

#### "Known Unknowns" and "Unknown Unknowns"

#### $A \rightarrow B$

- To safely plan a known reaction, we need access to solid thermodynamic data (e.g.,  $\Delta H_f$ , S°, C<sub>v</sub>) to understand and classify risks.
- This is a "known unknown" in that we know the reaction, A → B, but we need values for a few unknown variables.

#### $A \rightarrow ? \rightarrow B ; A \rightarrow B + ? ; A \rightarrow ?$

- A → ? → B, means that we know the net reaction, but there may be a consequential (e.g., potentially reactive) intermediate. Even if we have accurate thermodynamic data on A/B, neglecting the intermediate could be disastrous.
- The  $A \rightarrow B + ?$  (unknown side-reaction) and  $A \rightarrow ?$  (unknown main product), problems have similar "unknown unknown" characteristics.

#### "Known Unknowns" and "Unknown Unknowns"

 $A \rightarrow B$ 

# TAFFIComponent Increment Theory

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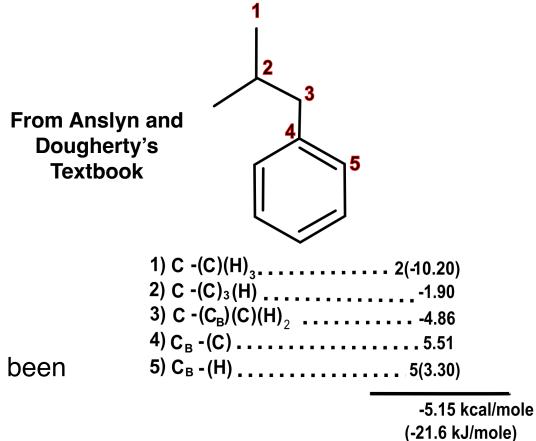
### **Challenges of Contemporary Group Theories**

#### **Benson Group Theory:**

- The idea is to decompose molecular properties ( $\Delta H_f$ , S°, C<sub>v</sub>) 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.



Experimental ΔH<sub>f</sub>: -5.15 +/- 0.34 kcal/mol

• Extensibility: because of the provenance and specificity problems, it isn't possible to develop new groups in a consistent way.

#### **Challenges of Contemporary Group Theories**

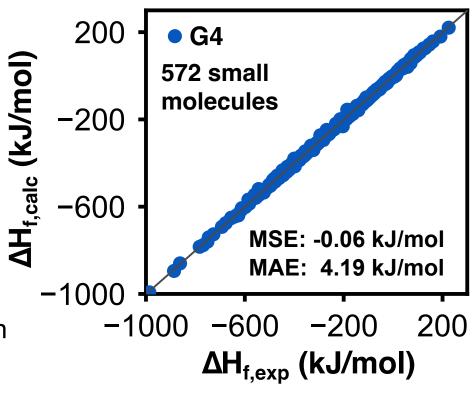
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#### ΔH<sub>f</sub> from modern quantum chemistry



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

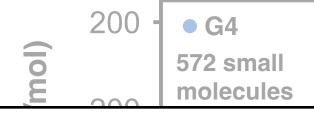
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ΔH<sub>f</sub> from modern quantum chemistry



• Groud on trust data, a

Prob

Speciformal

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

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#### The fundamental idea

• Systematize component-definitions and model compound selection with rigorous graph-based typing.

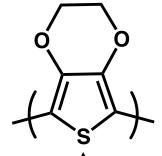
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P2SAC Publications

TCIT is a <u>component</u> theory (2-bond specific)



Topology Automated Force Field Interactions



graph/structure equivalence

Adjacency matrix for PEDOT monomer

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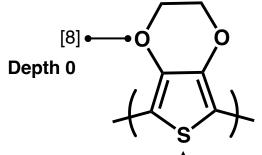
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S	Γ0	1	0	0	1	0	0	0	0	0	0	0	0٦
С	1	0	1	0	0	0	0	0	0	0	0	0	0
С	0	1	0	1	0	1	0	0	0	0	0	0	0
С	0	0	1	0	1	0	0	0	0	0	1	0	0
С	1	0	0	1	0	0	0	0	0	0	0	0	0
0	0	0	1	0	0	0	1	0	0	0	0	0	0
С	0	0	0	0	0	1	0	1	1	1	0	0	0
С	0	0	0	0	0	0	1	0	0	0	1	1	1
	_	_	_	_	_	_	_	_	_	_	_	_	$\sim$
Η	0	0	0	0	0	0	1	0	0	0	0	0	0
H H	0	0	0	0	0	0	1	0	0	0	0	0	0
	ľ	-	·	•	-	-		-	·	-	•	·	Ĭ
Н	0	0	0	0	0	0	1	0	0	0	0	0	0
H O	0	0	0	0	0	0	1 0	0	0	0	0	0	0

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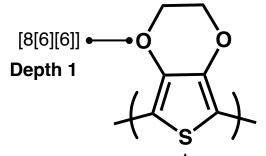
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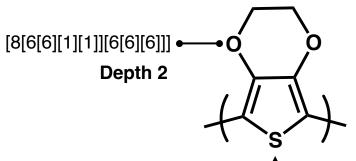
S	٥٦	1	0	0	1	0	0	0	0	0	0	0	0٦
С	1	0	1	0	0	0	0	0	0	0	0	0	0
С	0	1	0	1	0	1	0	0	0	0	0	0	0
С	0	0	1	0	1	0	0	0	0	0	1	0	0
С	1	0	0	1	0	0	0	0	0	0	0	0	0
0	0	0	1	0	0	0	1	0	0	0	0	0	0
С	0	0	0	0	0	1	0	1	1	1	0	0	0
С	0	0	0	0	0	0	1	0	0	0	1	1	1
Н	l٥	0	0	0	0	0	1	0	0	0	0	0	o l
		U	U	U	•	•	_	•	•	•	U	U	٠,
Н	0	0	0	0	0	0	1	0	0	0	0	0	0
	ľ	-	•	•	-	-		-	-	-	-	•	0
Н	0	0	0	0	0	0	1	0	0	0	0	0	٦
H O	0	0	0	0	0	0	1 0	0	0	0	0	0	0

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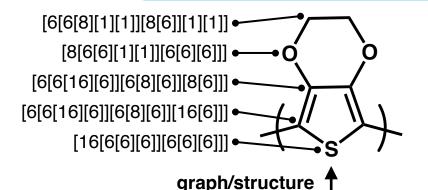
P2SAC Publications

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P2SAC Publications

Adjacency matrix for PEDOT monomer

#### The fundamental idea

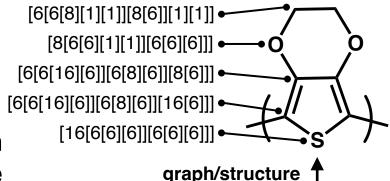
- Systematize component-definitions and model compound selection with rigorous graph-based typing.
- Two-bond specificity should improve both the accuracy and transferability of the resulting components.
- Parameterizing a component model would not be feasible with only experimental data.

