



The Rapidly Expanding
Frontier of the Possible in
Reaction Planning

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P2SAC Spring Conference, Purdue University, 5/8/24

Challenges of Contemporary Group Theories

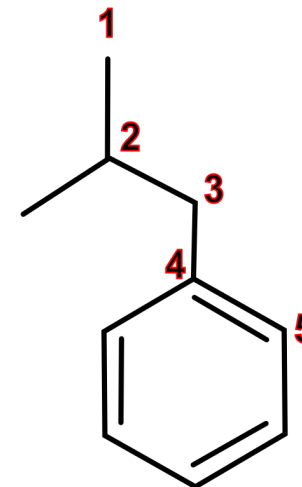
Benson Group Theory:

- The idea is to decompose molecular properties (ΔH_f , S° , C_v) as the sum of “group” contributions.
- Group contributions are calculated based on trusted experimental or computational data, and transferability is assumed.

Problems we want to address:

- **Specificity:** the definition of a “group” has never been formalized and inconsistent granularity is applied.
- **Provenance:** inconsistent thermodynamic data is available/used to determine group contributions.
- **Extensibility:** because of the provenance and specificity problems, it isn't possible to develop new groups in a consistent way.

From Anslyn and
Dougherty's
Textbook



1) C -(C)(H) ₃	2(-10.20)
2) C -(C) ₃ (H)	-1.90
3) C -(C _B)(C)(H) ₂	-4.86
4) C _B -(C)	5.51
5) C _B -(H)	5(3.30)
	<hr/>
	-5.15 kcal/mole (-21.6 kJ/mole)

Experimental ΔH_f : -5.15 +/- 0.34 kcal/mol

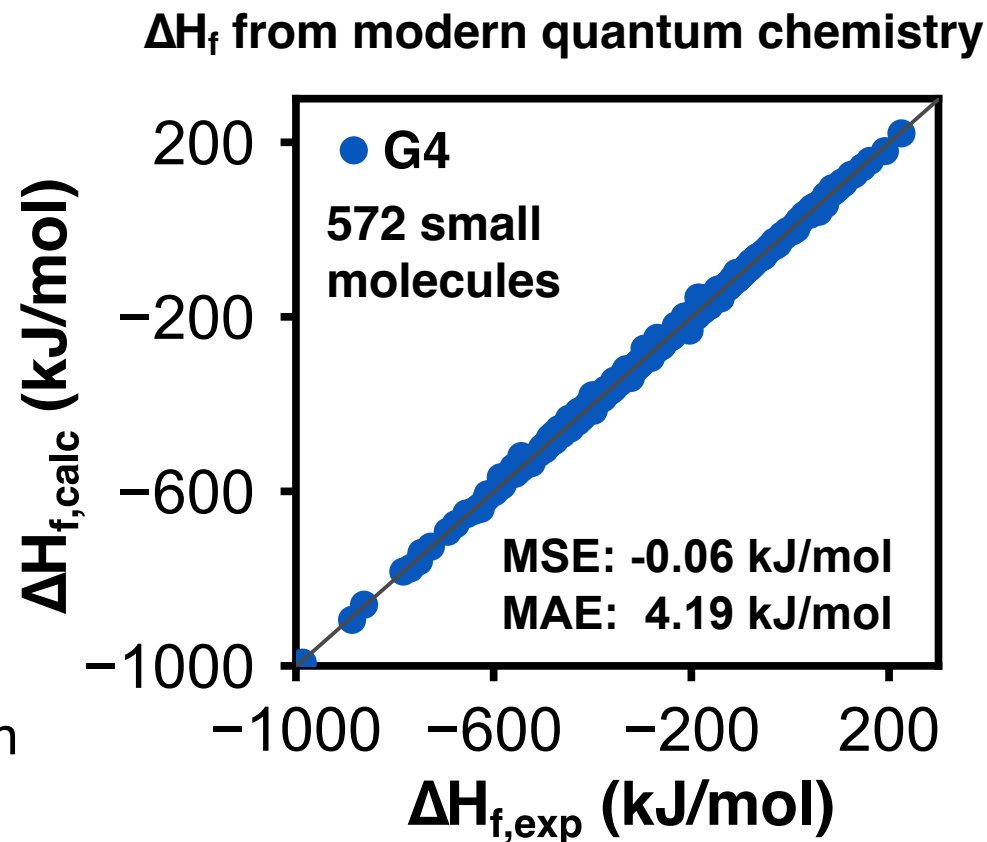
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Zhao, Q.; Savoie, B. M.; Enthalpy of Formation Prediction via a fully Self-Consistent Component Increment Theory. *J. Chem. Info. Model.* **2020**, 60, 2199-2207

Challenges of Contemporary Group Theories

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- Group
- on trust
- data, a

Prob

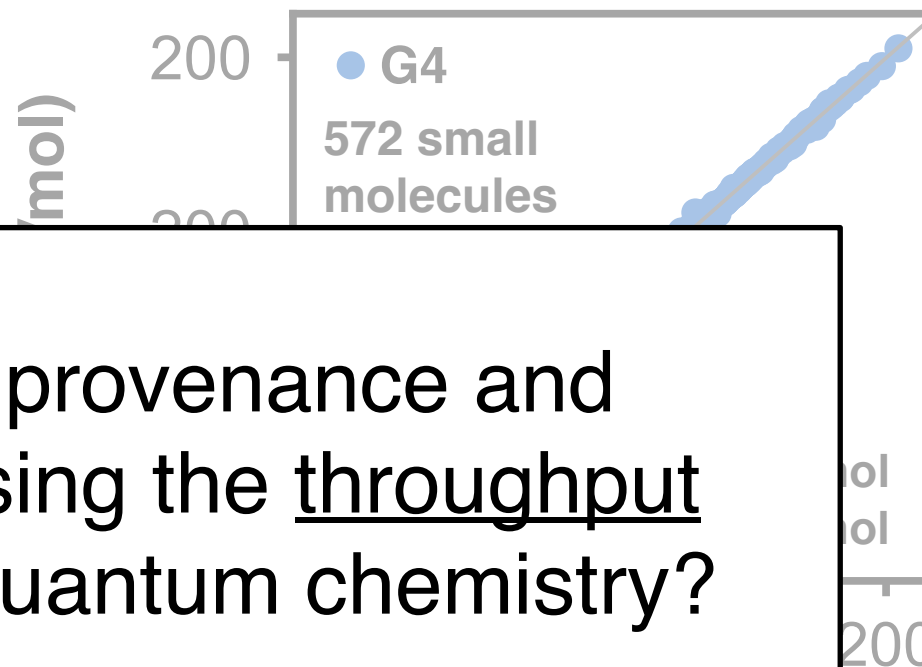
- Spec
- formal

Can we circumvent the provenance and extensibility challenges using the throughput and accuracy of modern quantum chemistry?

- **Provenance:** inconsistent thermodynamic data is available/used to determine group contributions.

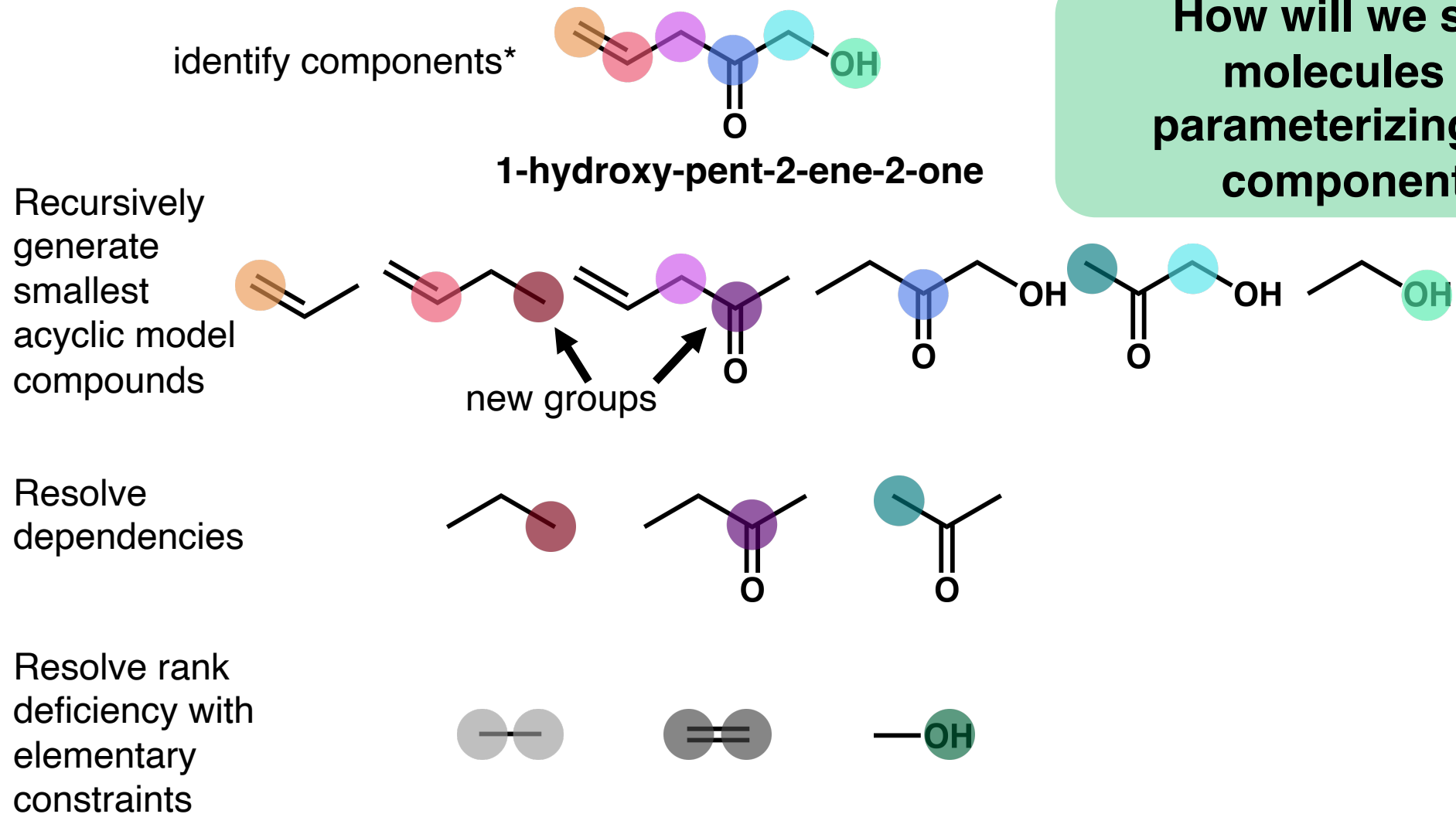
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ΔH_f from modern quantum chemistry



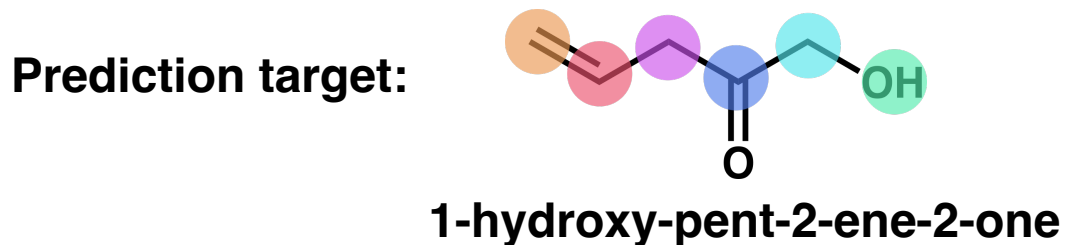
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Graphical Decomposition of Model Compounds

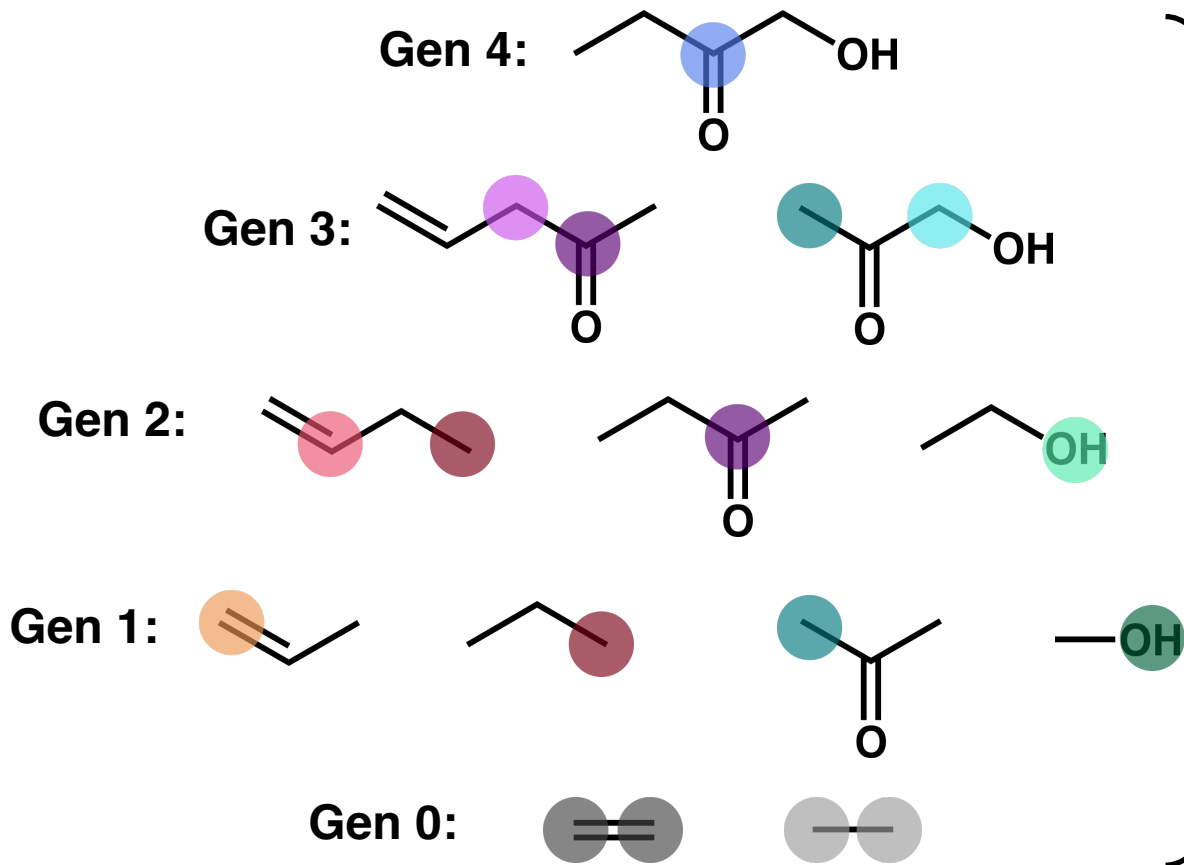
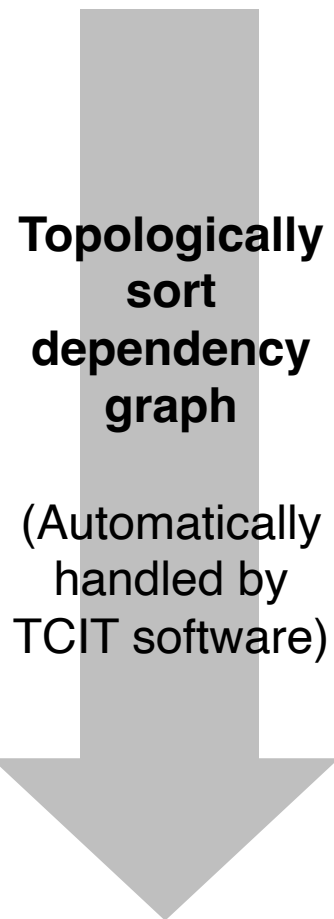


How will we select molecules for parameterizing TCIT components?

Graphical Decomposition of Model Compounds



$\Delta H_{f,G4} = -259.9 \text{ kJ/mol}$
 $\Delta H_{f,TCIT} = -259.3 \text{ kJ/mol}$
no experimental data



Model compounds are small enough to perform the highest quality quantum chemistry calculations (G4 throughout)

Graphical Decomposition of Model Compounds

Have we solved the specificity problem?

All components are unique out to a graph depth of two, no exceptions.

Have we solved the provenance problem?

All ΔH_f data is calculated at the G4 composite level, no exceptions.

Have we solved the extensibility problem?

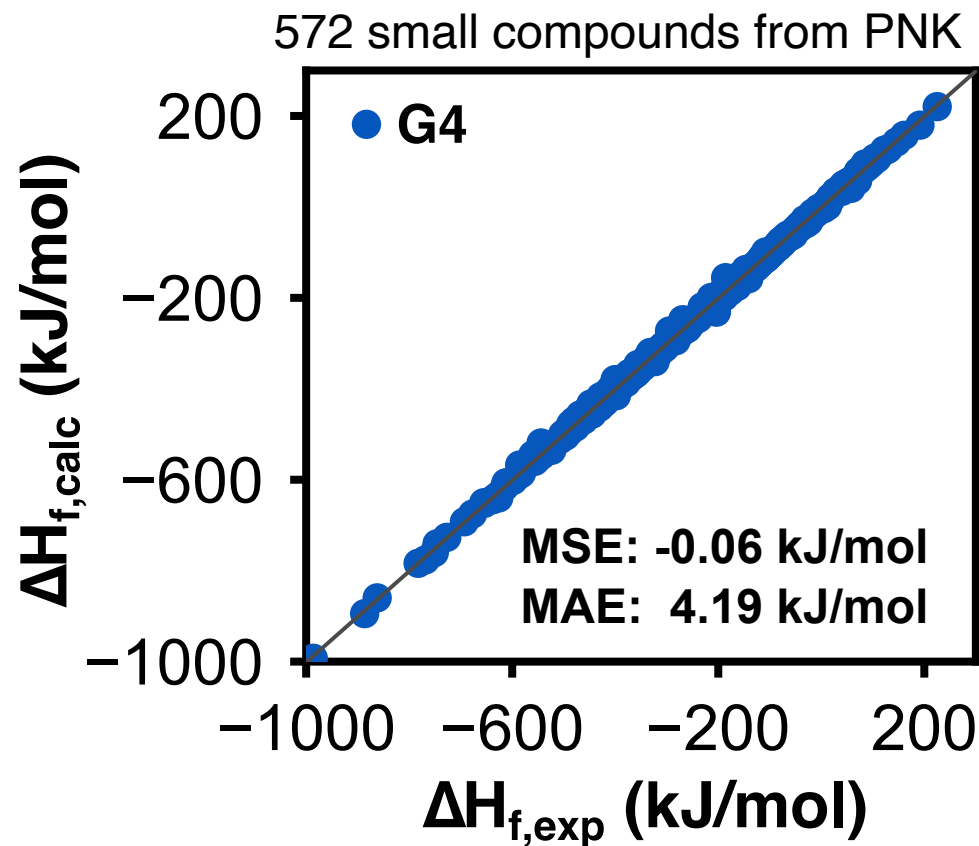
Model compounds exist for all conceivable components, no exceptions.

