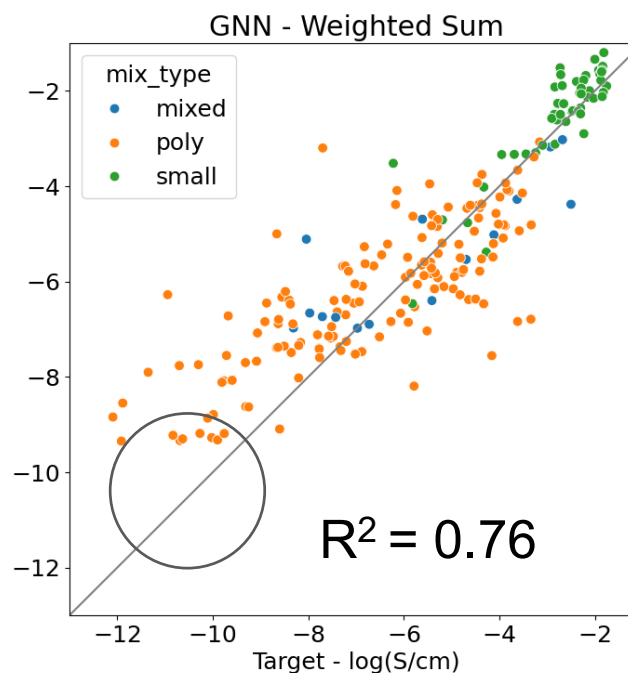
~50 Li<sup>+</sup> salts



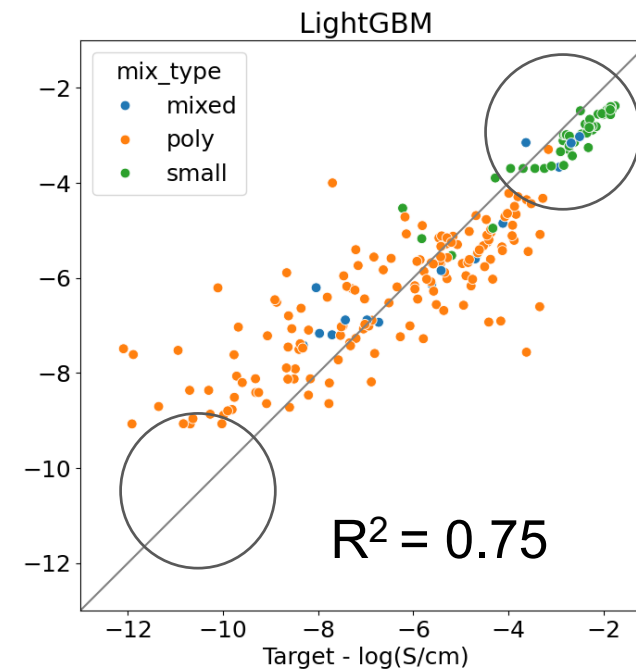
## MolSets



Ablation test:  
replace  $\oplus$  with weighted sum



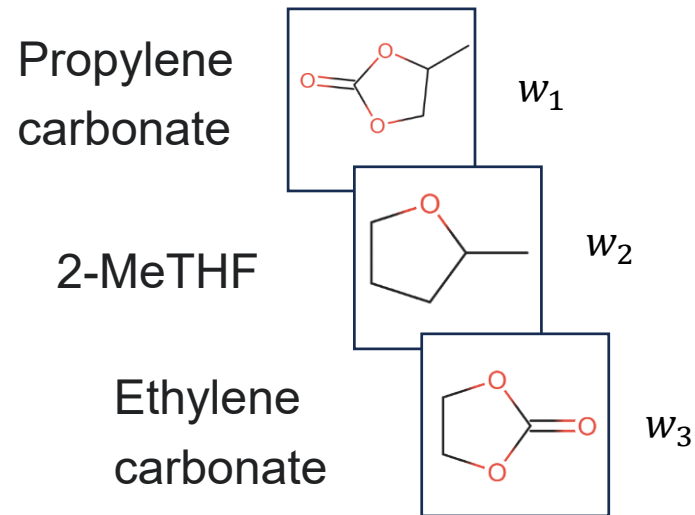
## Gradient boosting



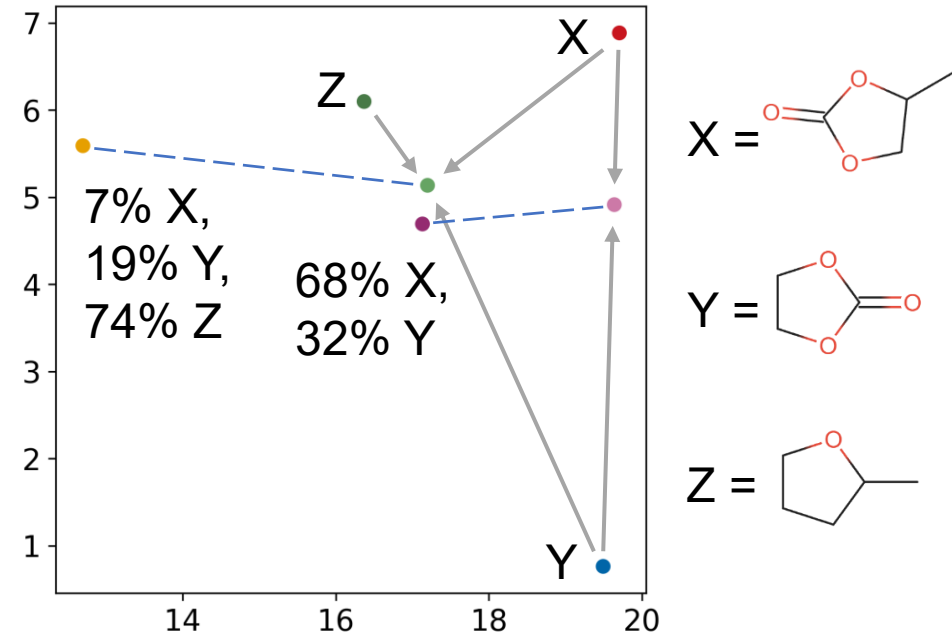
Tests performed on an electrolyte dataset collected in [1] from experimental literature  