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P2SAC Publications

Adjacency matrix for PEDOT monomer

How will we select molecules for parameterizing TCIT components?

identify components\*

1-hydroxy-pent-2-ene-2-one

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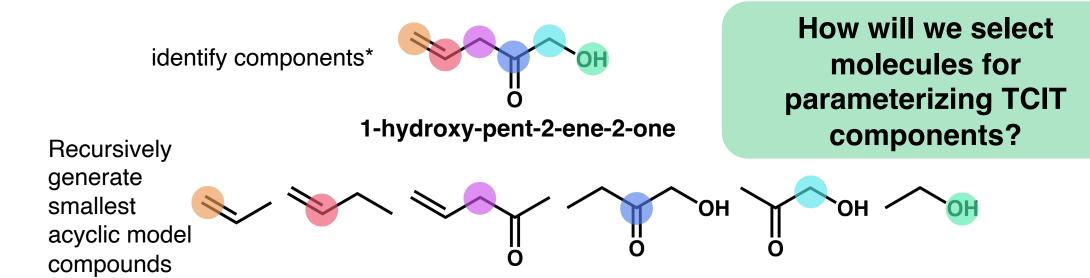
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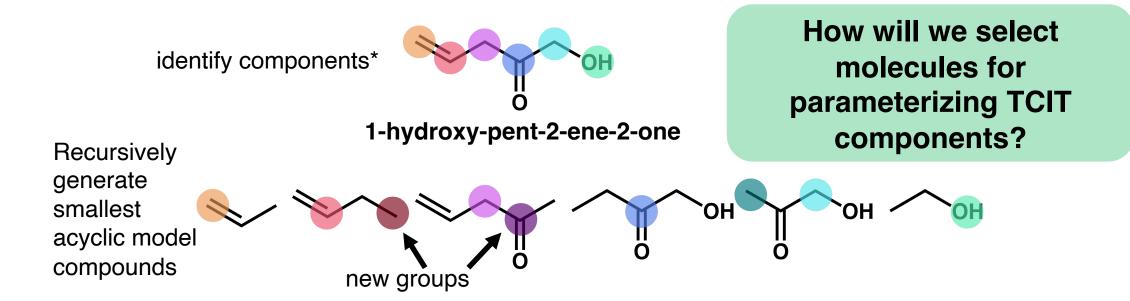


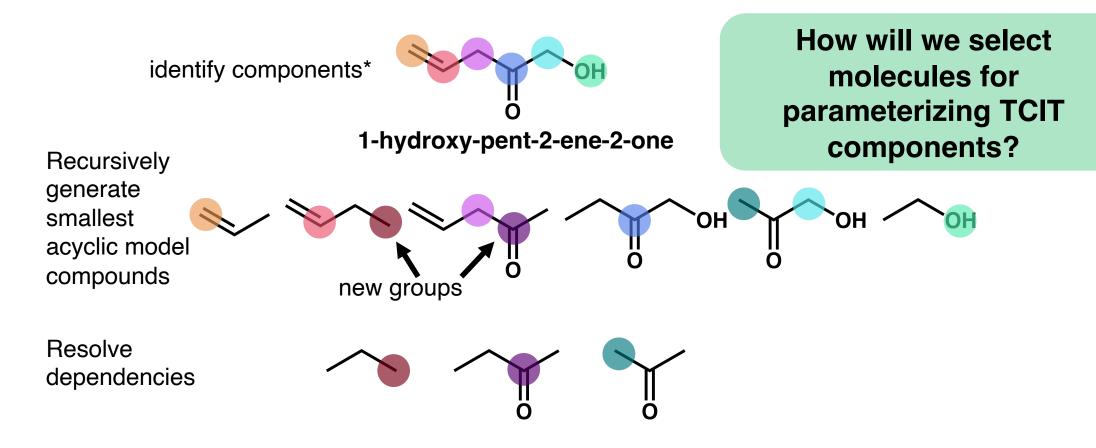
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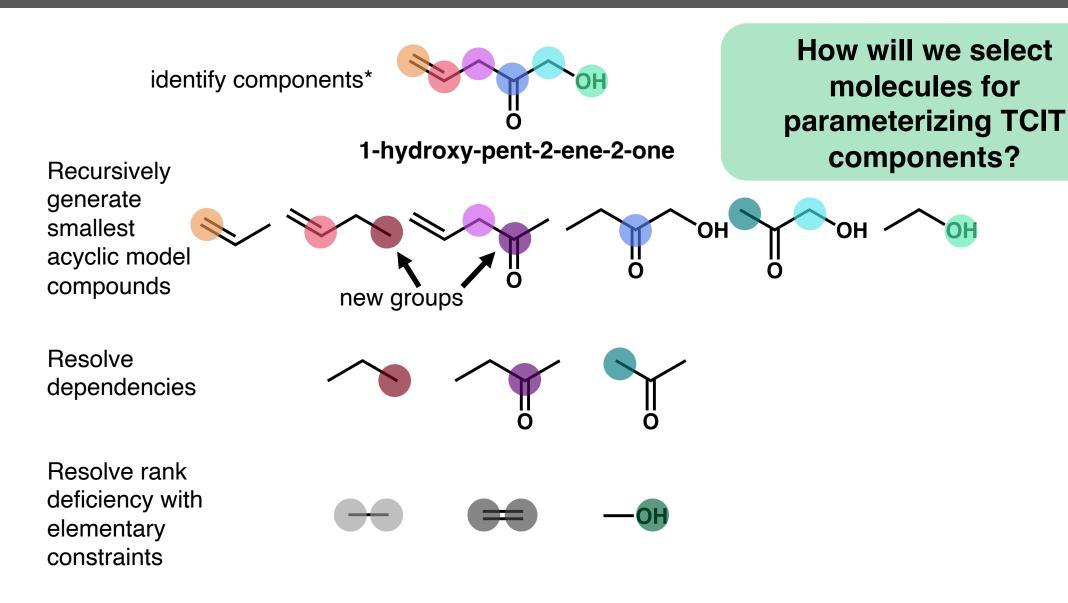
Recursively generate smallest acyclic model compounds

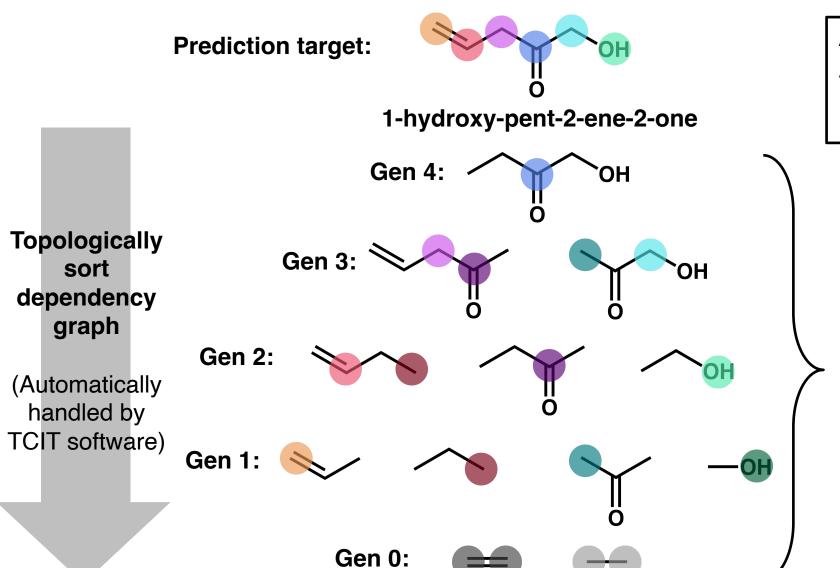
How will we select molecules for parameterizing TCIT components?











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

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

#### 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  $\Delta H_f$  data is calculated at the G4 composite level, no exceptions.

handled by

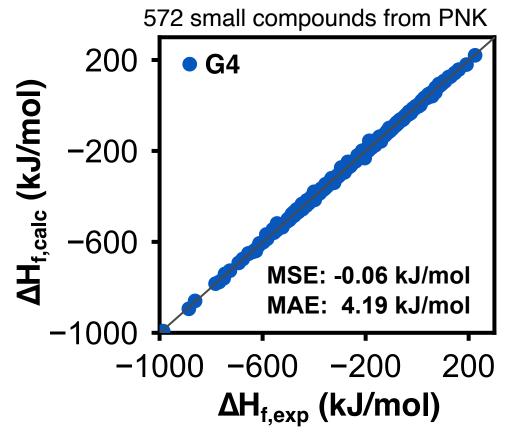
### Have we solved the extensibility problem?