Benchmarking $\Delta H_{f,gas}$ Predictions Against the PNK Dataset

- Initial benchmarking set consists of ~1100 **linear** C,H, and O containing compounds from PNK¹

(1) J. B. Pedley, R. D. Naylor, S. P. Kirby "Thermochemical Data of Organic Compounds" 2nd ed. 1986

- PNK is a core dataset for fitting Benson groups
- ~600 PNK compounds are small enough for G4 calculations and comparison with experiment.



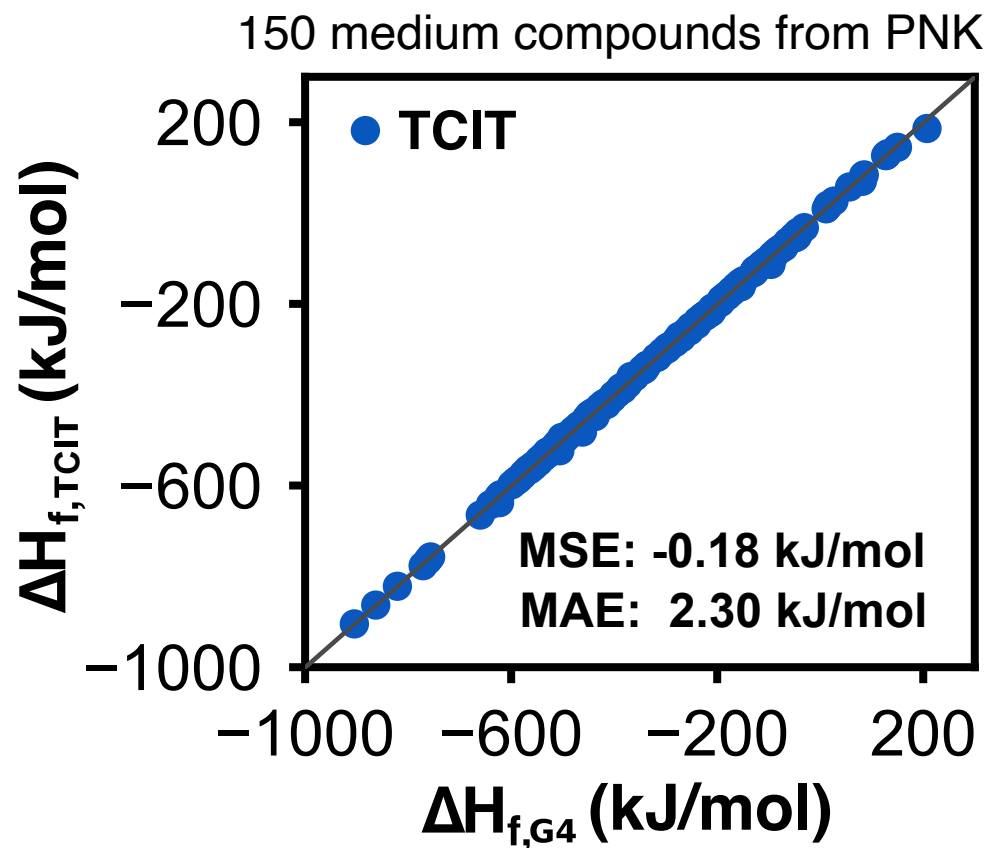
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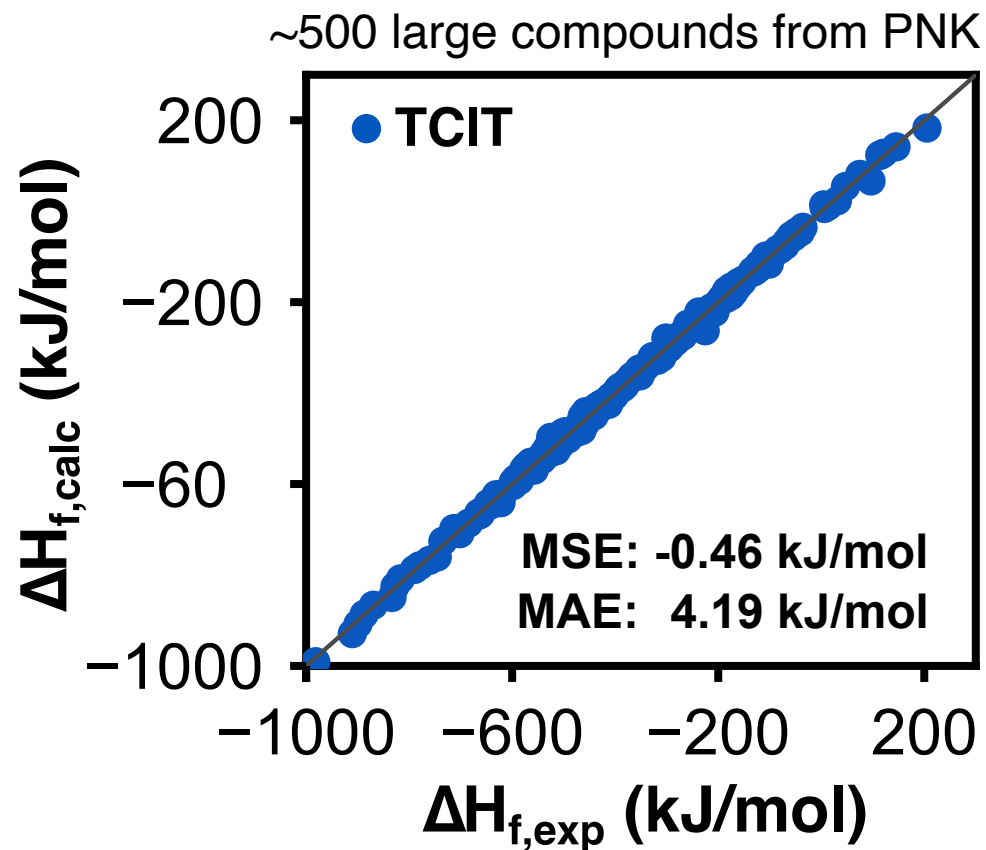
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- ~500 PNK compounds are large enough to evaluate the predictive accuracy of the increment theories.



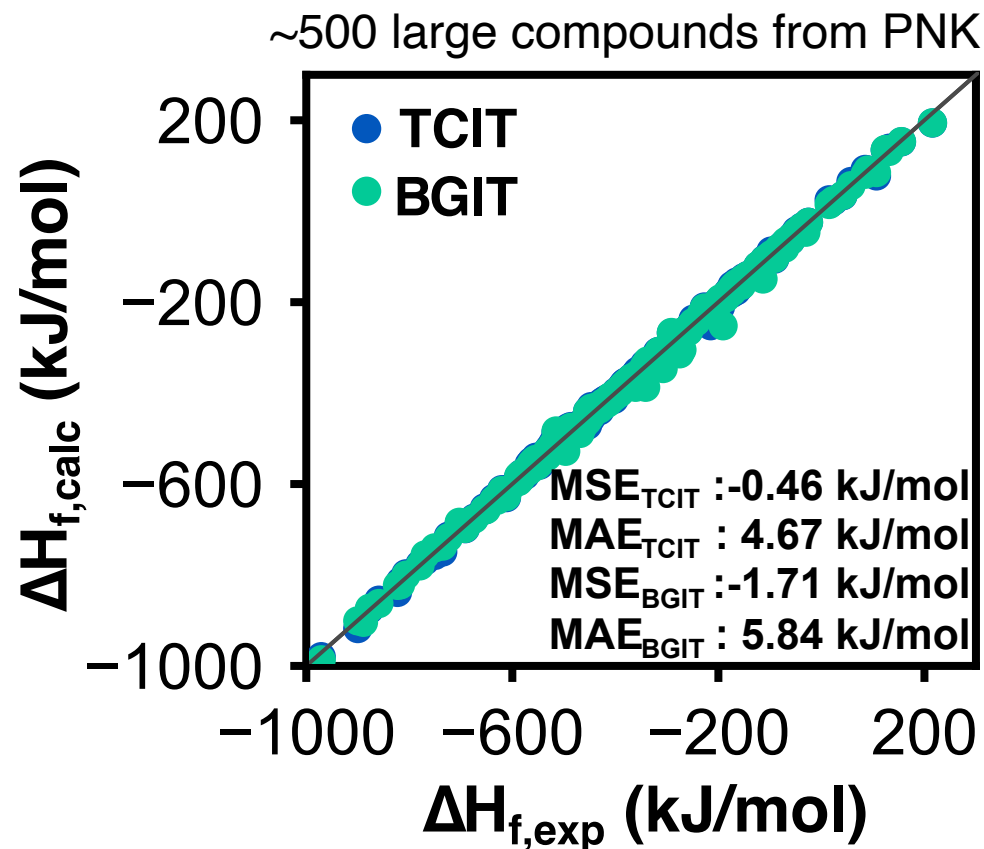
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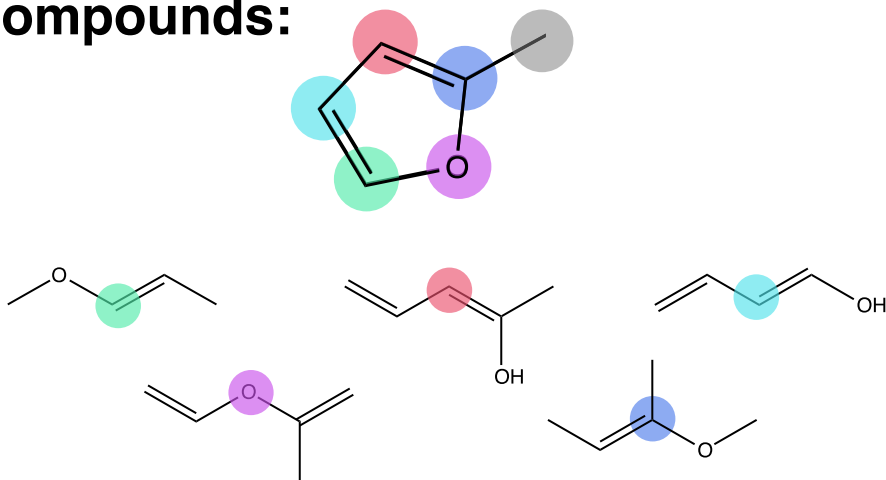
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TCIT shows comparable performance to BGIT/CHETAH but is derived exclusively from extensible G4 data.

Extension to Ring-Containing Molecules

- Ring-containing molecules have additional strain and/or conjugation corrections that exacerbate the extensibility issues of Benson Theory.
- In TCIT we are addressing this through chemically specific ring corrections that account for differences in substitution pattern and topology:

1. Decompose ring into acyclic model compounds:



2. Add ring correction (RC) to final prediction:

$$RC = H_f(\text{ring}) - H_f(\text{red}) - H_f(\text{cyan}) - H_f(\text{green}) - H_f(\text{purple}) - H_f(\text{blue}) - H_f(\text{grey})$$

Technical Developments

RC Model Compounds

○ : Depth 0 ○ : Depth 1 ○ : Depth 2

Method 1: Use RC₁ based model parameterized to G4 data.

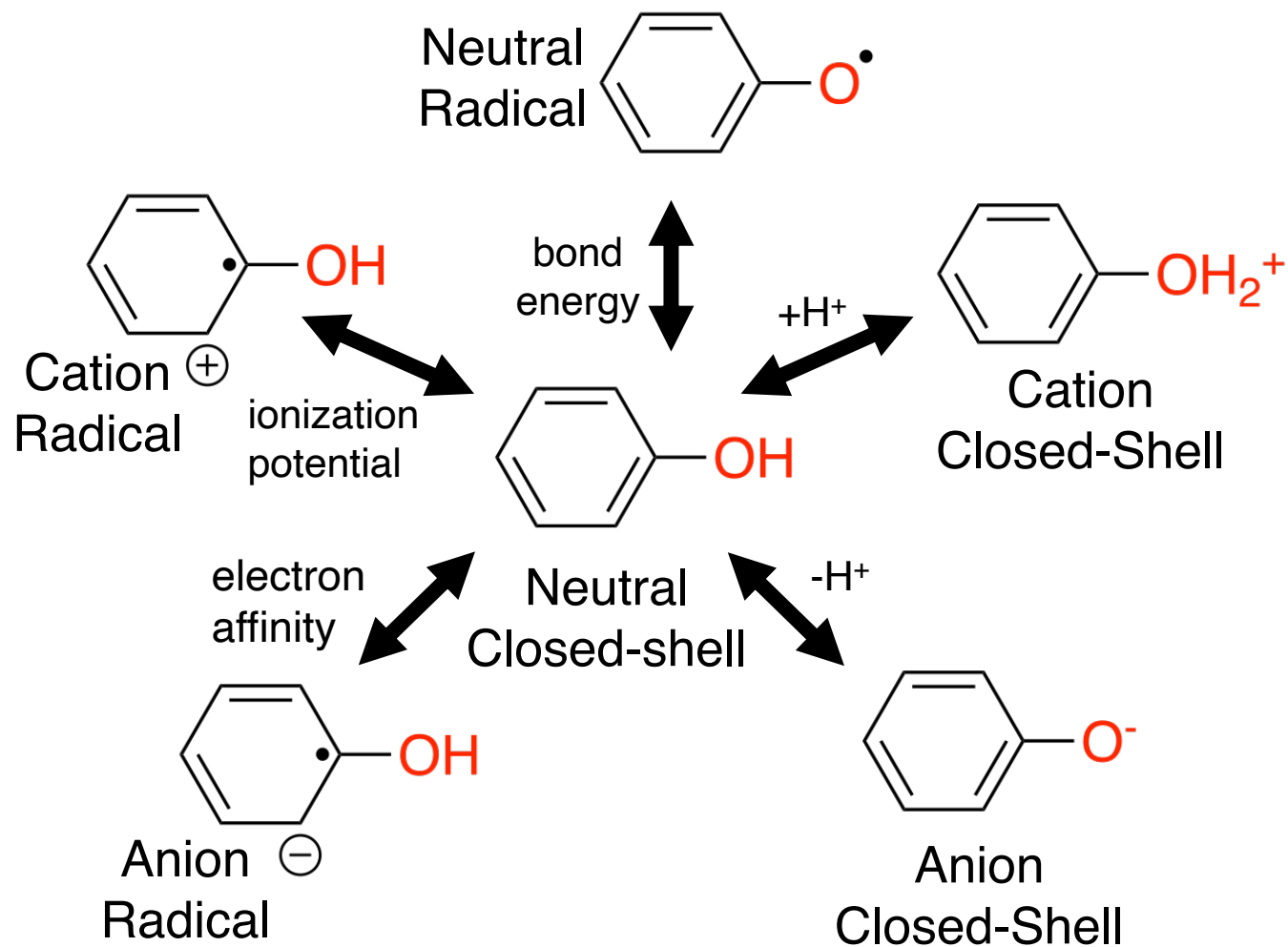
Method 2: Use graph-NN to predict RC₀-RC₂

Extending TCIT to Radicals and Ions

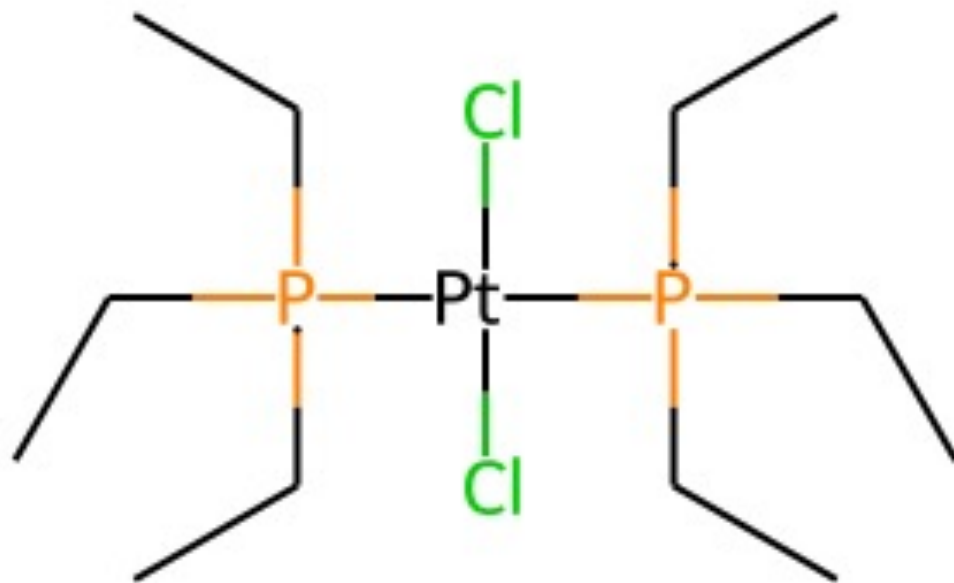
A recurring question is when will TCIT support predictions on **radicals** and **ions**?

TCIT already covers neutral close-shell species, so these extensions require us only to predict the difference between the target and the **nearest closed-shell neutral**.