~1100 unique mixtures; train : validation : test = 3 : 1 : 1

# Interpretation

- How is mixture representation different from linear combination of constituents?



$\phi$  and  $\oplus$   
t-SNE

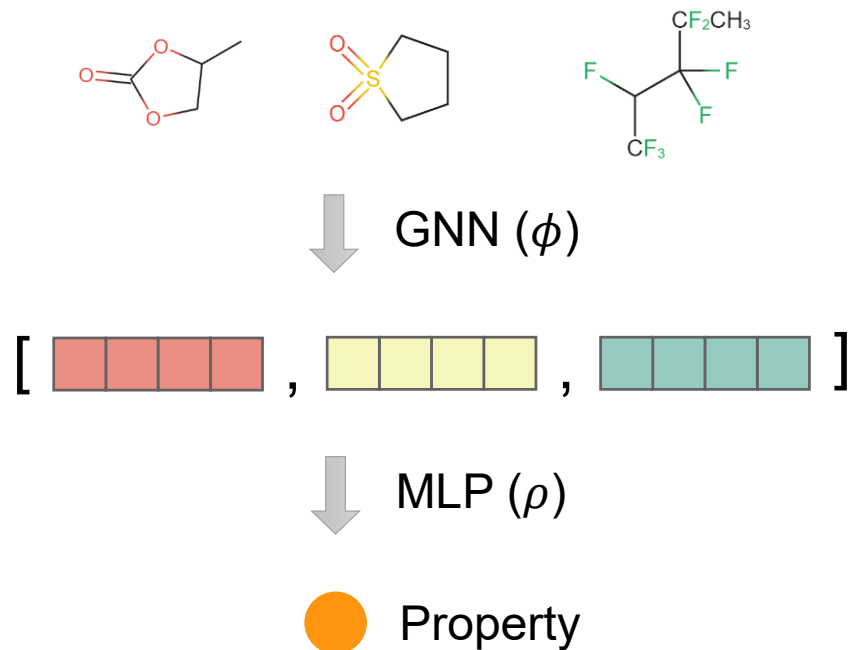


$\phi$  and  $\oplus$  learns a representation space of mixtures  
t-SNE projects the space into 2 dimensions