Model compounds exist for all conceivable components, no exceptions.

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

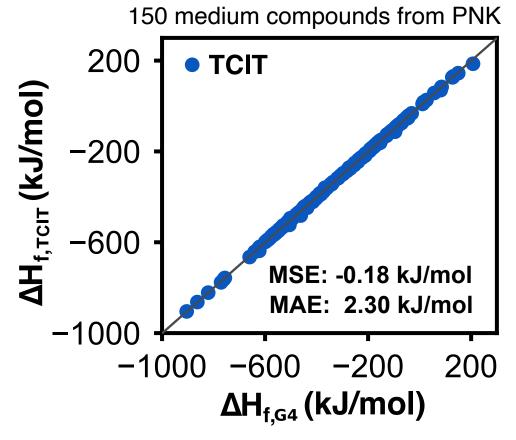
- Initial benchmarking set consists of ~1100 linear C,H, and O containing compounds from PNK¹
- (1) J. B. Pedley, R. D. Naylor, S. P. Kirby "Thermochemical Data of Organic Compounds" 2<sup>nd</sup> ed. 1986
- PNK is a core dataset for fitting Benson groups
- ~600 PNK compounds are small enough for G4 calculations and comparison with experiment.



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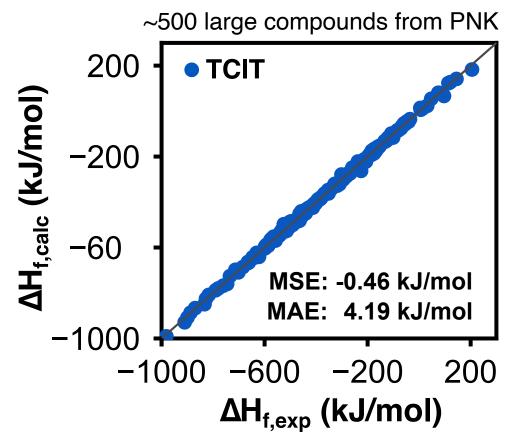
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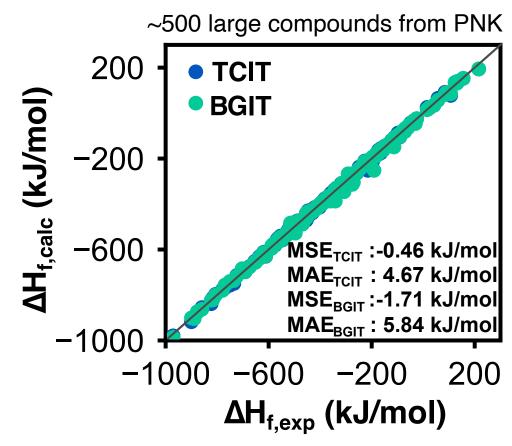
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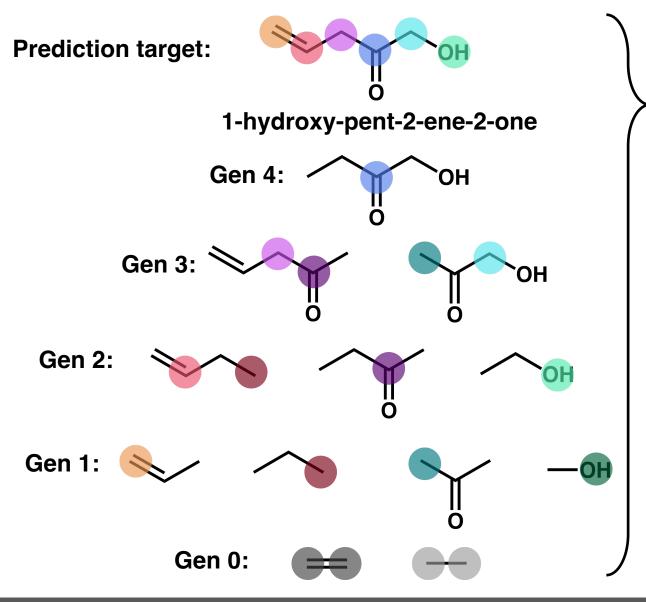
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TCIT shows comparable performance to BGIT/CHETAH but is derived exclusively from extensible G4 data.

### **How Many Components are Possible?**



We database all model compounds and components for reuse.

Over the past three years, we have parameterized new components in response to distinct project needs (many from P2SAC Pharma Members)

#### **Current Database:**

- $\sim$ 35k distinct components for  $\Delta H_f$  relevant to organic chemistry
- ~35k distinct G4 calculations on organic molecules.
- ~450 distinct ring corrections

### **How Many Components are Possible?**

**Prediction target:** 



We database all model compounds

How many components are required to predict the  $\Delta H_f$  of **all** (physically relevant) organic molecules?

Gen 3:



(many from P2SAC Pharma Members)

How many P2SAC funding periods would it take to make a "complete" or "gapless" component theory?

Gen 0:

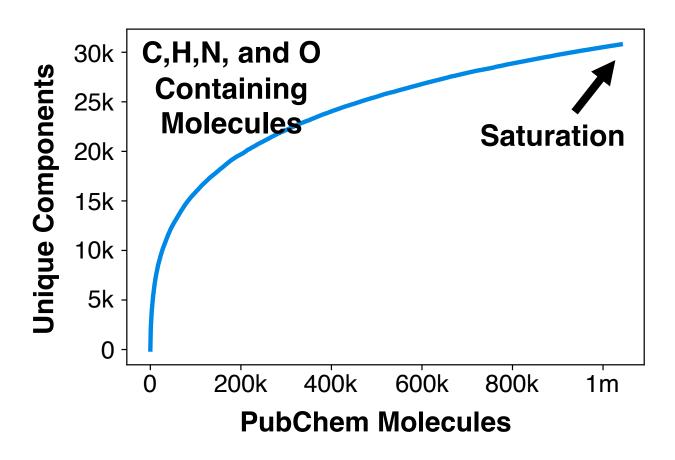




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**PubChem** is a repository of chemical properties that contains many millions of organic species ranging from small molecules to oligonucleotides.