This amounts to developing models to predict IP/EA/+H⁺/-H⁺



Extending TCIT to Organometallics



High level calculations on transition metals are going to get expensive, so we want to avoid calculating components directly

$$\Delta H_f^\circ = \sum_i C_{\text{TCIT},i} + \sum_{i \in M-X} \text{BDE}_i$$

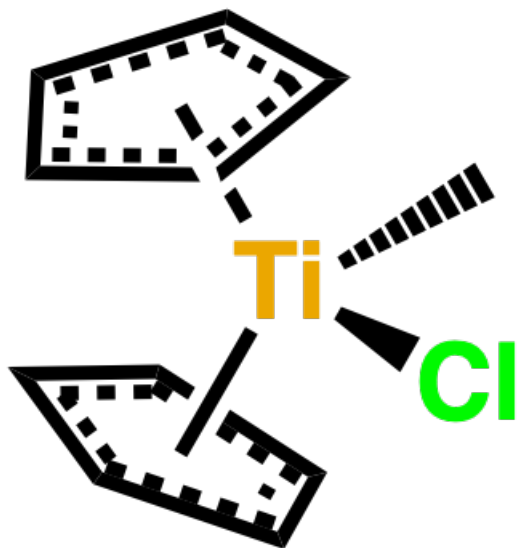
Idea: Only use TCIT for the ligands and approximate the remainder with bond energies

Species	ΔH_f Contribution (kcal/mol)	Source
PtCl ₂ (g)	26.41	Schafer 1975
Pt-P 1 (B3LYP)	-39.91	Craciun 2010
Pt-P 2 (B3LYP)	-34.81	Craciun 2010
PEt ₃ (TCIT)	-35.21 * 2	
Predicted	-118.73	
<i>trans</i> -Pt(PEt ₃) ₂ Cl ₂	-118.3 ± 1.9	Takhin 1984

Sanity checks work surprisingly well

(Note: PtCl₂ availability probably makes it better than it ought to be)

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Species	ΔH_f (kcal/mol)	Source
Cp ⁻ (TCIT/G4)	19.9 (x2 for compound)	TCIT/G4
Ti-Cl (TiCl ₄ /4)	-43.6	Calhorda 1986
Ti-Cp (Ti(Cp) ₂ Cl ₂)	-7.9 (x2 for compound)	Calhorda 1986
CH ₃ ⁻ (TCIT/G4)	32.41	TCIT/G4
Ti-CH ₃ (Ti(Cp) ₂ (CH ₃) ₂)	-38.6	Calhorda 1986
Predicted	-25.8	
Ti(Cp) ₂ (Cl)(CH ₃)	-29.8 +/- 3.0	Calhorda 1986

Extending TCIT to Organometallics

TCIT now contains all CAVs necessary to predict ΔH_f of all N, H, O, and C-containing molecules in pubchem. **This is the largest repository of G4 calculations on large molecules in the world.**

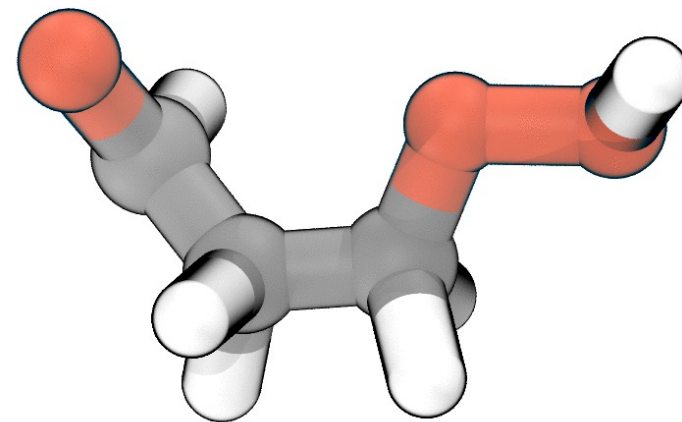
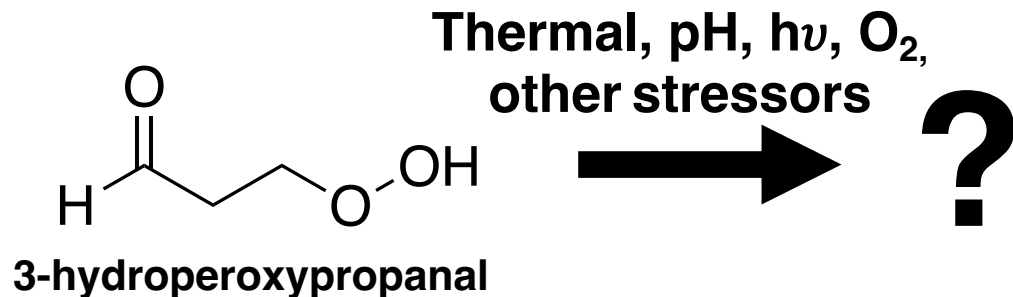
It is foreseeable that we could complete all B, F, Cl, S, and P containing structures over the next few years.

Species	ΔH_f (kcal/mol)	Source
Cp ⁻ (Ti)		
Ti-Cl (Cp)		
Ti-Cp		
CH ₃ ⁻ (Ti)		
Ti-CH ₃		
Predicted	-25.8	
Ti(Cp) ₂ (Cl)(CH ₃)	-29.8 +/- 3.0	Calhorda 1986

The Reaction Prediction Problem

A → B : When we know the reactants and products, mature quantum chemistry tools exist to characterize transition states and establish pathways

A → ? : For degradation reactions, plausible reactions are often unknown.

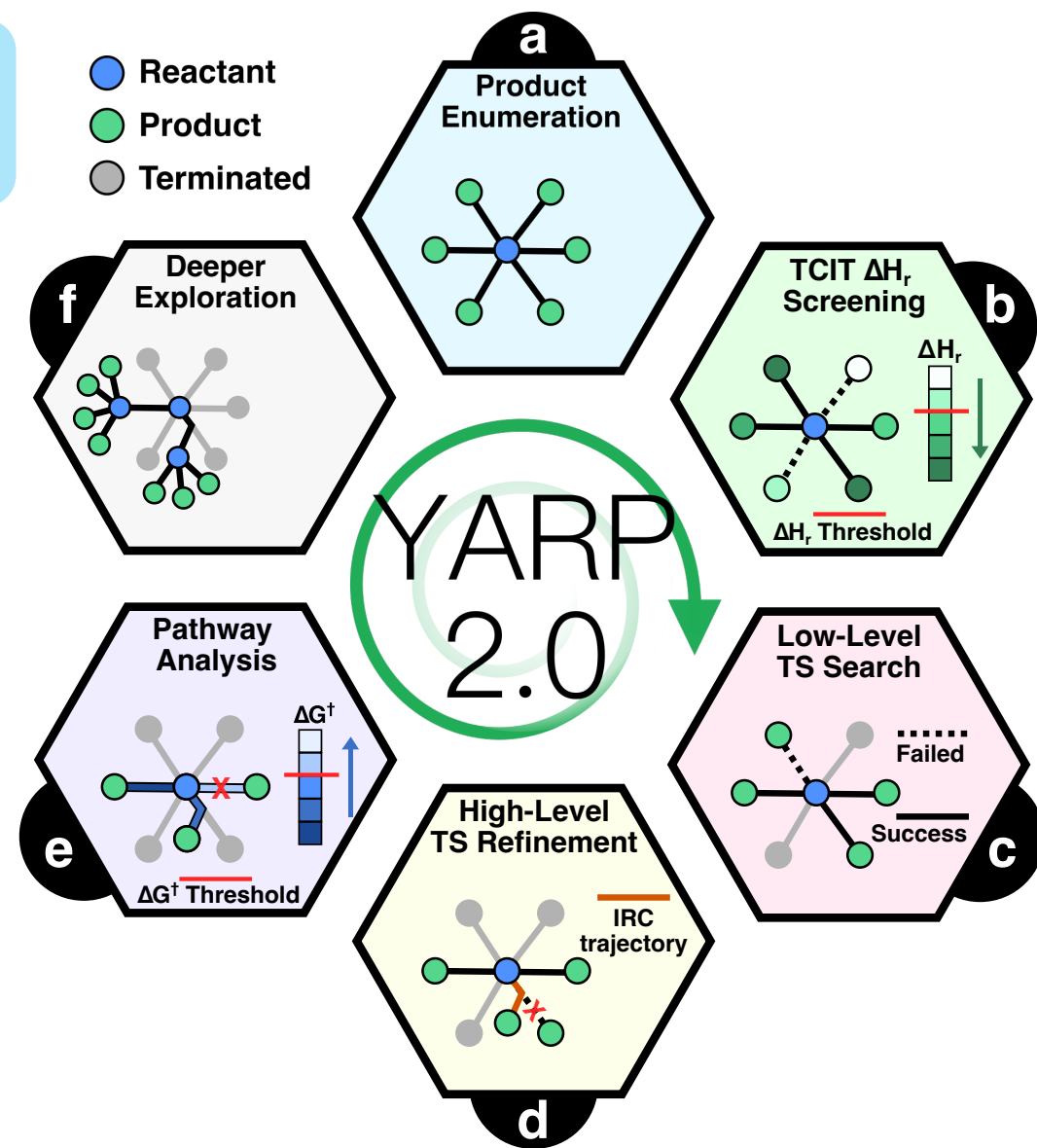


Yet Another Reaction Program (YARP)

Idea: Turn the $A \rightarrow ?$ problem into tractable (and parallelizable) $A \rightarrow B$ problems.

Observations:

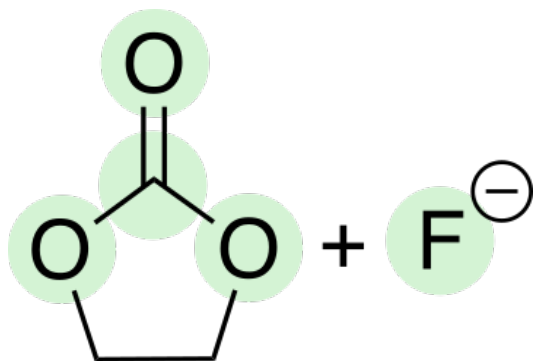
- Product enumeration is easier than transition state enumeration.
- Transition state algorithms for $A \rightarrow B$ problems are mature. Let the TS algorithm identify physical reactions.
- Recent developments in semi-empirical models and ML create opportunities.
- Solving the $A \rightarrow ?$ problem is the prerequisite for reaction network prediction.



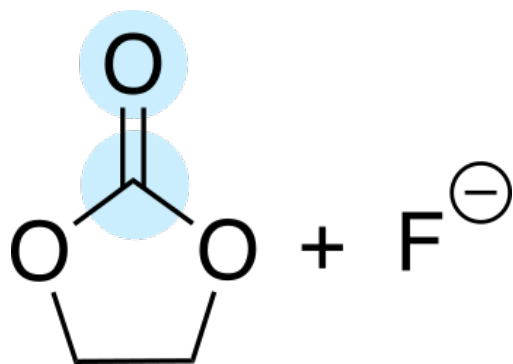
YARP: Elementary Reaction Step(s)

Polar and pericyclic organic reactions are decomposed into elementary electron donor and acceptor reactions with concomitant σ -bond breaks

bnfn
will refer to
 σ -bond
changes,
 π -bonds are
allowed
to arbitrarily
rearrange.

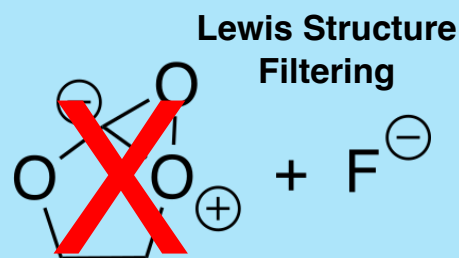
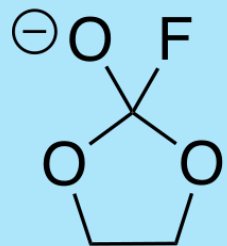
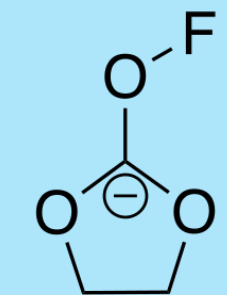


Lone-Pair Donors

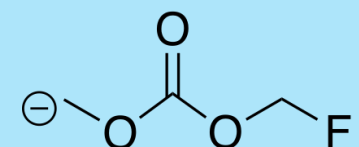
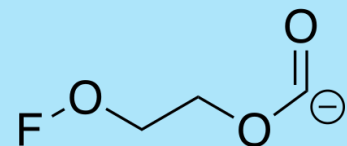
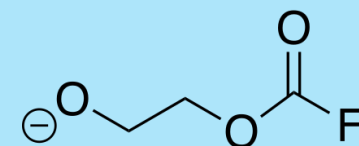
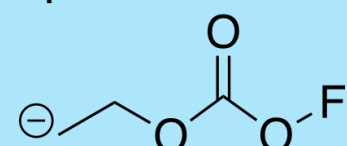
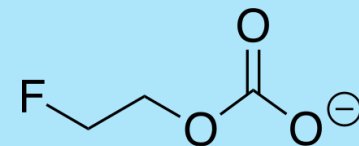
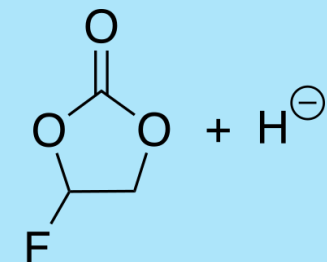
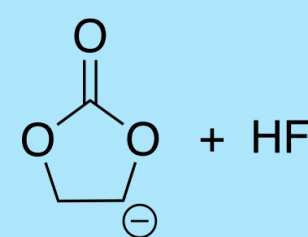


Lone-Pair Acceptors

Form 1 Products



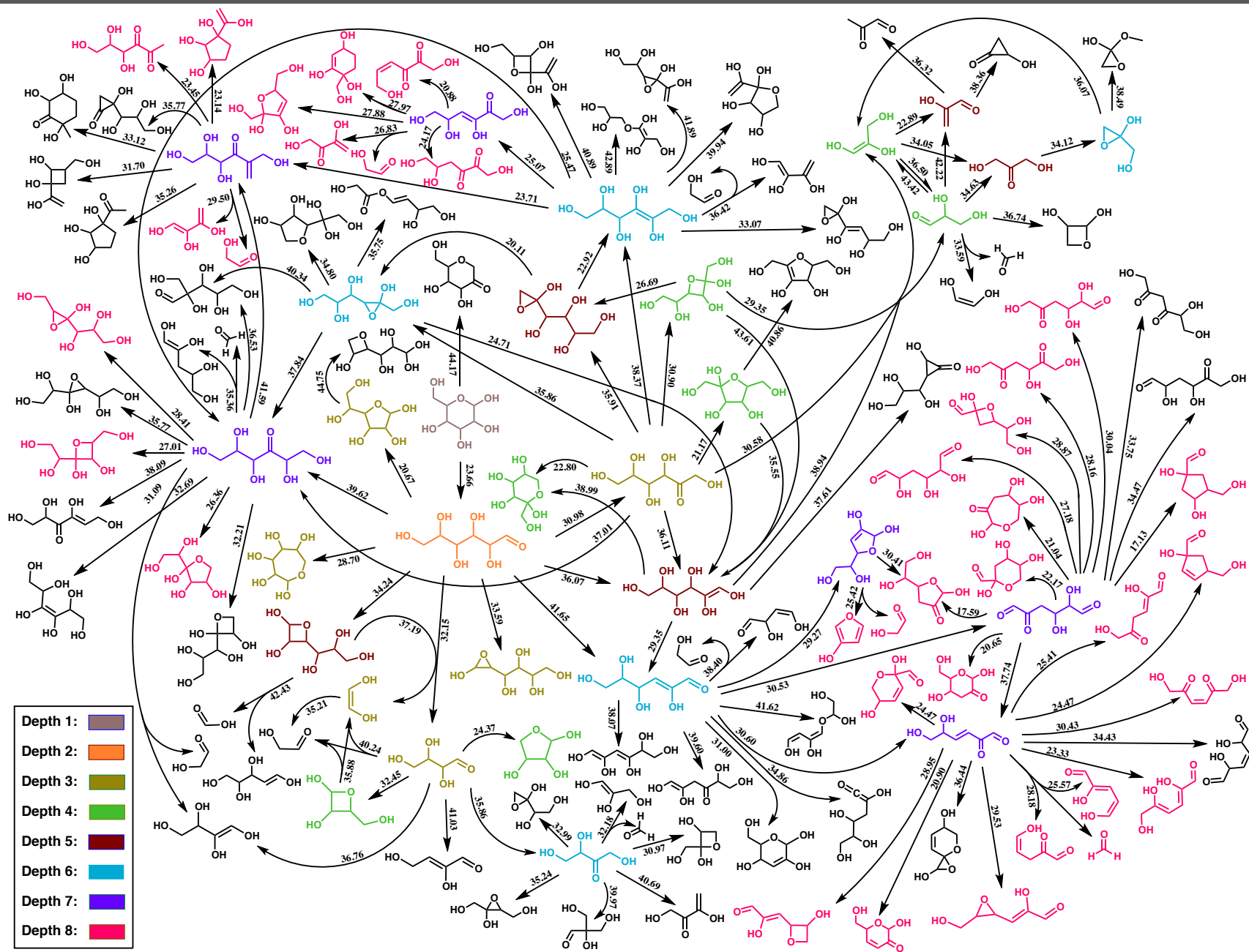
Break 1 Form 1 Products



+ 28 others

β -D-Glucose Pyrolysis Network Exploration

Zhao, Q.; Savoie, B. M. Proc. Nat. Acad. Sci. 120, 2023



To perform a deep network exploration, we've implemented a modified version of Dijkstra's algorithm

At each iteration:

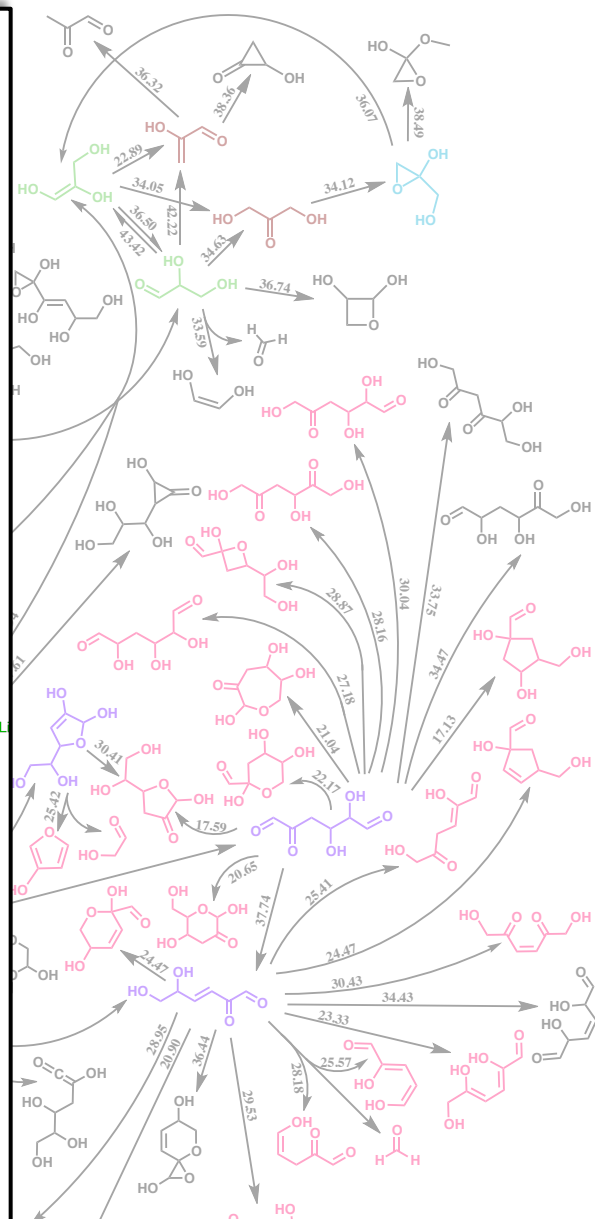
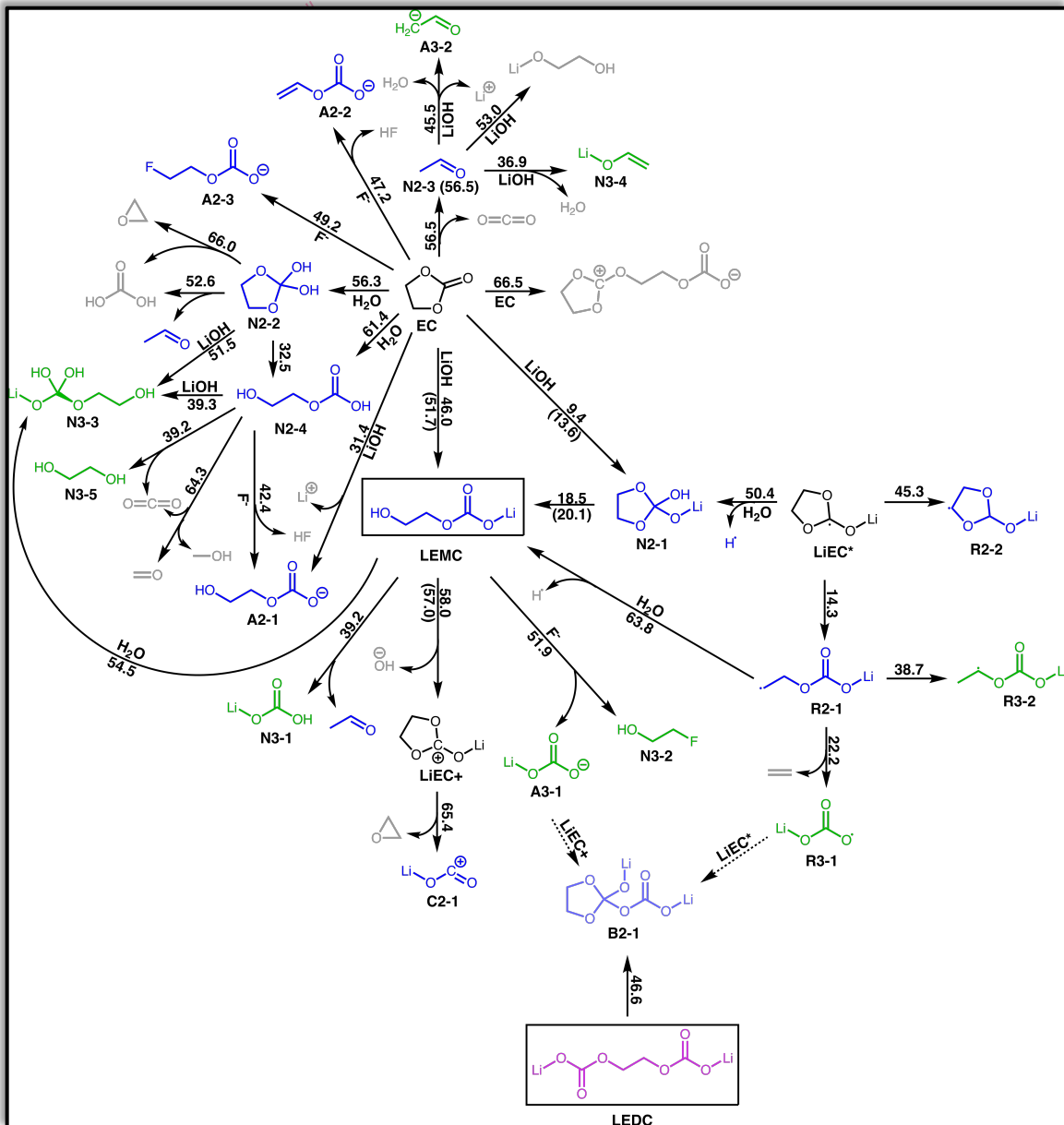
(1) all b2f2 reactions are explored for active nodes.