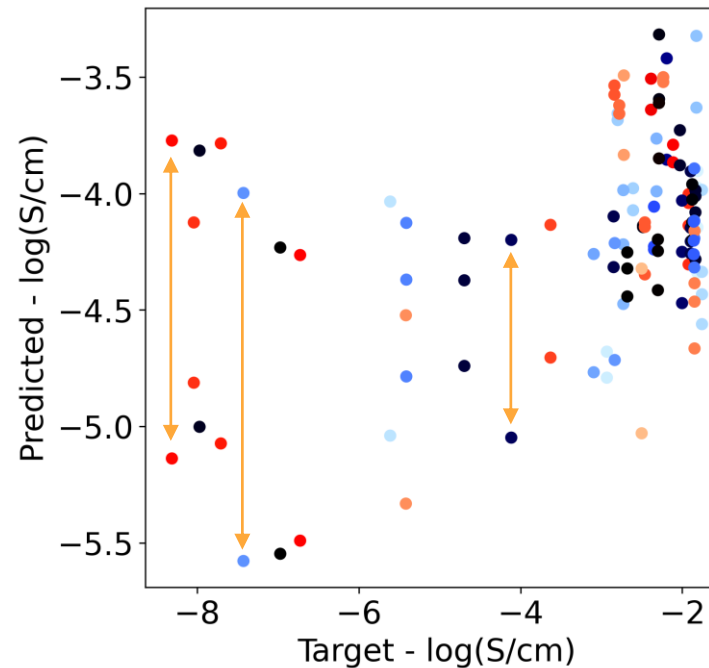
→ Linear combination of ingredients  
--- Mixture representation

# Interpretation

- What happens if a model doesn't preserve permutation invariance?



Concatenate molecular representations  
instead of aggregating



Same mixture, change sequence →  
predicted differently



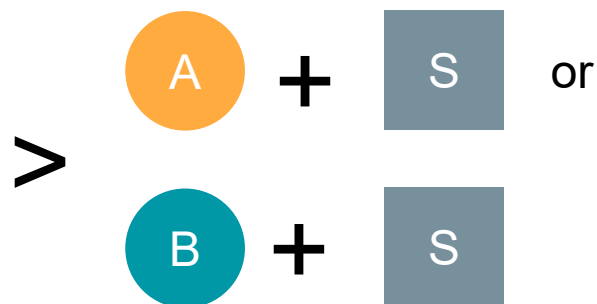
# Virtual screening

- Train on all data from [1], predict 298 K conductivity for 11,340 new candidates
  - Equal-weight binary mixture among 28 molecules  $\times$  30 salts (1 mol/kg)
- Data available on [Dryad](#)

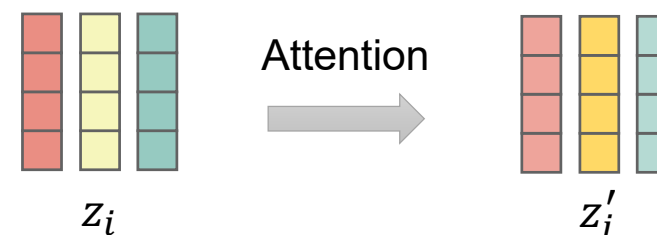


Dryad dataset

“Bowing effect”  
(nonlinear mixing behavior)



Relative importance



Magnitude change  $\|z'\|/\|z\|$  indicates importance

Molecules found important include:

- Cyclic carbonate esters (ion solvation ability)
- Benzene, toluene (incorrect)



A limitation: current model didn't learn solubility

Reason: limited data availability

- In experimental datasets, salt never exceeds solubility

⚠ Rationally choose salt molality in application

Future work directions

- Synergy with high-throughput experiments: “robots” make mixtures and measure properties
    - e.g., Coulomb efficiency, order parameters
  - Build a data platform: access MolSets' predictions
- ▶ “AlphaFold” or “matterverse” for mixtures



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# Thank you!



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