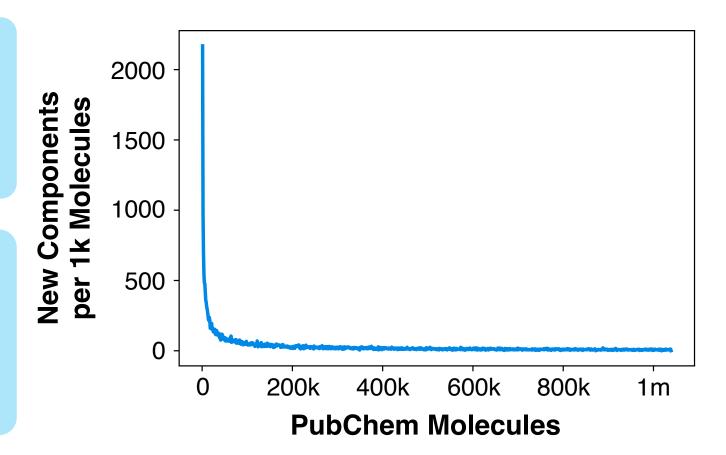
We recently started mining PubChem's H,C,N, and O containing molecules for distinct components and the model compounds necessary to predict ΔH<sub>f</sub>



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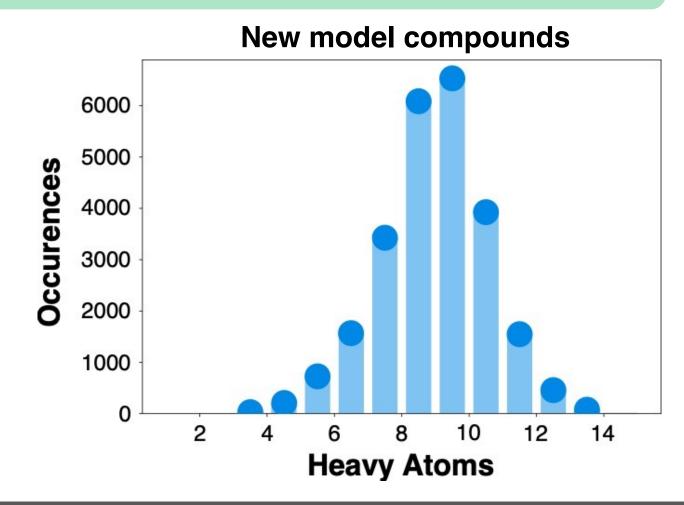
The derivative plot shows that TCIT initially generates ~2 new components per molecule, but by the end of the sampling ~100 molecules need to be sampled to find a new component.



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TCIT now contains all CAVs necessary to predict ΔH<sub>f</sub> 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.

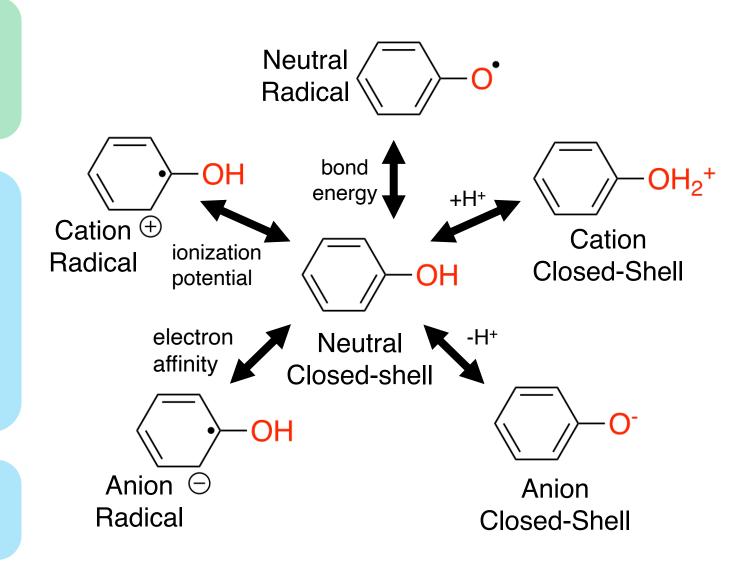
**Heavy Atoms** 

#### **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+

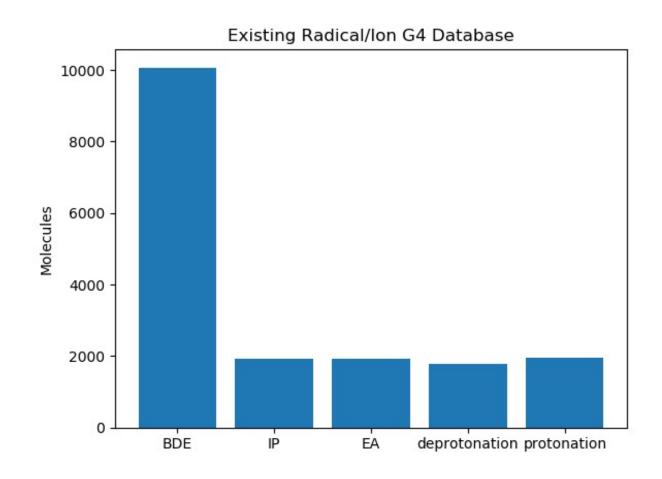


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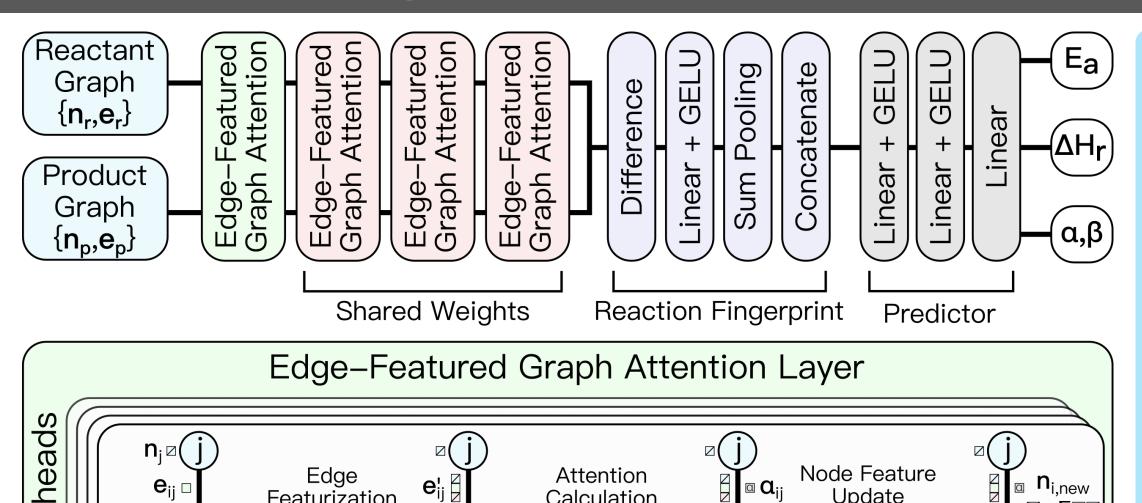
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#### Graph → IP/EA/+H+/-H+ Models



Attention

Calculation

# been activation **EGAT** developed architectu energy by prediction has group for already

 $\mathbf{n}_{\mathsf{i},\mathsf{new}}$ 

 $=\sum \alpha /$ 

Node Feature

Update

**X** 

 $n_i \square$ 

 $e_{ij}$ 

Edge

Featurization

e'ij

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#### **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

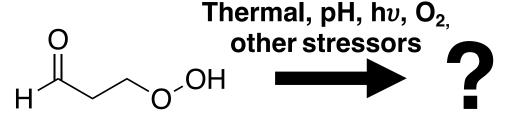
A → ?: For degradation reactions, plausible reactions are often unknown.