(2) Active nodes are determined by the minimum barrier to a given product (with a window)

(3) Water catalyzed reactions are considered for all H-transfers

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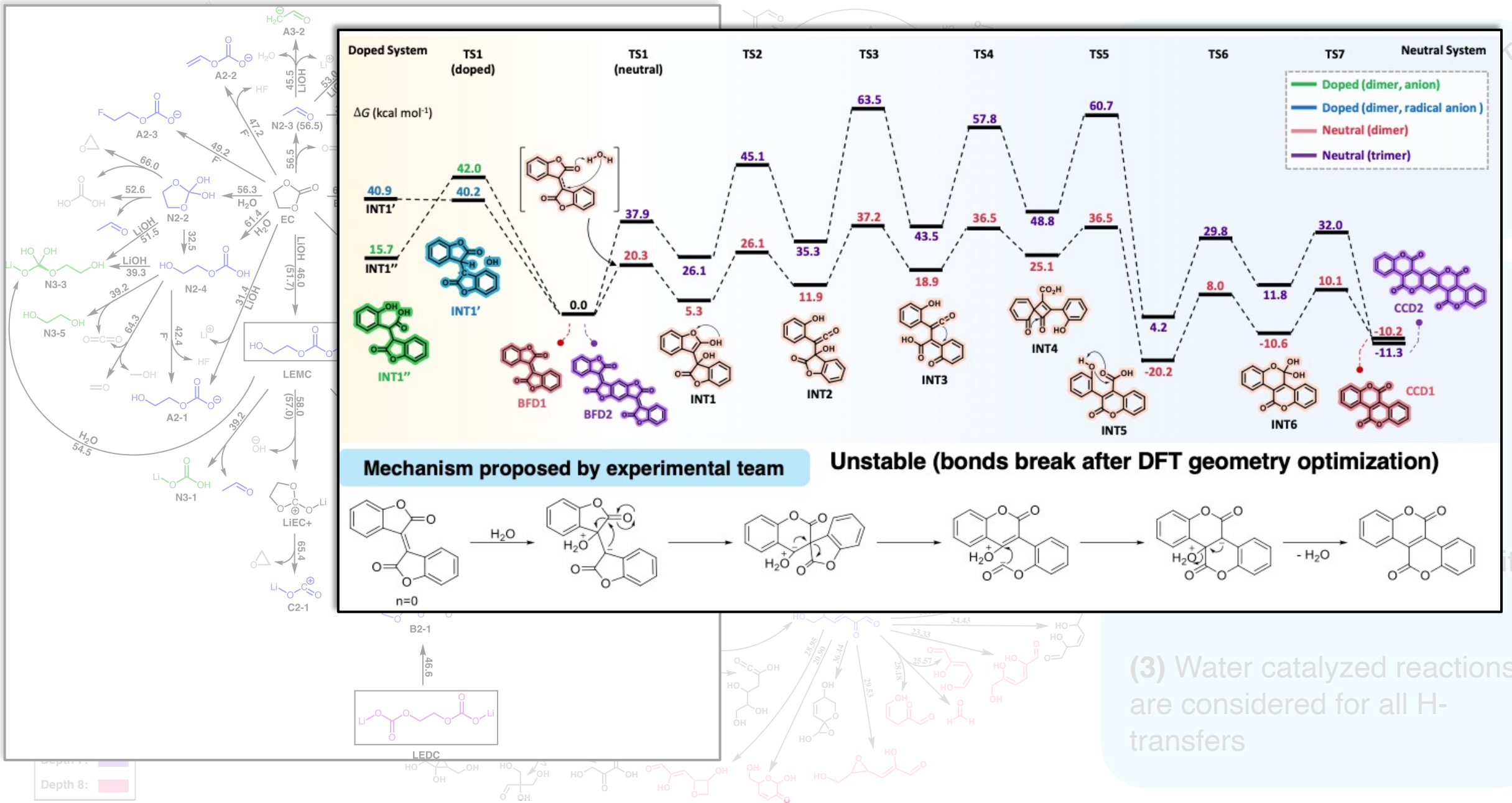


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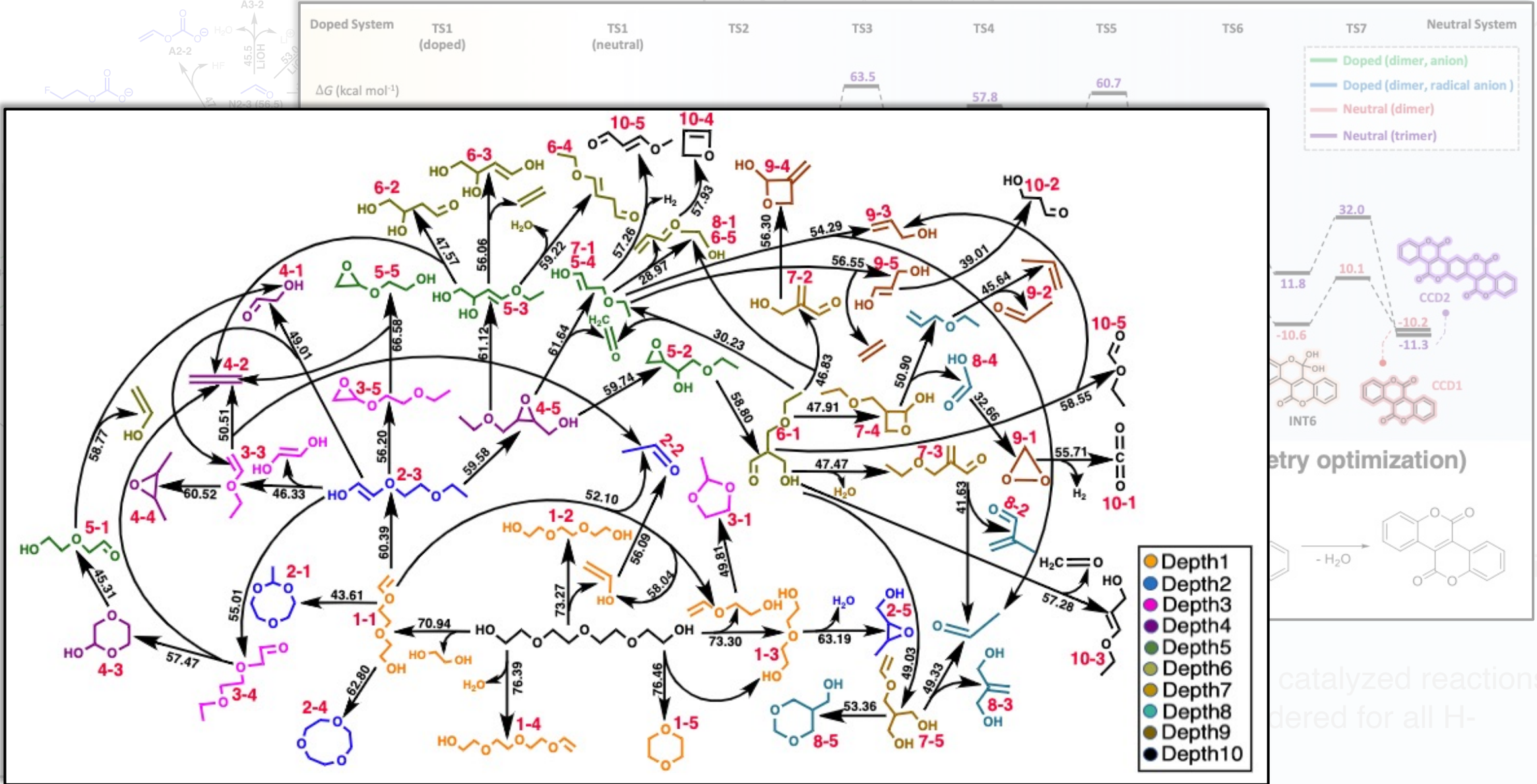
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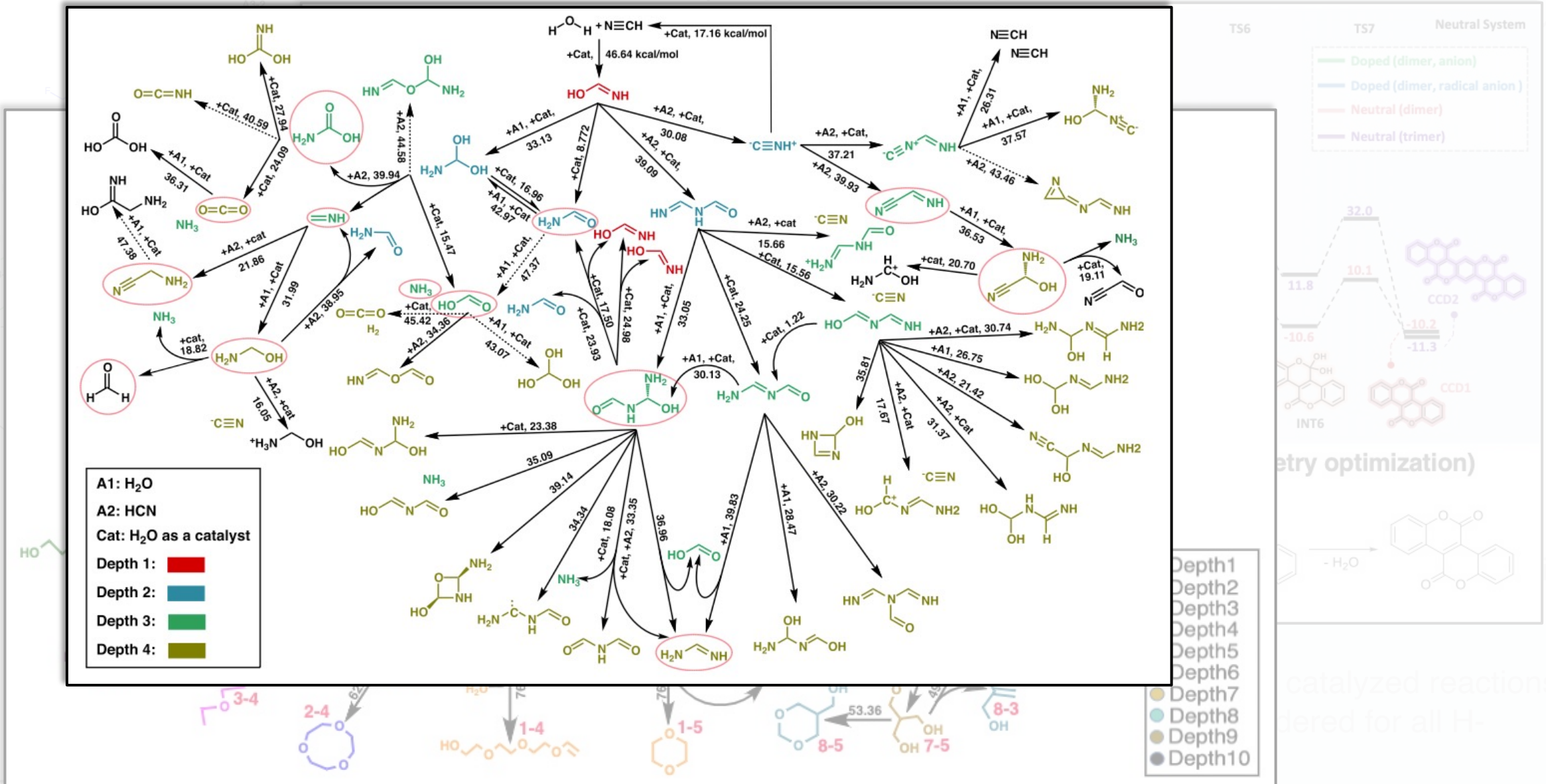
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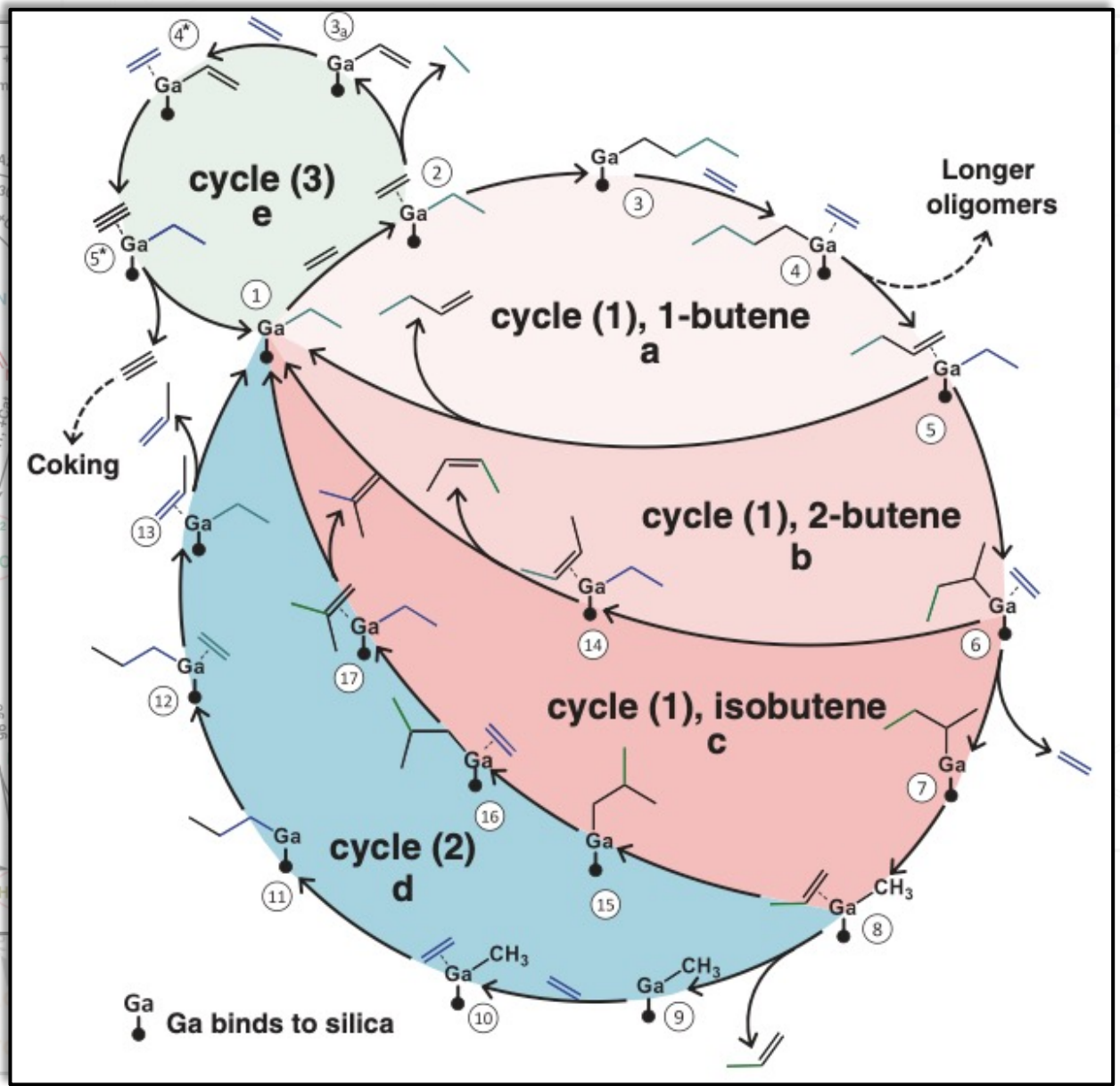
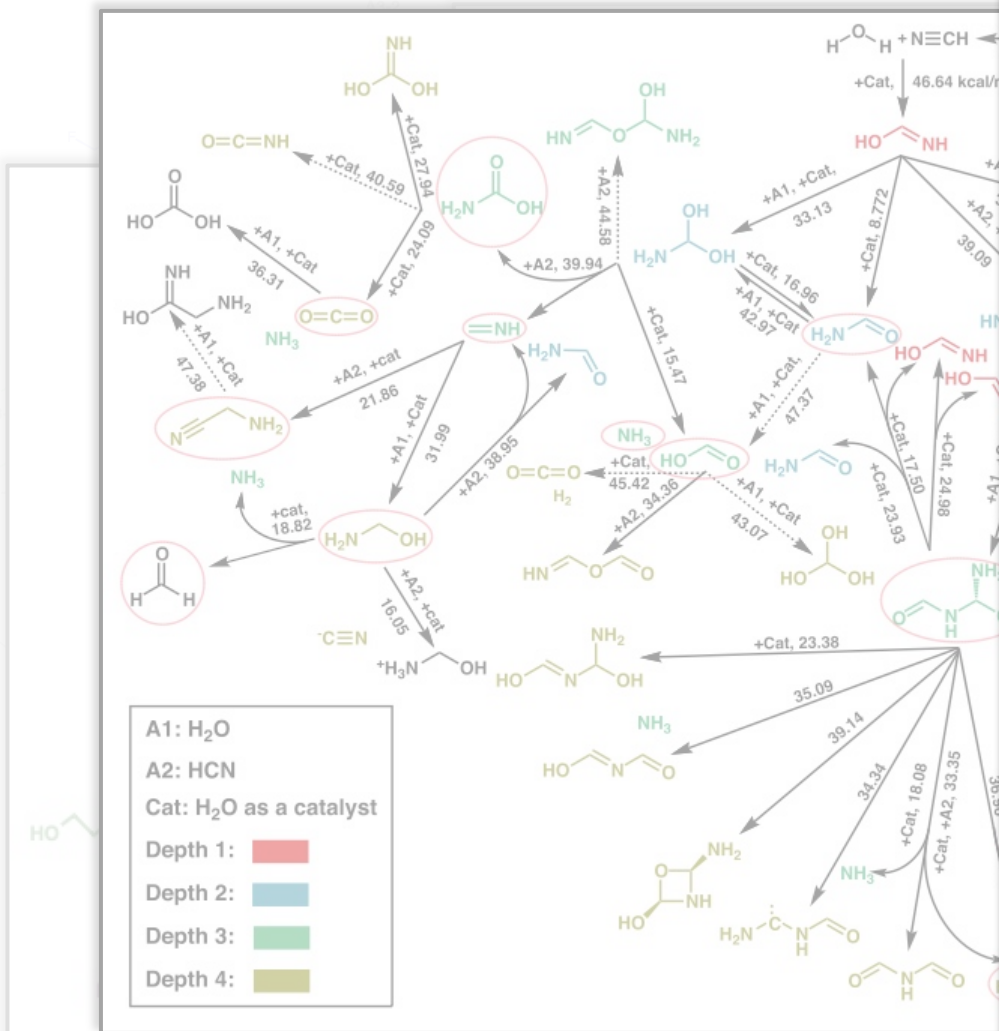
Depth 8: [pink box]

β -D-Glucose Pyrolysis Network Exploration



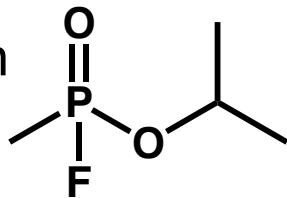
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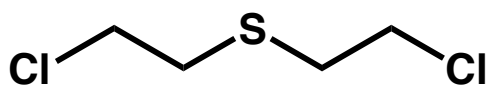


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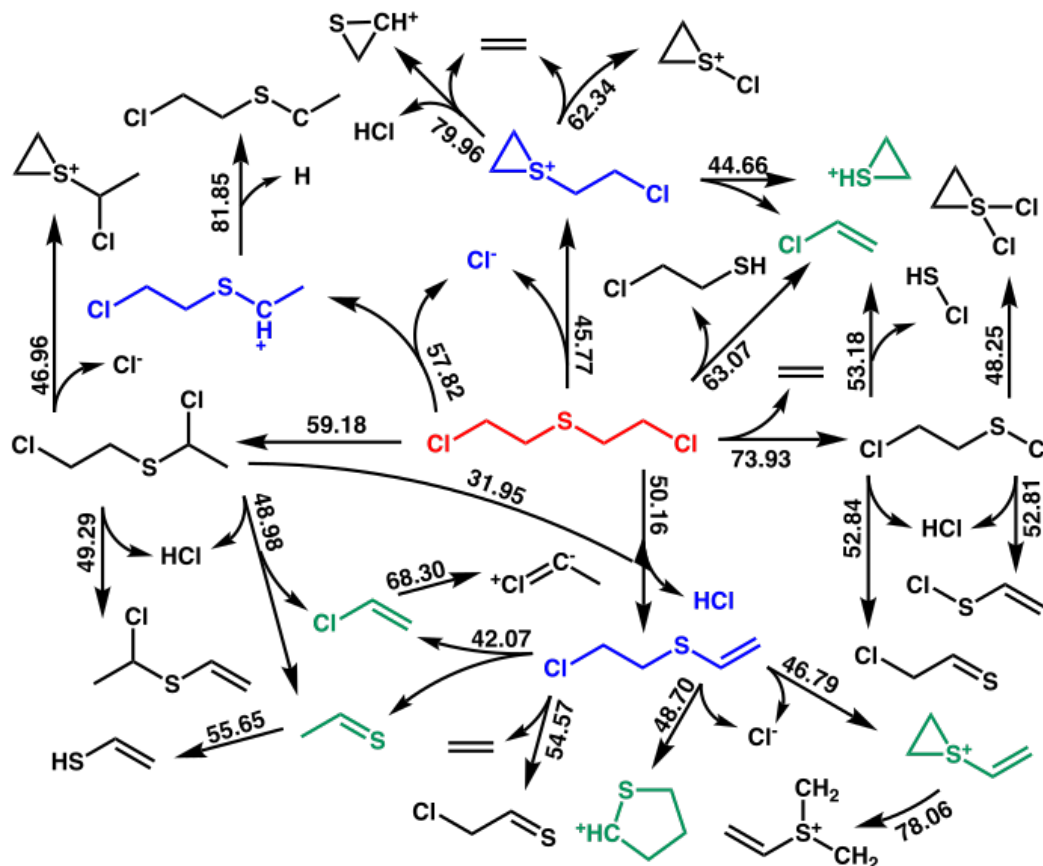
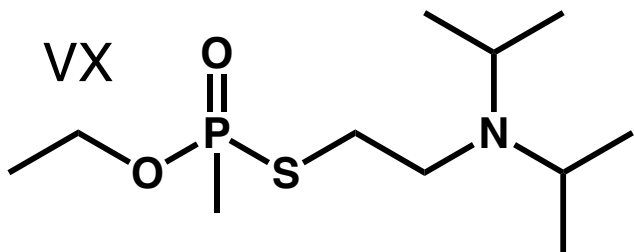
Sarin



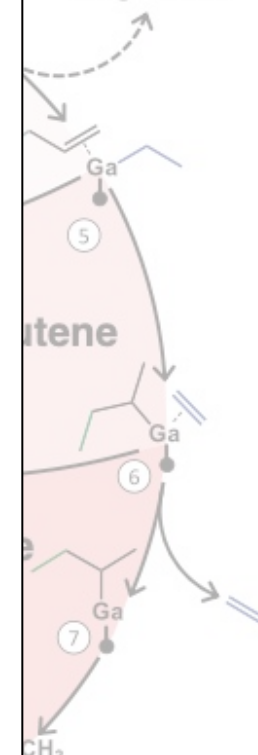
Mustard Gas (HB)



VX



Longer oligomers



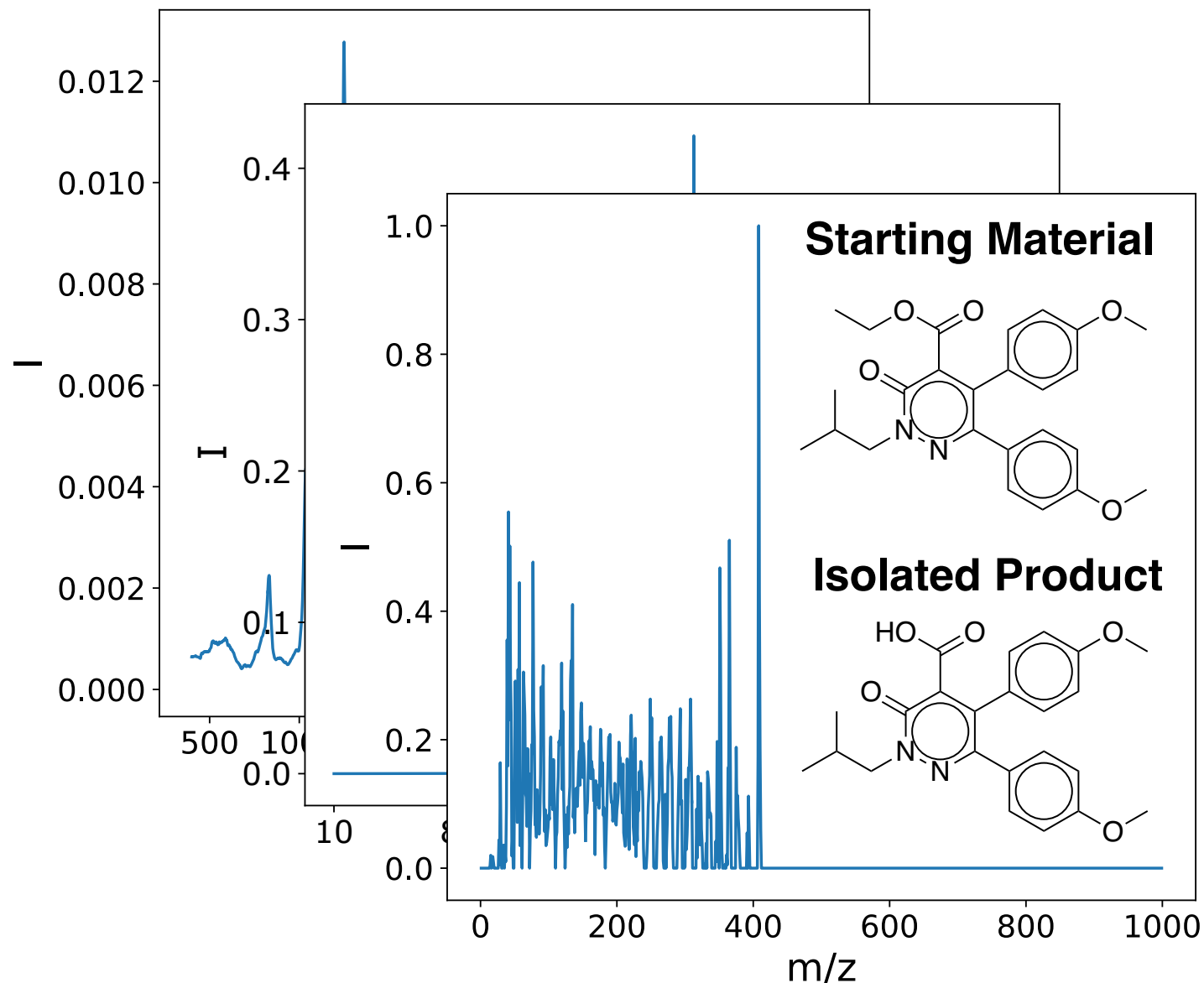
Ga binds to silica

(Pop Quiz) The Product Identification Problem

Suppose we've isolated an unknown degradant and I start offering you spectra.

How long until you can unequivocally identify the structure?

What if I told you the starting material?

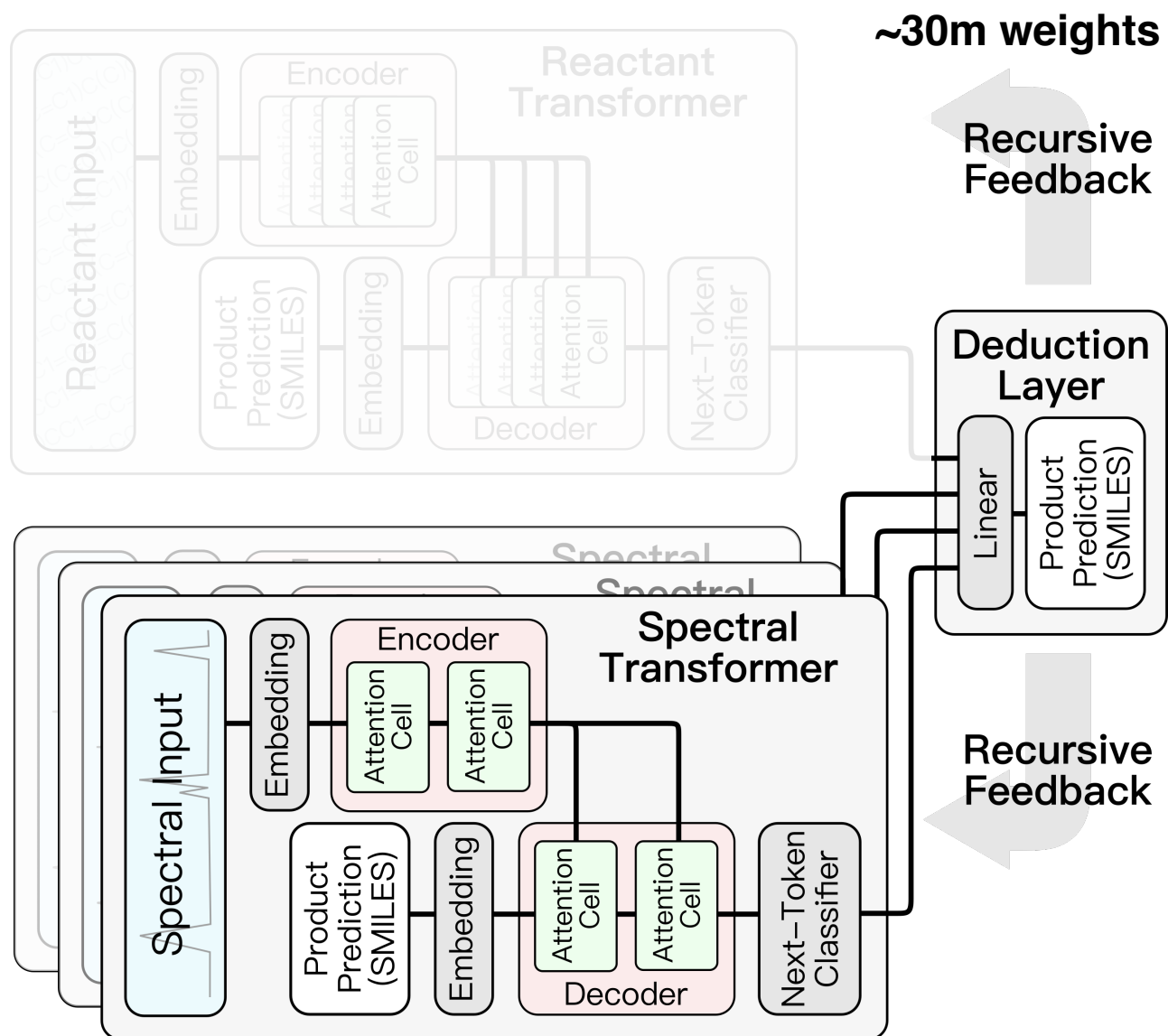


Leveraging Mixture of Experts Transformer Architectures

We developed this architecture to emulate the deductive process that experts use during product identification

The **product identification** architecture, corresponding to the full network, accepts information about the **reactants** and **spectral information of products** to predict the corresponding product identity.

The **spectrum to structure (StS)** identification architecture corresponds to just the bottom half. This architecture doesn't use any knowledge of reactants.



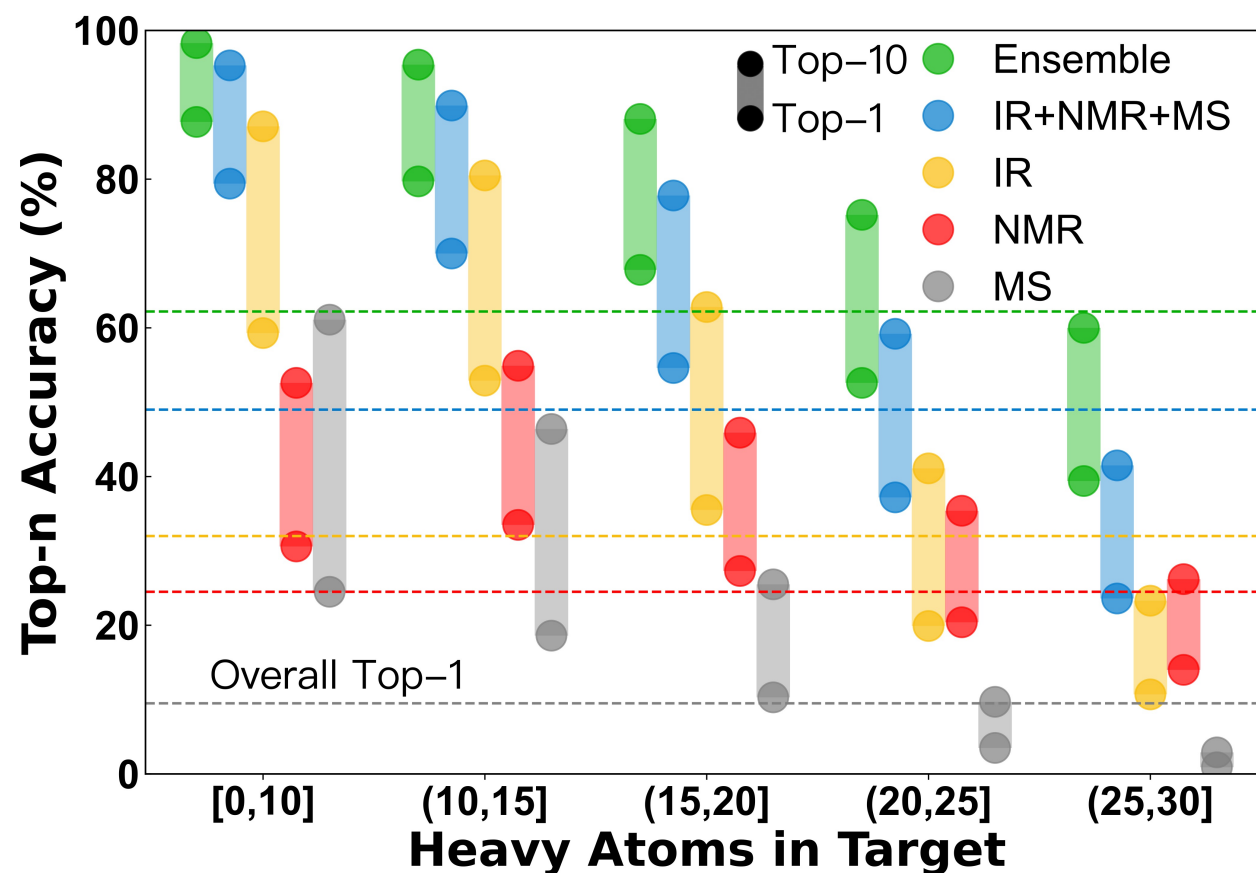
Is there sufficient information for the StS task?