Savoie Research Group | | | |

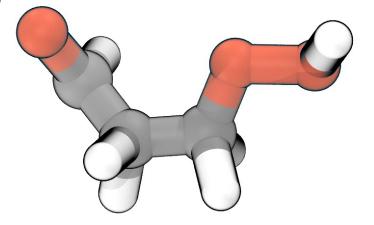
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3-hydroperoxypropanal



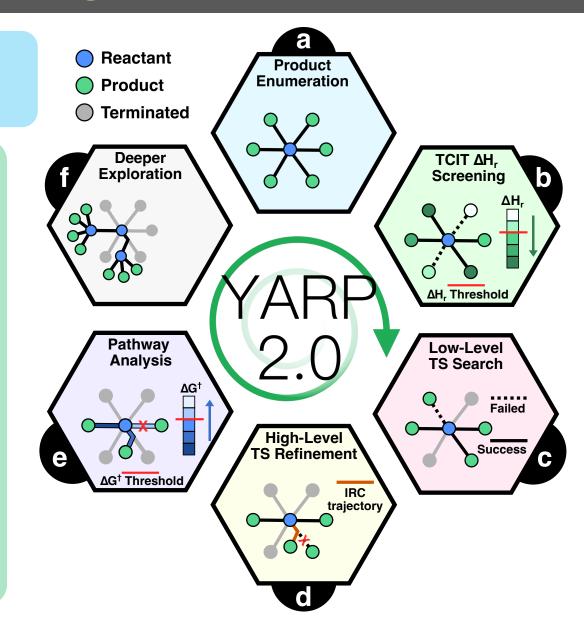
Savoie Research Group

## Yet Another Reaction Program (YARP)

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

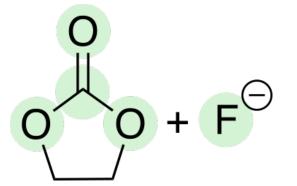
#### **Observations:**

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

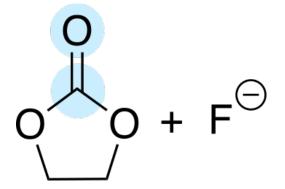


Polar and pericyclic organic reactions are decomposed into elementary electron donor and acceptor reactions with concomitant σ-bond breaks

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



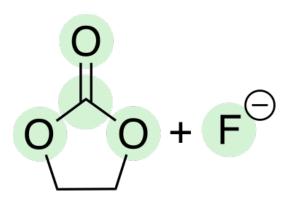
Lone-Pair Donors



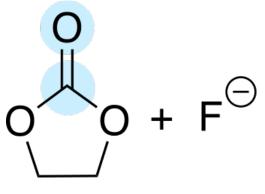
**Lone-Pair Acceptors** 

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

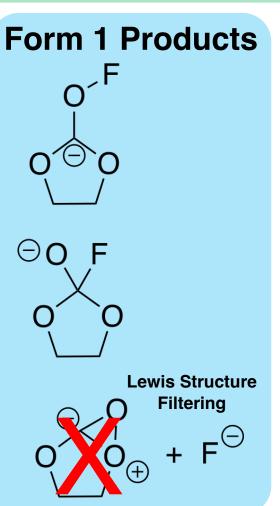
bnfn
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rearrange.



**Lone-Pair Donors** 

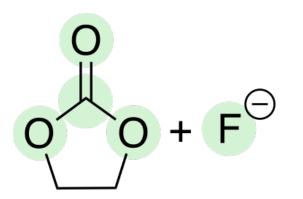


Lone-Pair Acceptors

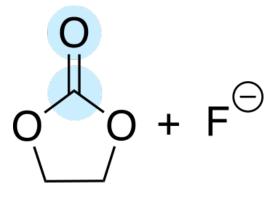


Polar and pericyclic organic reactions are decomposed into elementary electron donor and acceptor reactions with concomitant σ-bond breaks

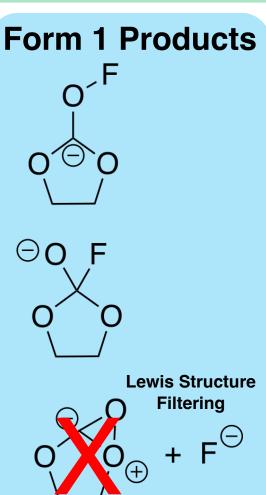
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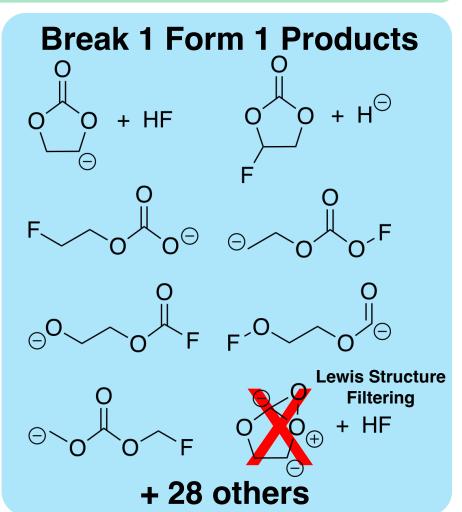


**Lone-Pair Donors** 



Lone-Pair Acceptors





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

Form 1 Products

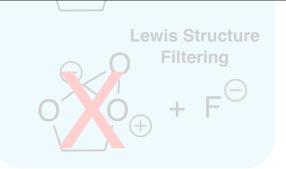
**Break 1 Form 1 Products** 

## All bnfn products are b(n-1)f(n-1) decomposable

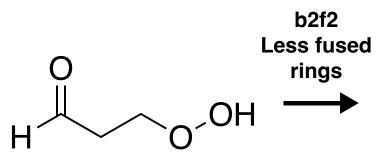
This means that using only "break 1 bond form 1 bond" (b1f1) for radicals and ions won't miss any products, but it will potentially miss important transition states (i.e., by predicting a sequential mechanism when a concerted mechanism is favored)

o arbitrarily rearrange.