1H-NMR, IR, and EI-MS spectra were simulated for 305,493 USPTO species and 524,860 from PubChem.

Sufficient information is probable for small molecules:
~85% top-1

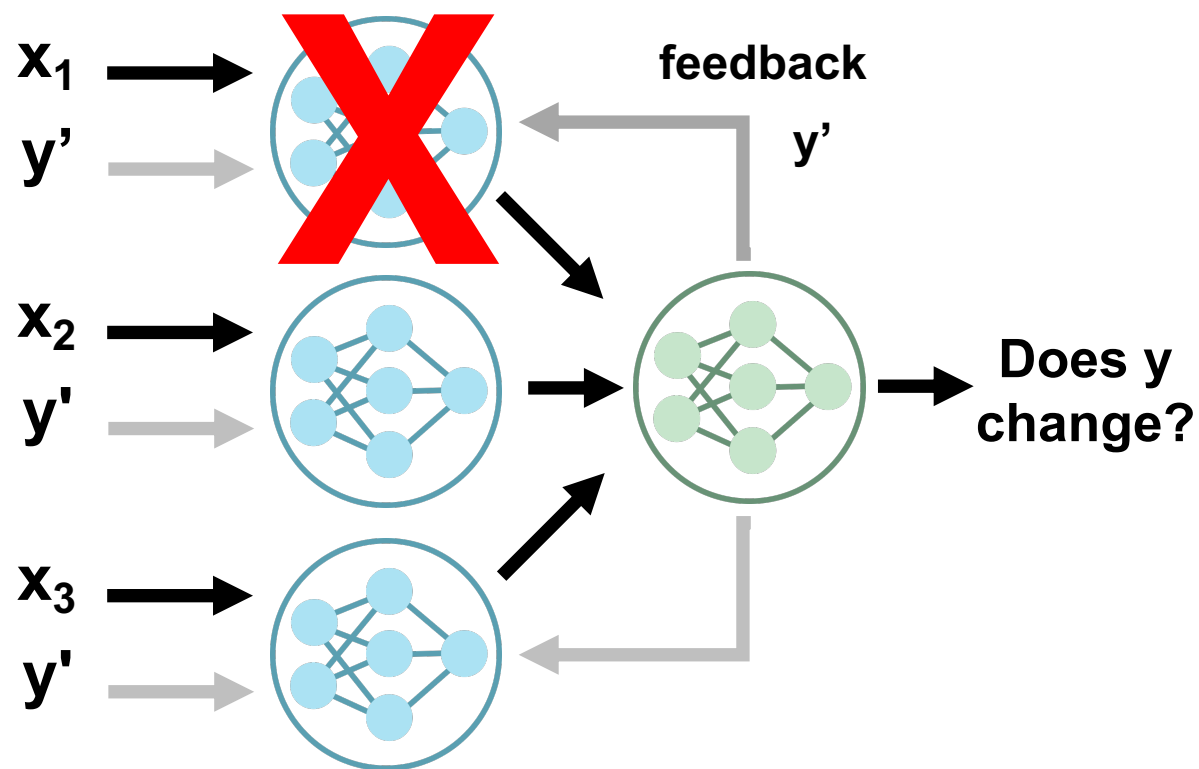
Potential deductive bottleneck for large molecules:
~38% top-1

Top-10 predictions obtained with beam-search. Performance is shown on a random 10% testing split.



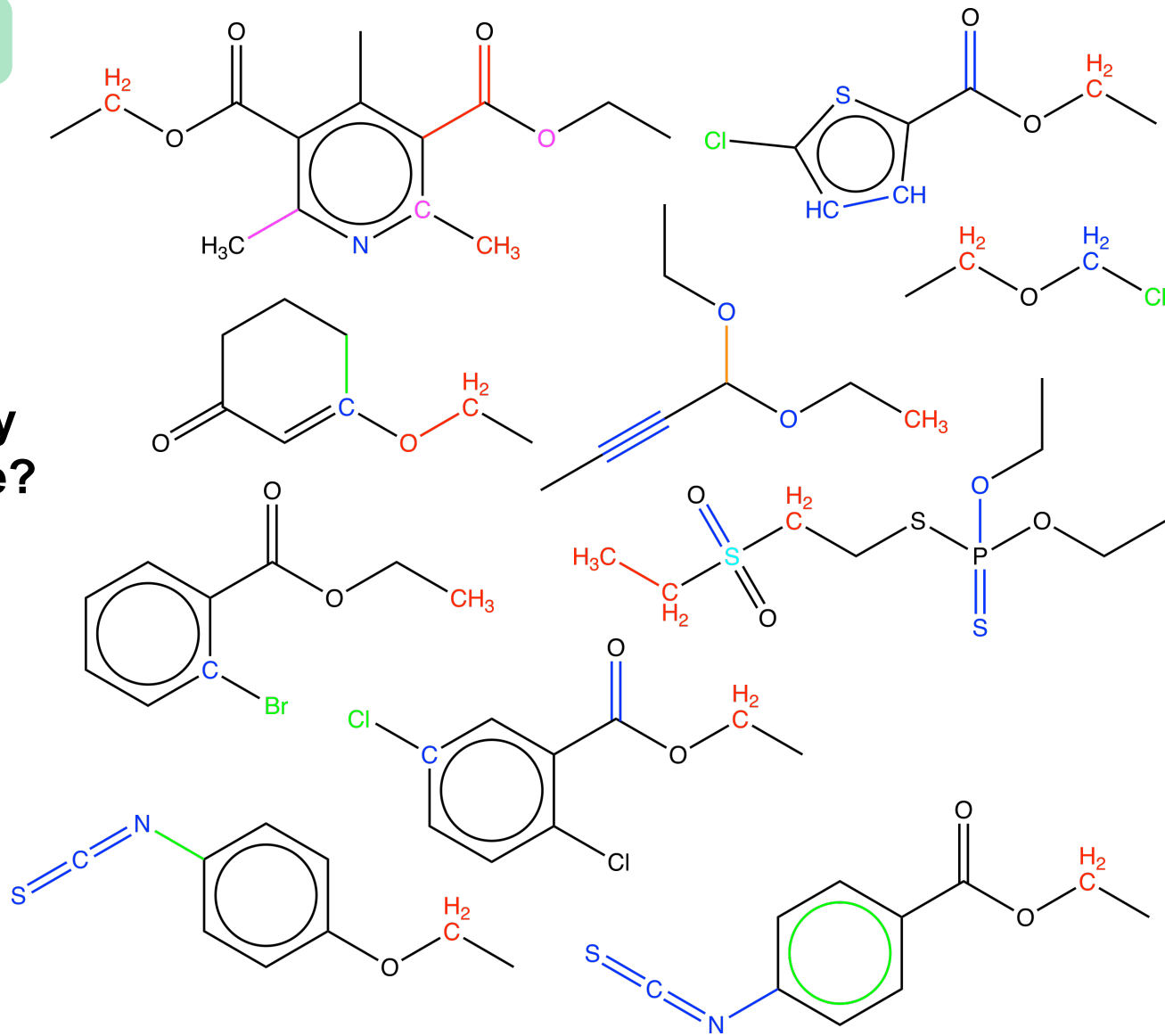
Direct Evidence of Dynamic Deduction by these Models

Visualizing Per Token Spectral Decisiveness



Color Legend:

- $^1\text{H-NMR}$ (Red)
- IR (Blue)
- EI-MS (Green)

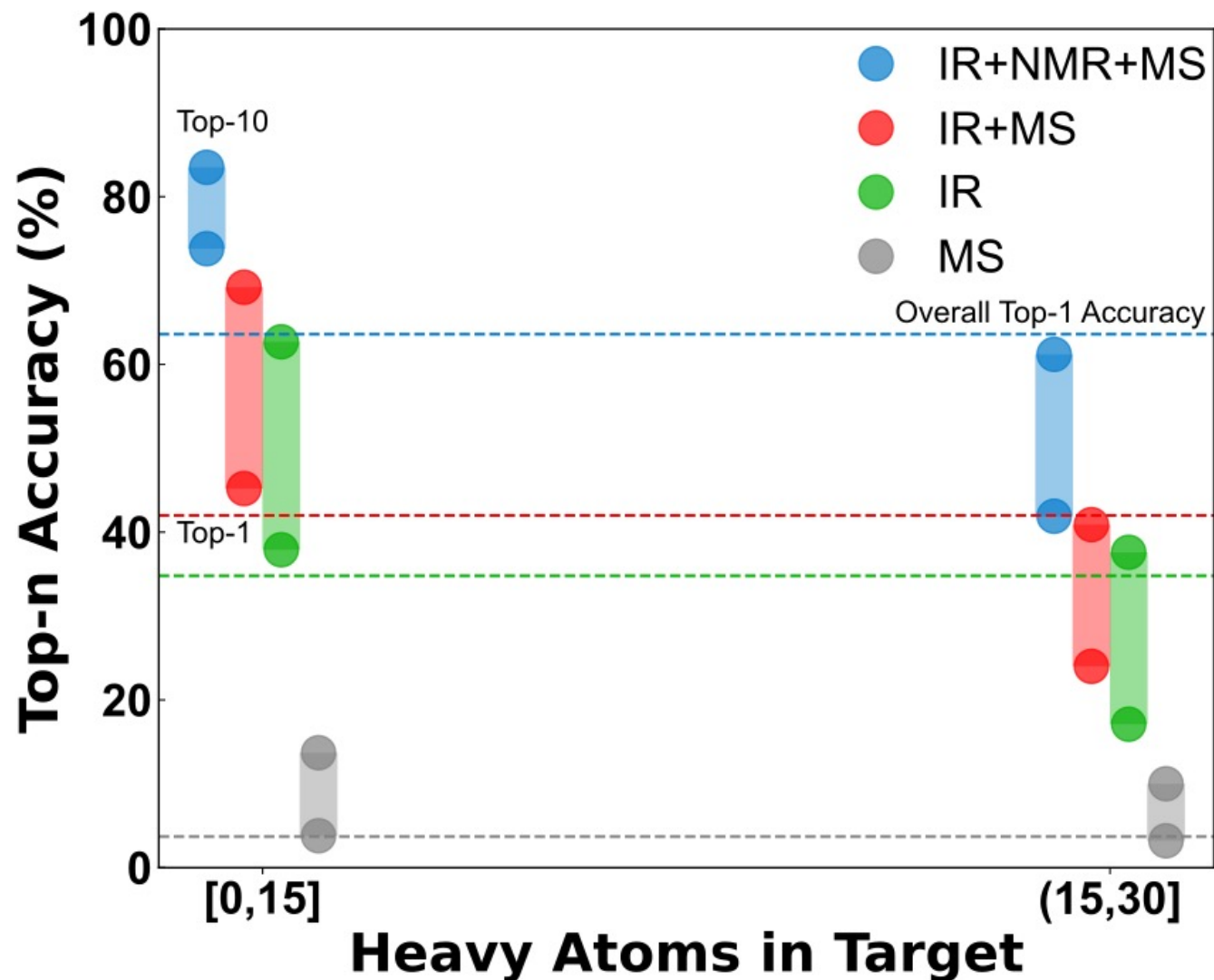


Does the StS Model Transfer to Real Spectra?

We used 5544 molecules from NIST with experimental IR/H-NMR/EI-MS* as an external test for the model

None of the NIST molecules were in the training data, and all training was on simulated not experimental spectra.

External NIST Testing Set

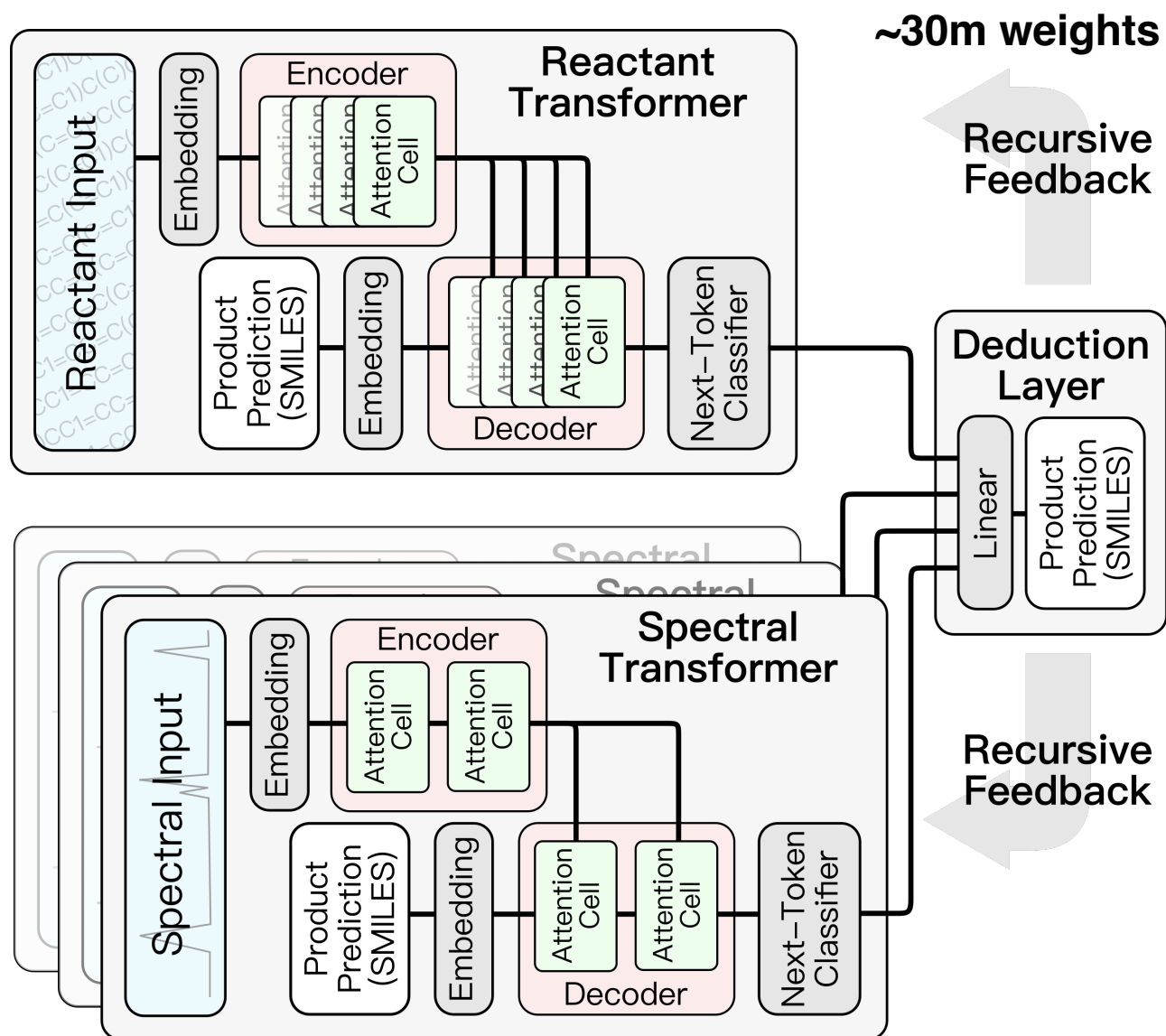


Leveraging Mixture of Experts Transformer Architectures

We developed this architecture to emulate the deductive process that experts use during product identification

The **product identification** architecture, corresponding to the full network, accepts information about the **reactants** and **spectral information of products** to predict the corresponding product identity.

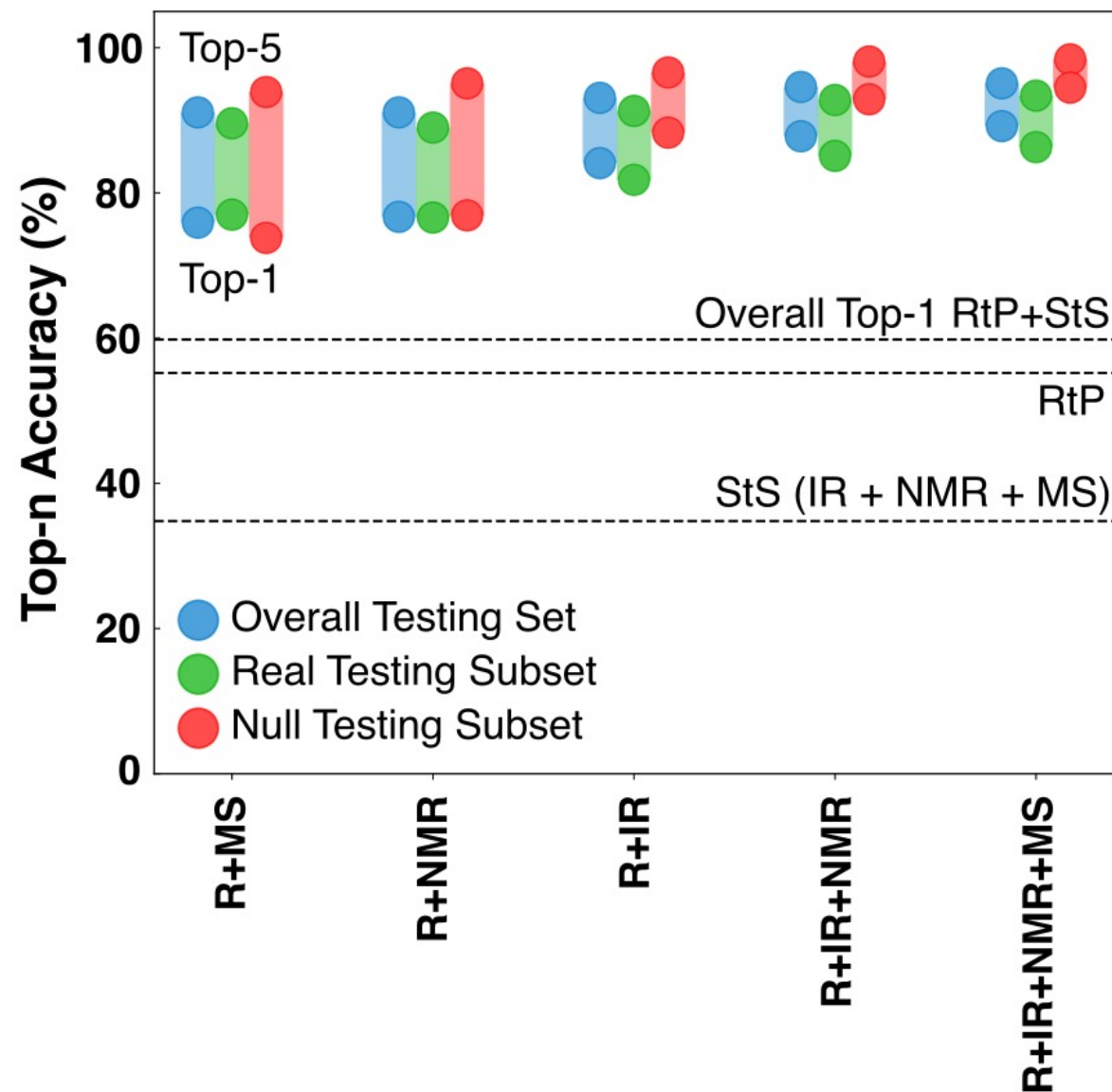
The **spectrum to structure (StS)** identification architecture corresponds to just the bottom half. This architecture doesn't use any knowledge of reactants.