WI

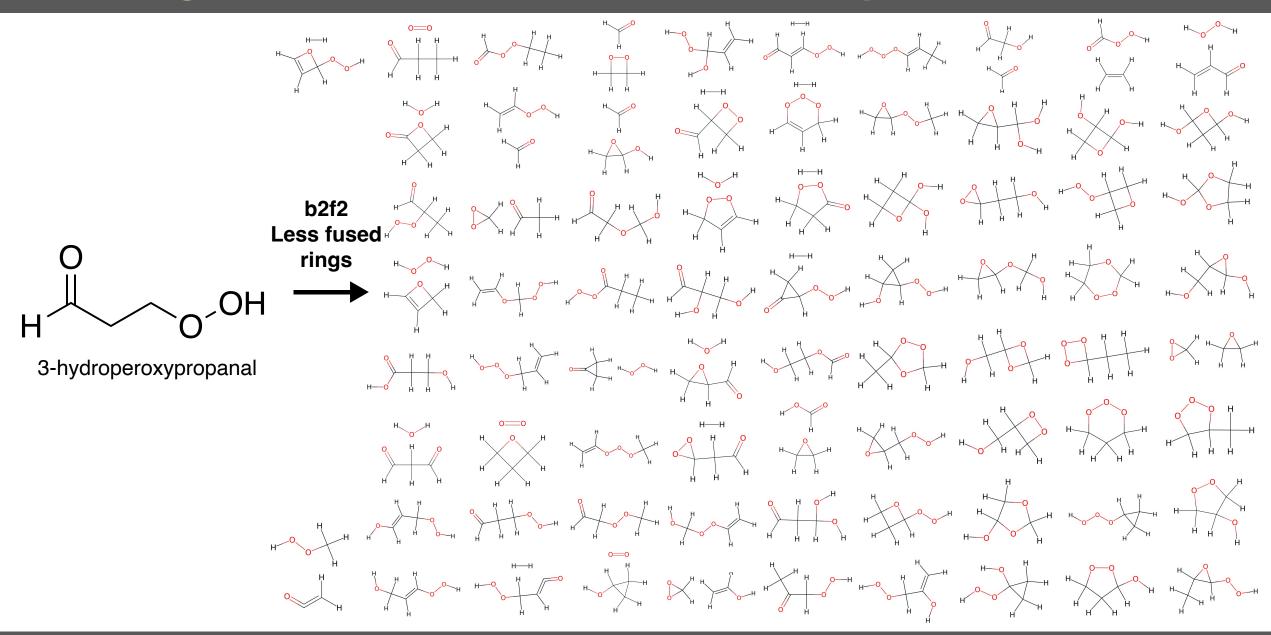


## Testing YARP on a Unimolecular Decomposition Problem

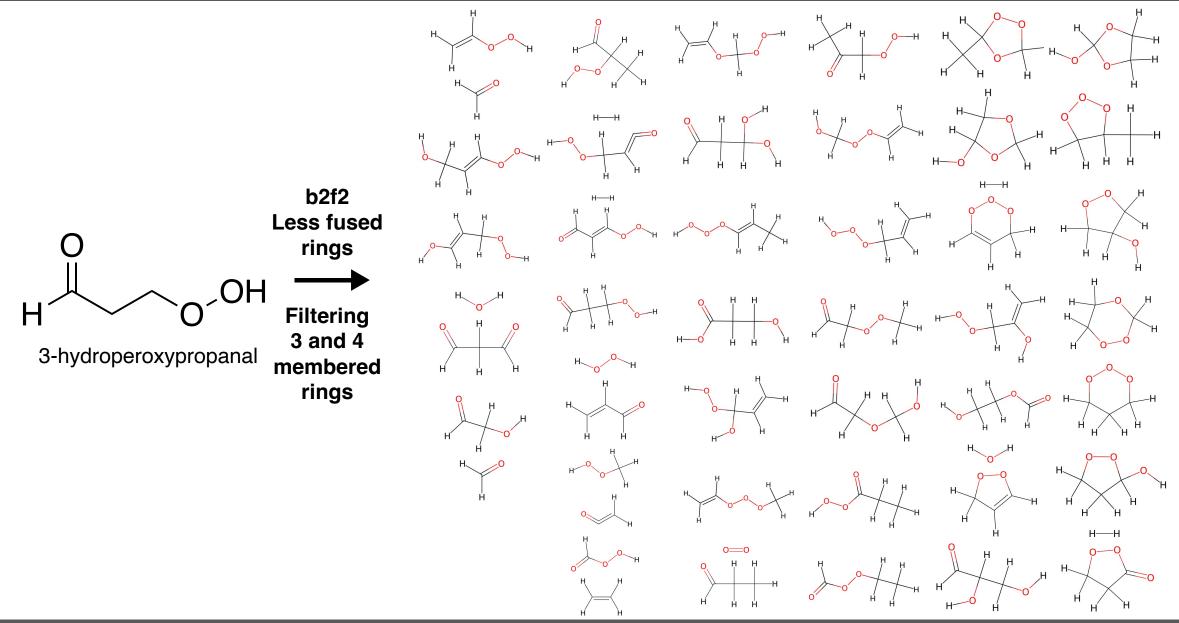


3-hydroperoxypropanal

## Testing YARP on a Unimolecular Decomposition Problem



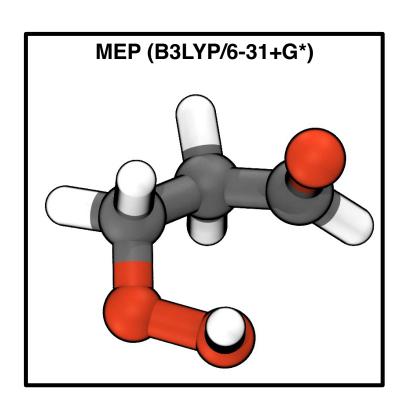
# Testing YARP on a Unimolecular Decomposition Problem



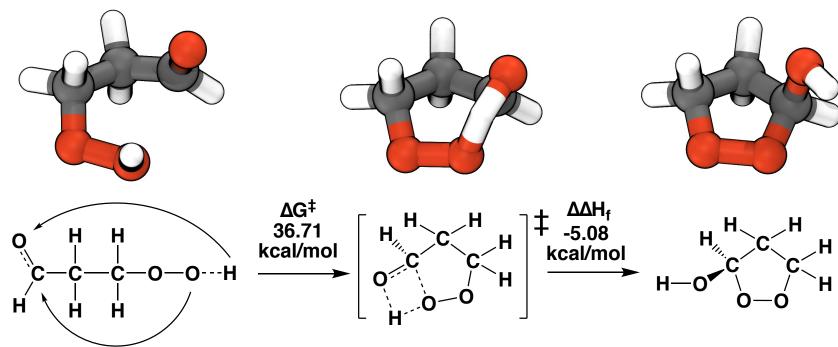
### What Happens First?

Jensen, R. K.; Korcek, S.; Mahoney, L. R.; Zinbo, M. JACS 1979, 101, 7574

#### The Korcek Mechanism

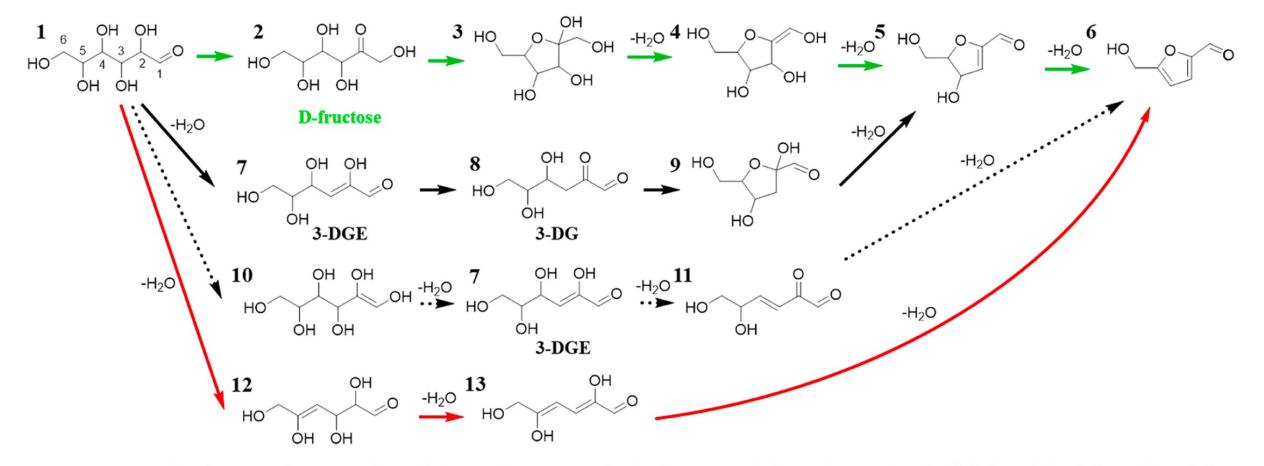


According to YARP, this is the lowest barrier unimolecular reaction.



Fully resolved (along with subsequent ROOH and R=O formation) 30 years later by Green and Truhlar: Jalan, A.; Alecu, I. M.; Meana-Pañeda, R.; Aguilera-Iparraguirre, J.; Yang, K. R.; Merchant, S. S.; Truhlar, D. G.; Green, W. H. *JACS* 2013, 135 (30), 11100–11114.

## Reaction Network Case Study: β-D-Glucose Pyrolysis



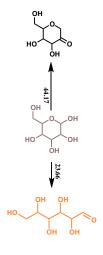
**Figure 1.** Proposed pathways in literature from glucose to HMF, namely the fructose path (green), 3-DG paths (black and black dotted), and direct path (red). The molecules are indicated by numbers and some key molecules are named as follows: **1.** D-glucose; **2.** D-fructose; **3.** D-fructofuranose; **6.** 5-hydroxymethylfurfural (5-HMF); **7.** 3-deoxyglucos-2-ene (3-DGE); **8.** 3-deoxyglucosone (3-DG); and **10.** hex-1-ene-1,2,3,4,5,6-hexaol (enol form of glucose).

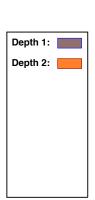




To perform a deep network exploration, we've implemented a modified version of Dijkstra's algorithm

- (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



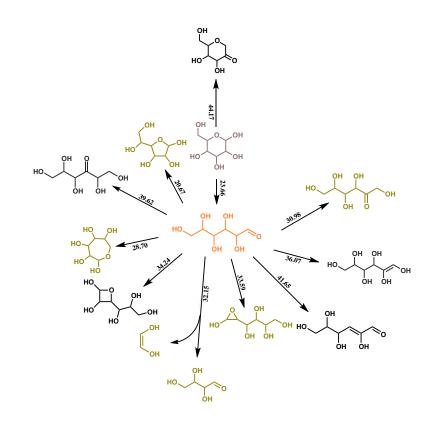


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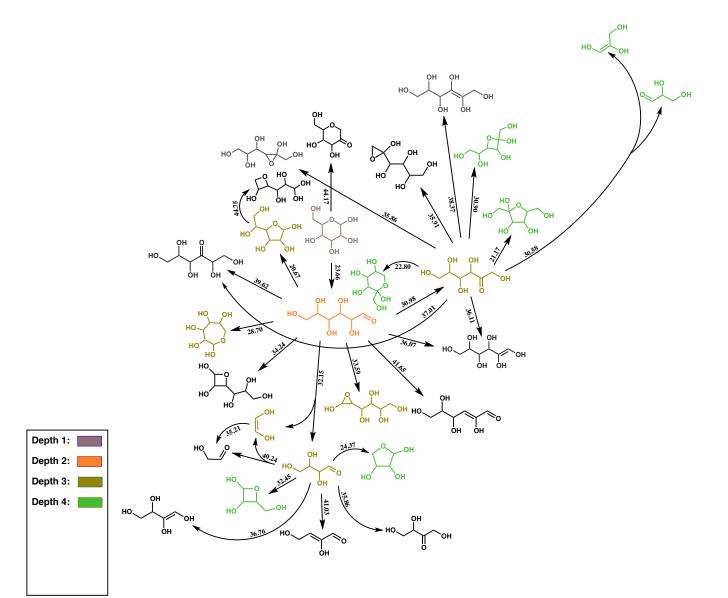
Depth 1: Depth 2: Depth 3:

## **β-D-Glucose Pyrolysis Network Exploration**



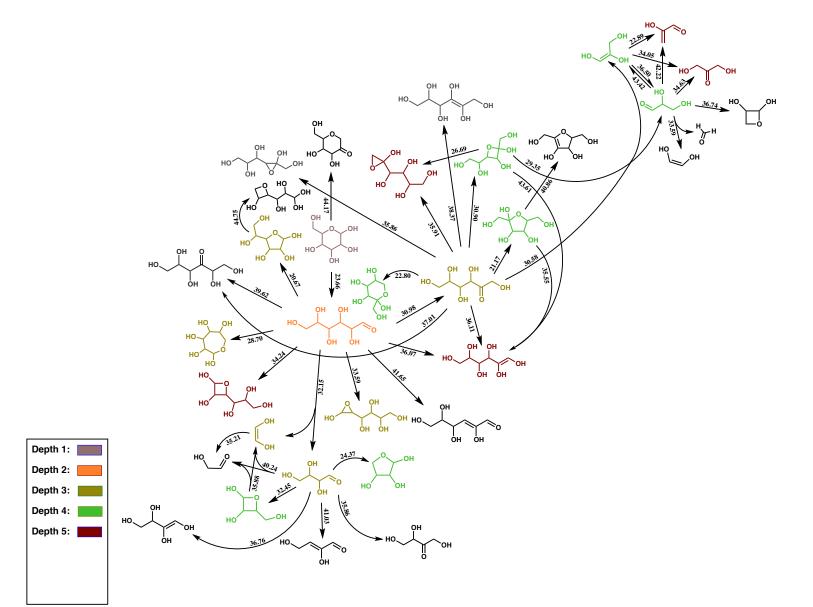
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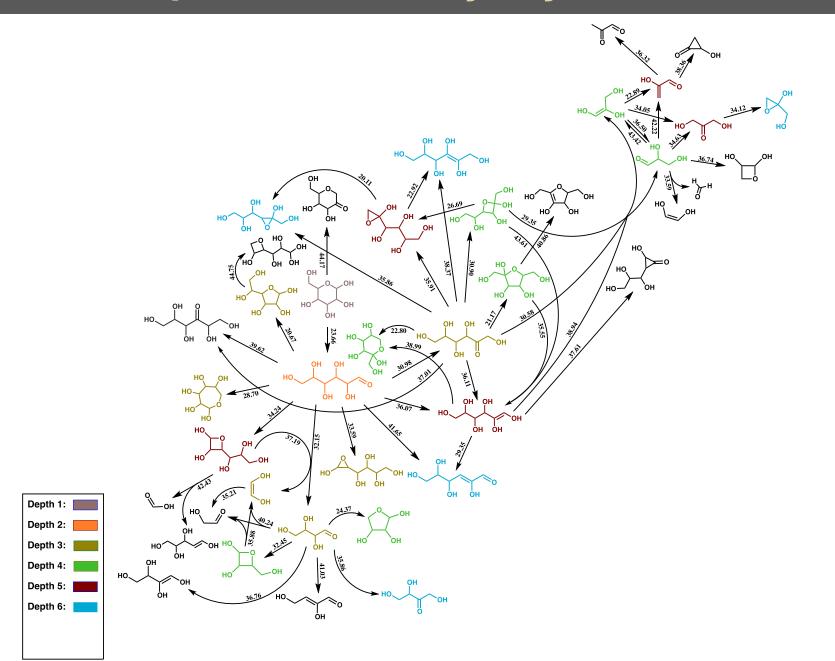
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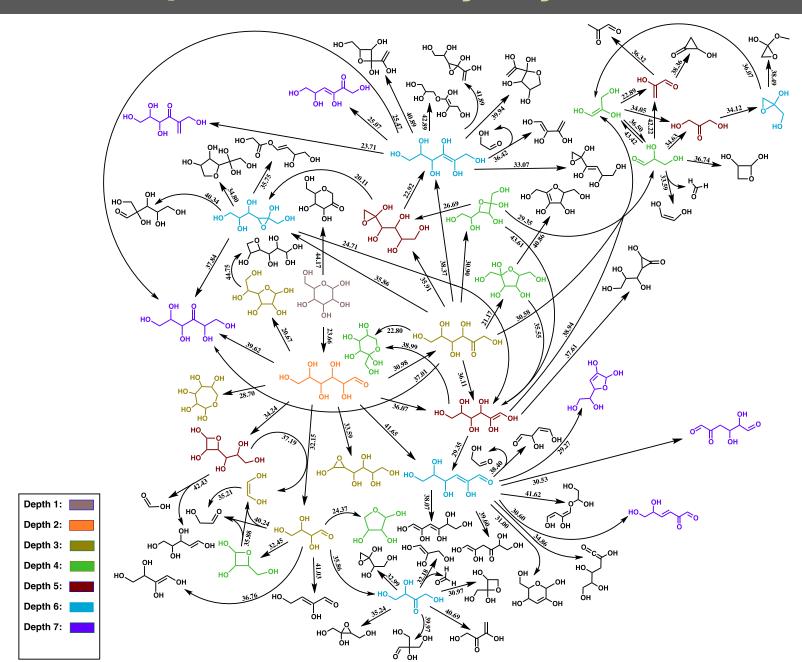
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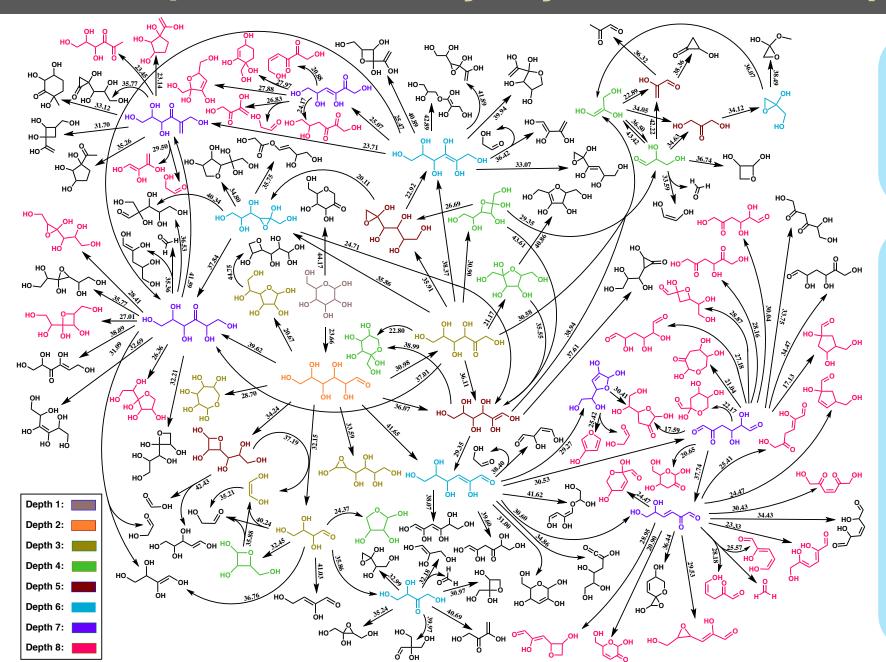
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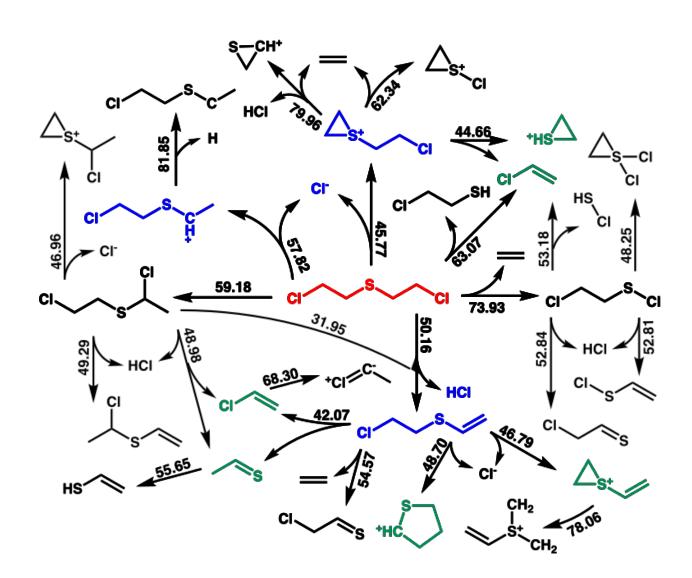
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## Studying Chemical Warfare Agents (CWAs) with YARP