Adding Reactant Info Qualitatively Improves Performance

Testing these Reactant + Spectra models on product identification:

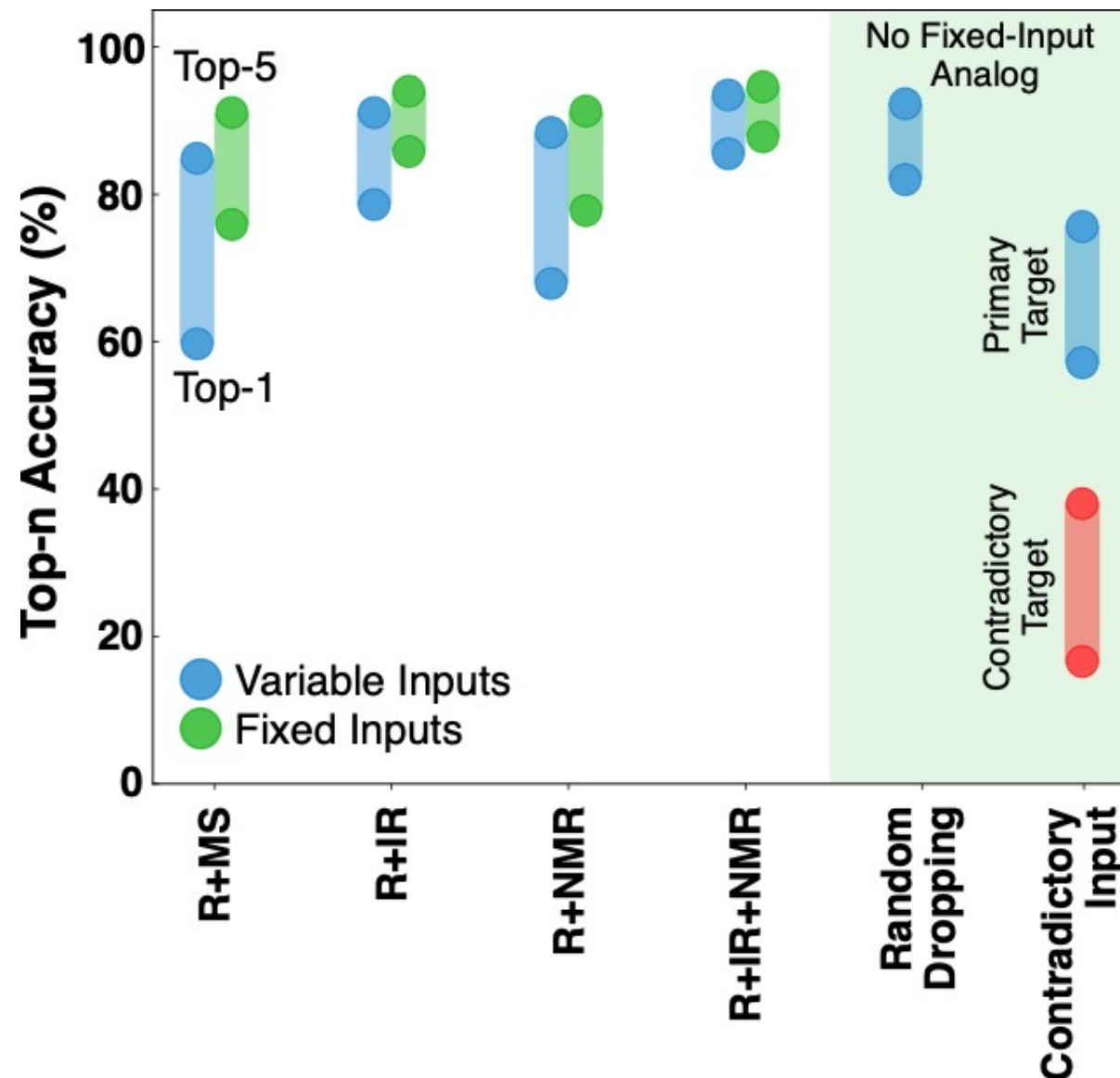
1. 466,330 Reactions from D. Lowe
2. 2:1 Split between real products and starting material
3. 80:10:10 Train:Val:Test
4. Simulated IR/1H-NMR/EI-MS for all 305,493 species



Can the Mixture of Expert Models Handle Contradictory Information?

A second aspect of deductive behavior is to reason with partial, or even, contradictory inputs.

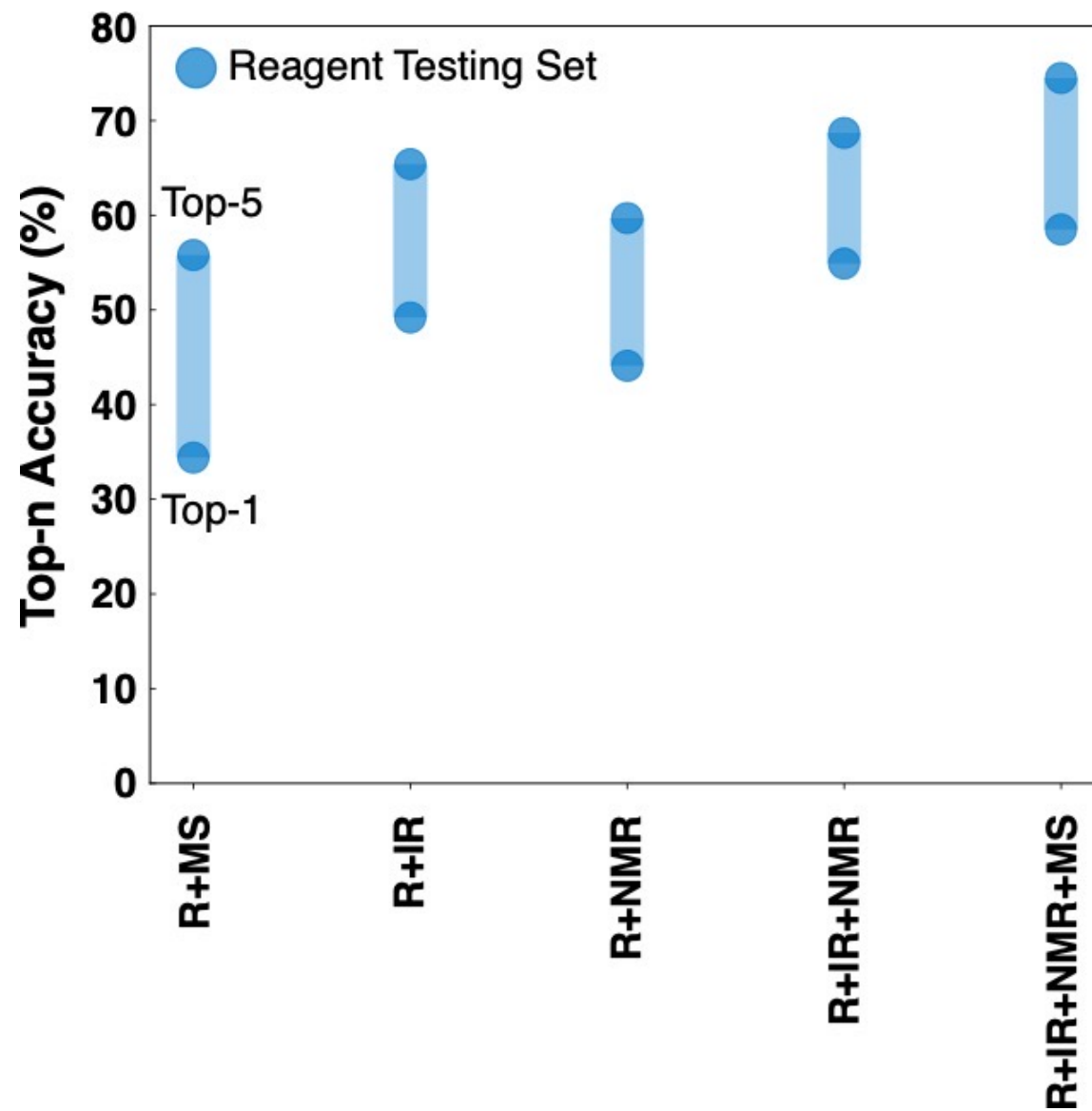
To test the performance under contradictory information we supplied a contradictory spectrum to one transformer (at random)



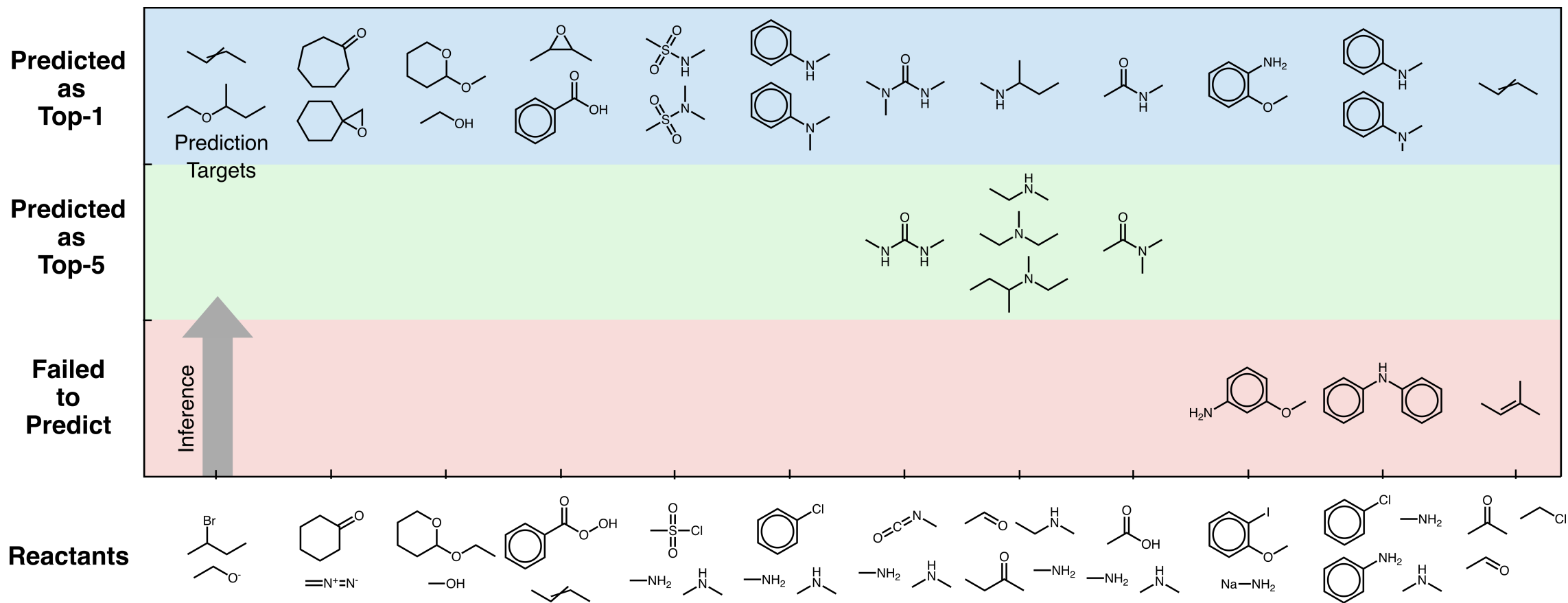
Can the Mixture of Expert Models Perform Out-of-Distribution?

Reagent identification is an untrained task that has the same inputs as product/null characterization

We curated ~1k reagents involved in ~3k reactions from the USPTO dataset. None of these species were seen as prediction targets during training.



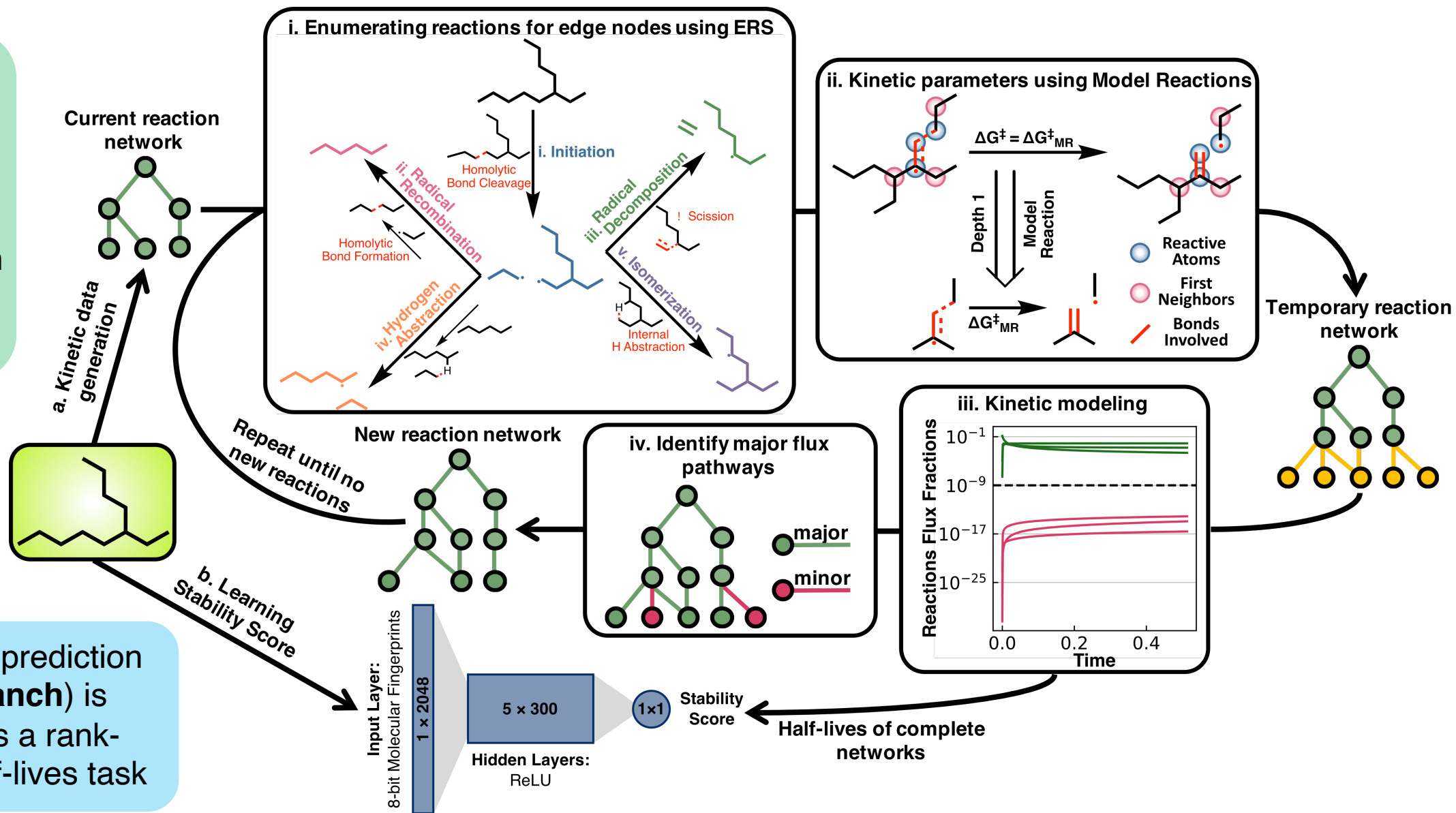
Can the Mixture of Expert Models Perform Out-of-Distribution?



Minor product identification was tested using 18 reactants with 50 distinct products from Grossman's textbook and Hartenfeller et al.

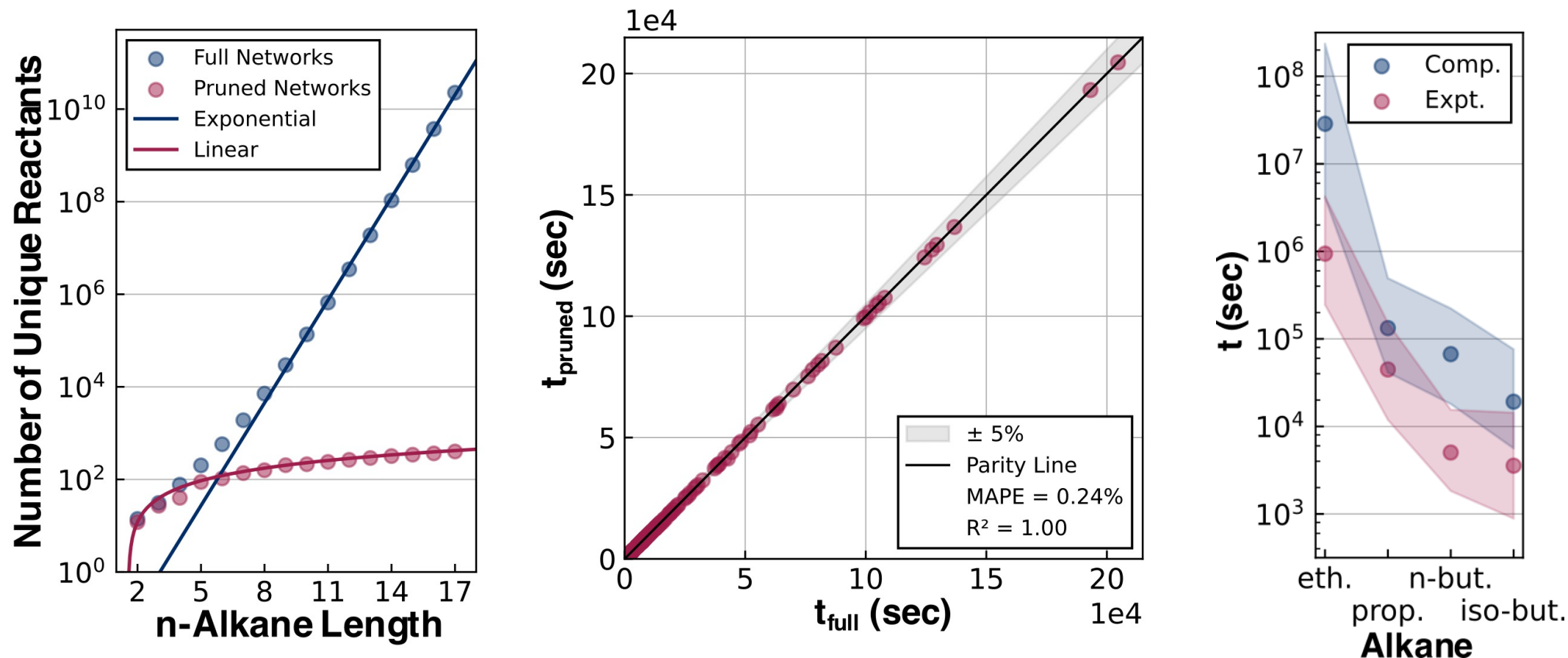
Actionable Design Information: Thermal Stability Scores

Computational half-life generation (**top-branch**) is expensive and incompatible with on-the-fly design needs.



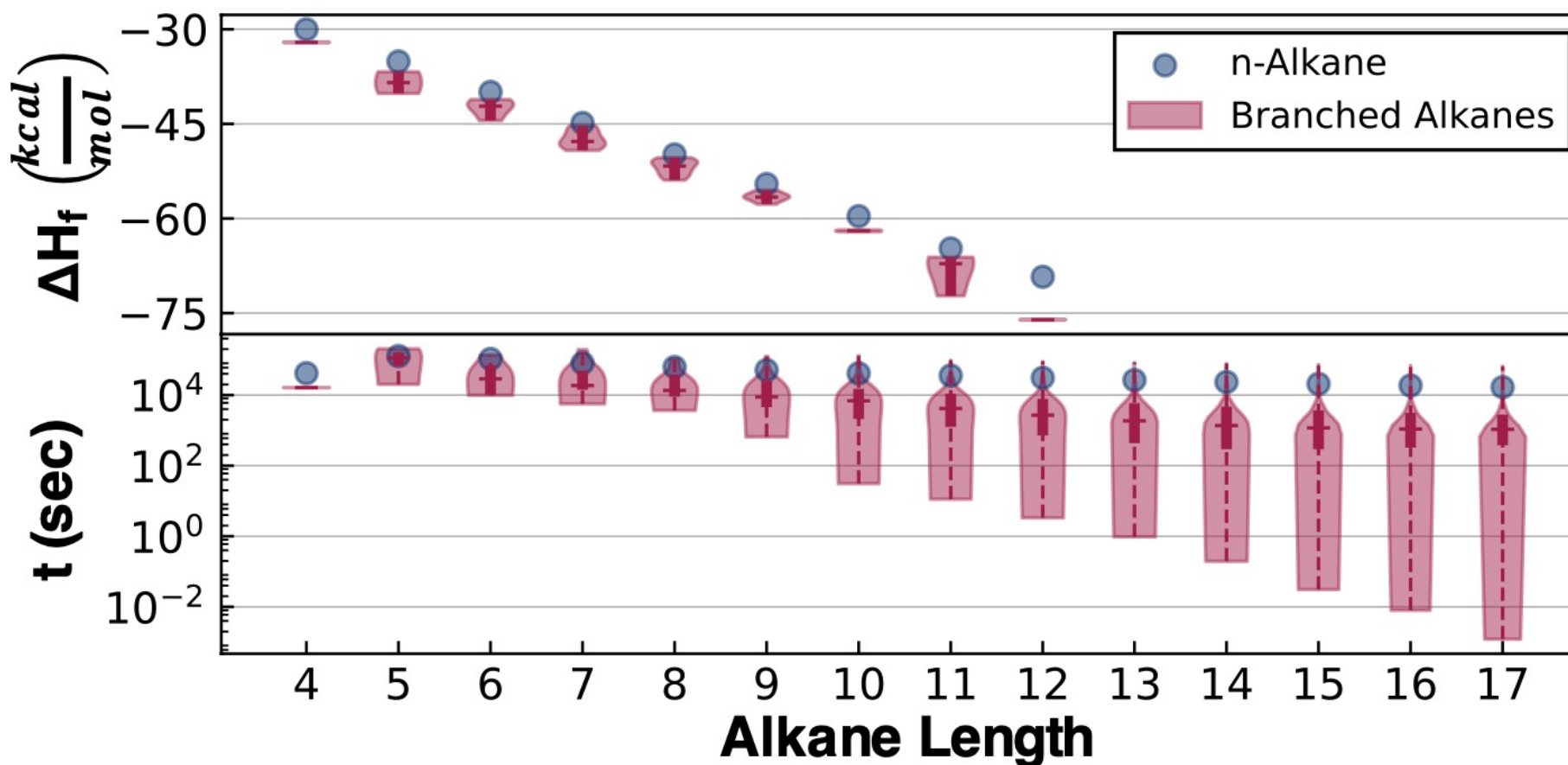
Stability score prediction (**bottom branch**) is formulated as a rank-ordering of half-lives task

Half-Life Simulations are Non-Trivial



The half-lives of $\sim 32\text{k}$ alkanes with varying topology were simulated under pyrolysis conditions as a surrogate for thermal stability

Kinetics Matter for Stability



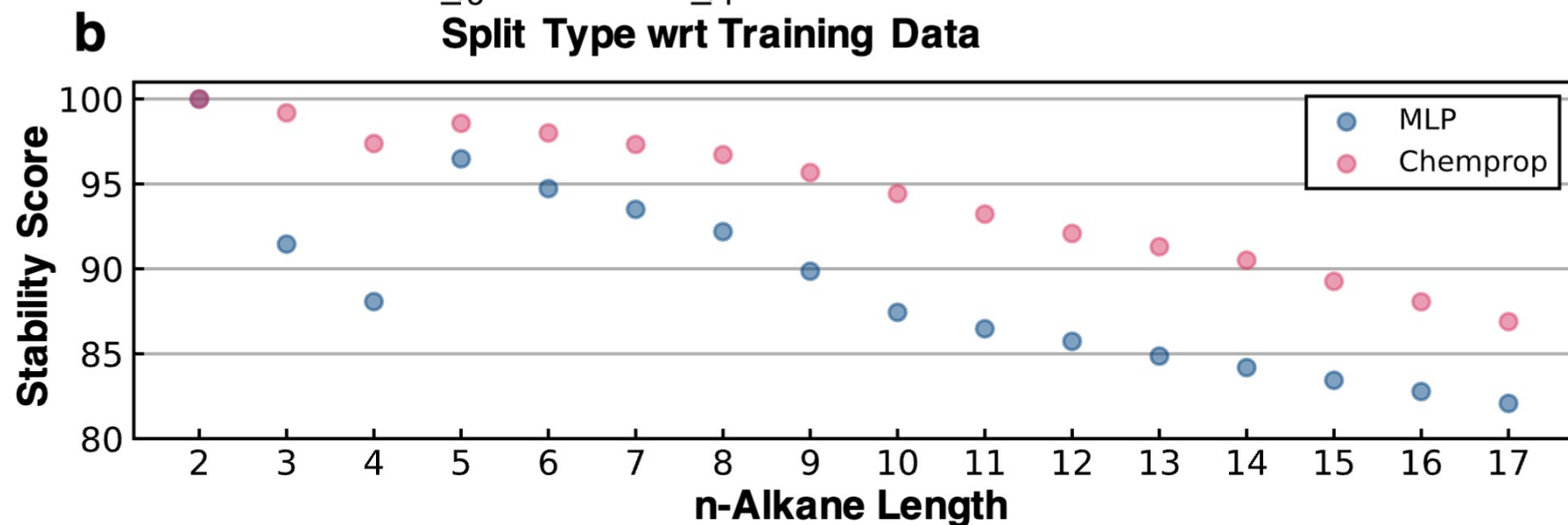
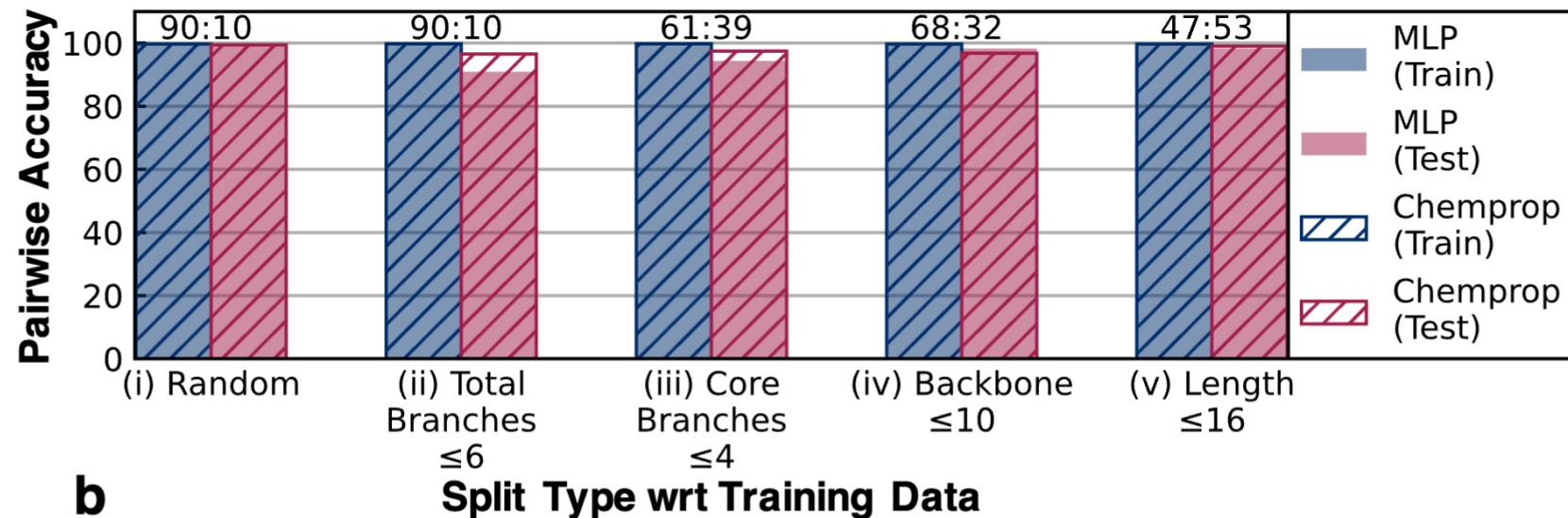
Heats of formation are completely non-predictive for relative stability (even for the simplest organic materials class).

Kinetics Matter for Stability

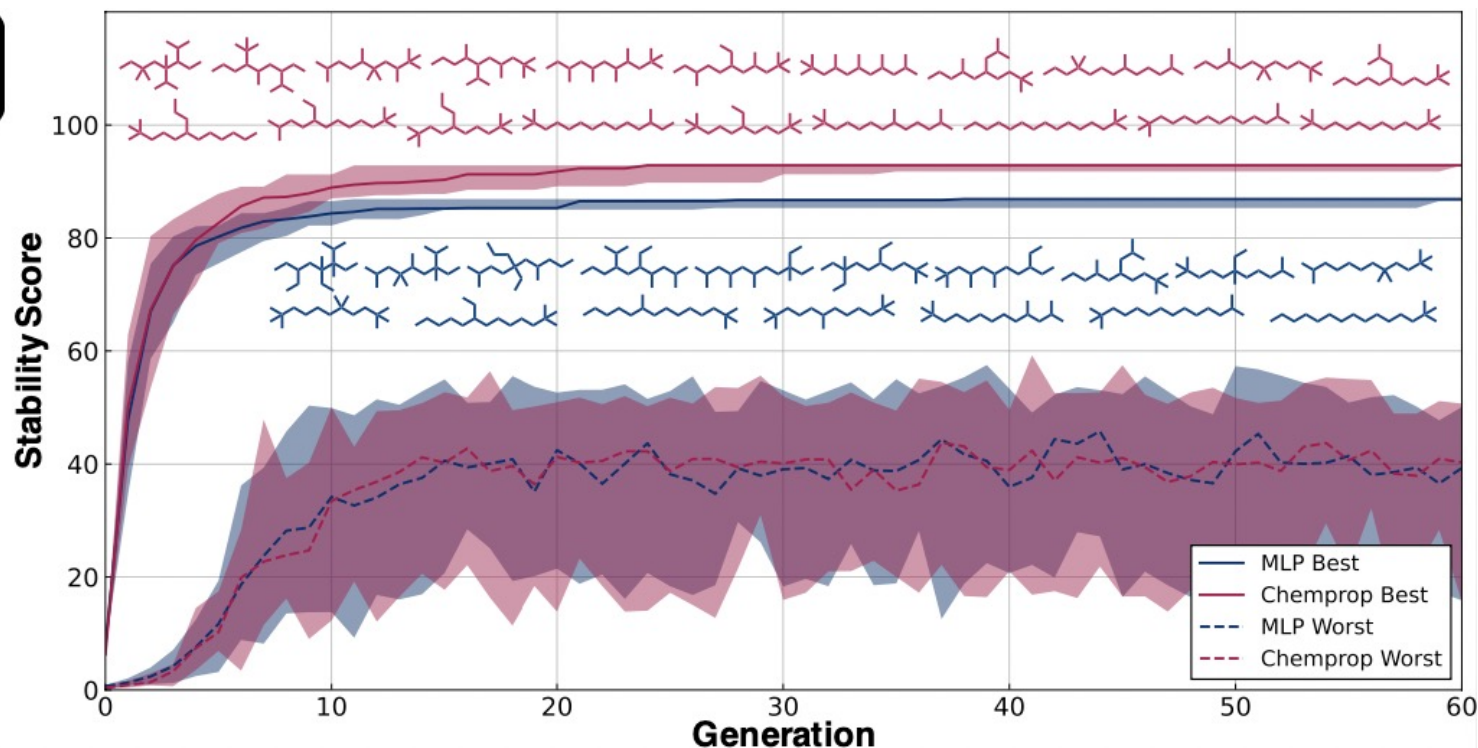
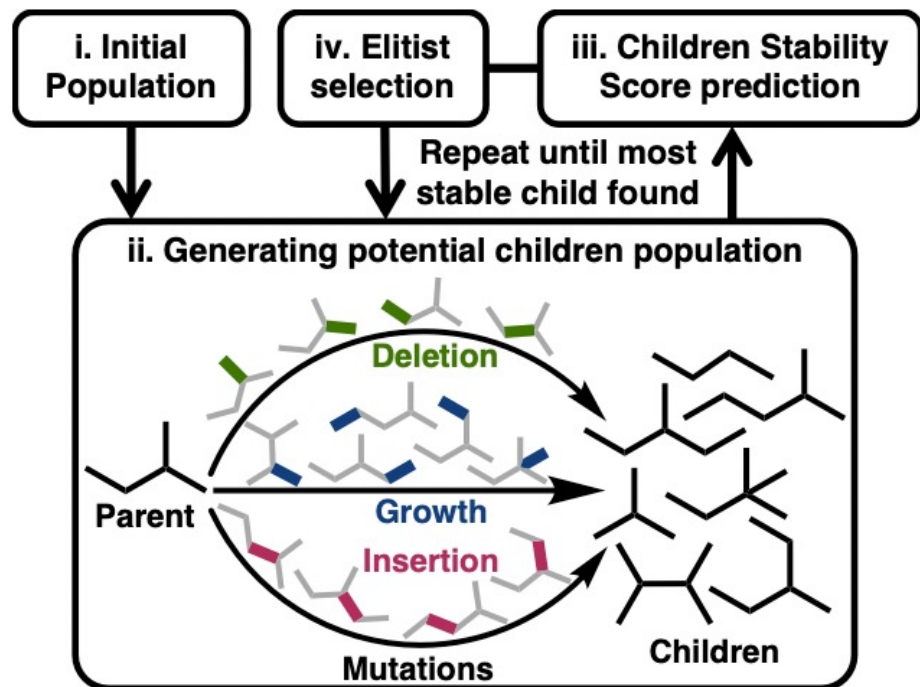
The stability score is trained to predict the rank order of the log half-life using a hinge-loss function

The relative stability is really easy to learn (basically every architecture we tried >95% accurate)

Non-trivial kinetic stability trends are learned in testing set



Actionable Design Information: Thermal Stability Scores



Genetic algorithms are used to explore chemical space, using the stability score as an objective function

The simple (ML) and complex (Chemprop) models are both extremely proficient at guiding the search towards stable alkanes with a minimal number of branches

Great, so you have an alkane stability score...

This is just the foundation showing that half-life to stability score is a learnable task. It now becomes a data problem.

Teasers:

- Stability is stressor-specific and multi-dimensional, so half-lives under other conditions are coming.
- We already see evidence that thermal stability scores are predictive of degradation temperature for polymers.

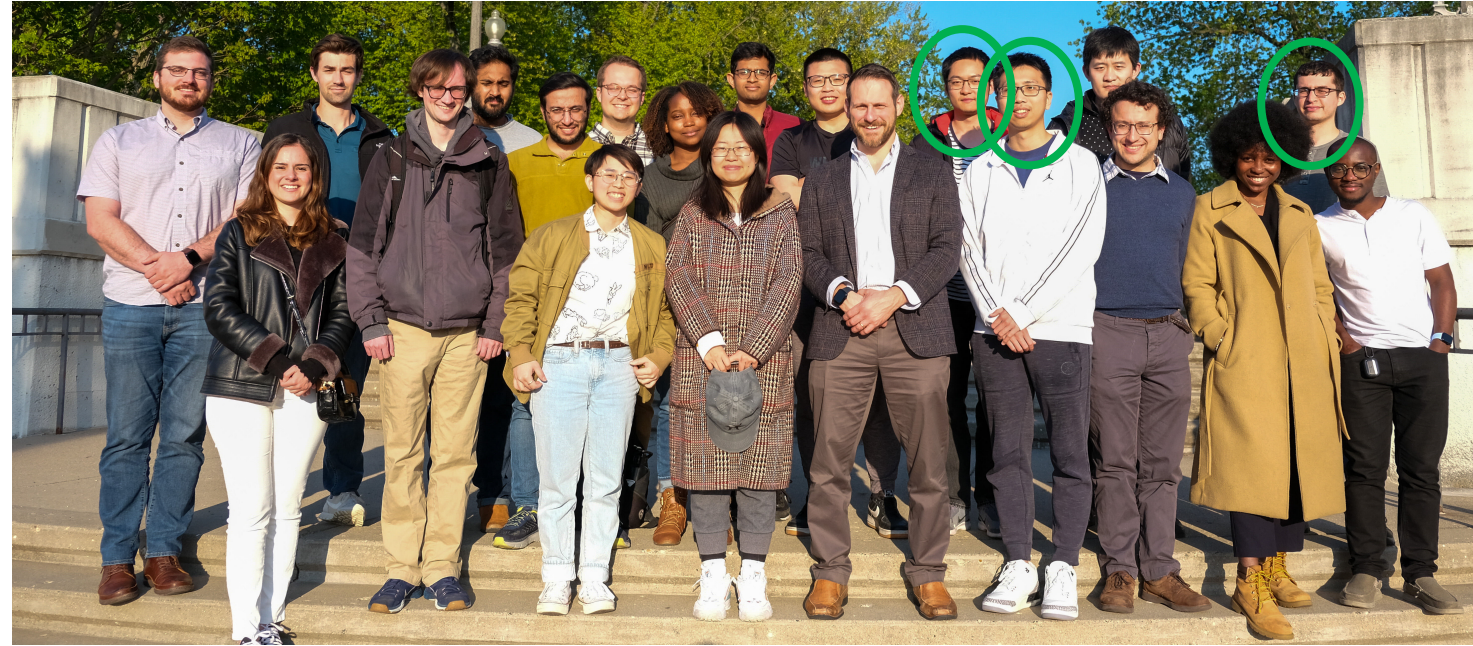
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