Degradation products are often the only evidence of CWA use or existence. Establishing mechanistic pathways provides evidentiary value to investigators.

CWA type	Chemical agents	Method of exposure	Clinical symptoms
Nerve agents	G-agents (sarin, cyclosarin, tabun, soman)	Inhalation	SLUDGE, miotic pupils, bradycardia, bronchospasm, bronchorrea, muscle spasms/fasciculations, weakness, flaccid paralysis, tachycardia, seizures, respiratory failure
	V-agents (VE, VG, VM, VR, VX)		
Blistering agents	Nitrogen mustard & sulfur mustard (mustard gas)	Inhalation	Acute: Skin, eye and lung damage (pulmonary edema and pulmonary hemmorhage), erythematous rash, skin blistering
			Chronic: Lung damage (chronic obstructive pulmonary disease, asthma, bronchiolitis obliterans), neutropenia, pancytopenia
Asphyxiants	Carbon monoxide, chlorine, phosgene, hydrogen sulfide gases	Inhalation	Upper airway distress, skin and eye irritation, fatal pulmonary edema and acute respiratory distress syndrome
Blood agents	Cyanide	Skin absorption, inha- lation and ingestion	Severe distress, tachycardia, cyanosis, hypotension, severe metabolic acidosis, seizures, cardiac arrest
Hydrofluoric acid		Skin absorption, inhalation and ingestion	Severe pain in exposed area, gastrointestinal distress, vomiting, cardiac arrhythmias, hypocalcemia, hyperkalemia

## YARP Prediction for Sulfur Mustard (HD) Reactivity

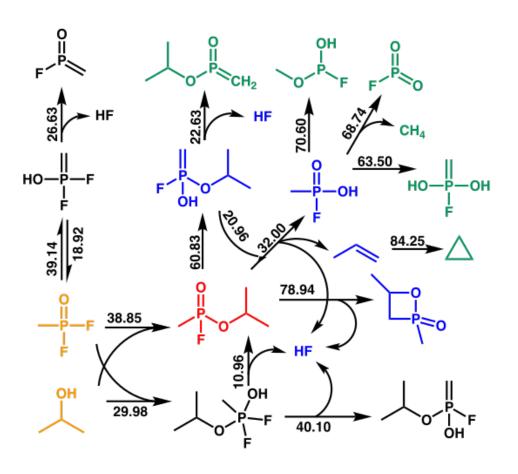


Mechanism of Action

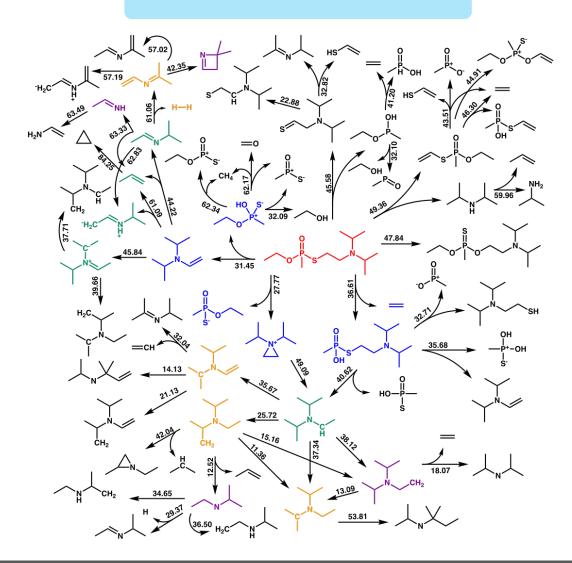
Lowest barrier bimolecular reaction

## Predicted Reactivity for Organophosphorus Nerve Agents

Sarin (GB)



VX



## **Outlook and Acknowledgements**

**Students:** Qiyuan Zhao, Tyler Pasut, Michael Woulfe

#### State-of-the-art:

- The accurate calculation of thermodynamic properties has become routine in many scenarios. Major opportunities lie in automation, systemization, and low-cost models.
- Practical solutions to the  $A \rightarrow ? \rightarrow B$ ,  $A \rightarrow B+?$ , and  $A \rightarrow ?$  problems are now available. We envision black-box tools for non-experts in the near future that will assist in hypothesis generation and potentially reactivity screening.



- P2SAC and ONR for funding.
- Ray Mentzer (Purdue)
- Spencer Goldrich(PMP)

