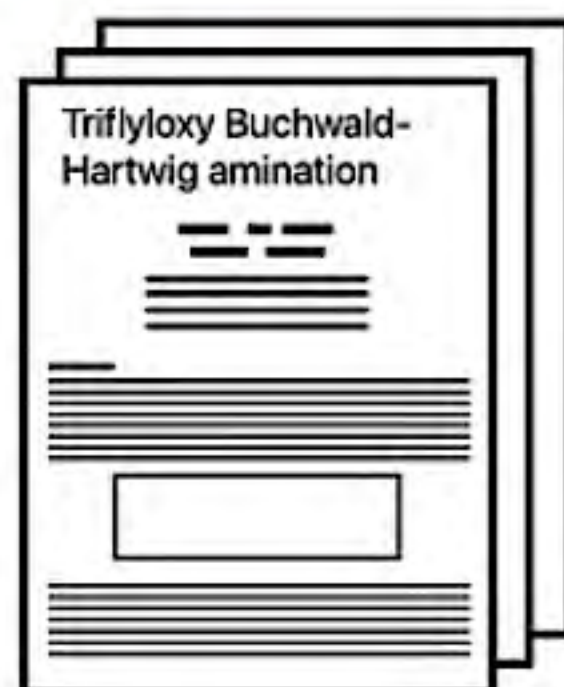
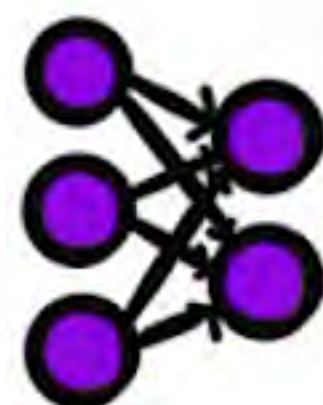
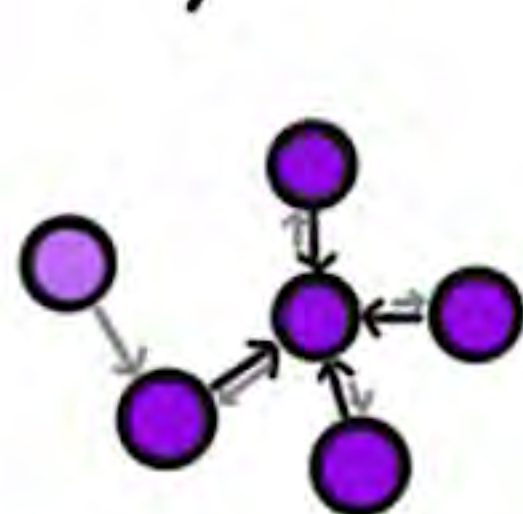




US20030166932A1: General Procedure
 H A solution of trifluoromethanesulfonic acid 3,5,8,8-tetramethyl-7,8-dihydronaphthalen-2-yl ester (Compound 35, 0.41 g, 1.2 mmol), Pd(OAc)₂ (0.027 g, 0.12 mmol), BINAP (0.11 g, 0.18 mmol), Cs₂CO₃ (0.56 g, 1.72 mmol), ethyl 4-aminobenzoate (0.25 g, 1.5 mmol) and 5 mL of toluene was flushed with argon for 10 min, then stirred at 100° C. in a sealed tube for 48 h. ...



EPFL
LIAC



Language models for accelerated discovery and synthesis

Philippe Schwaller (he/him/his)

Laboratory of Artificial Chemical Intelligence (LIAC) – SB-ISIC



@SchwallerGroup



philippe.schwaller@epfl.ch



NCCR
Catalysis

Grew up in Fribourg, Switzerland
 - French
 - Swiss German / German



NCCR
Catalysis

TT Assistant Professor
 in Digital Chemistry
 since Feb 2022

EPFL
LIAC

u^b

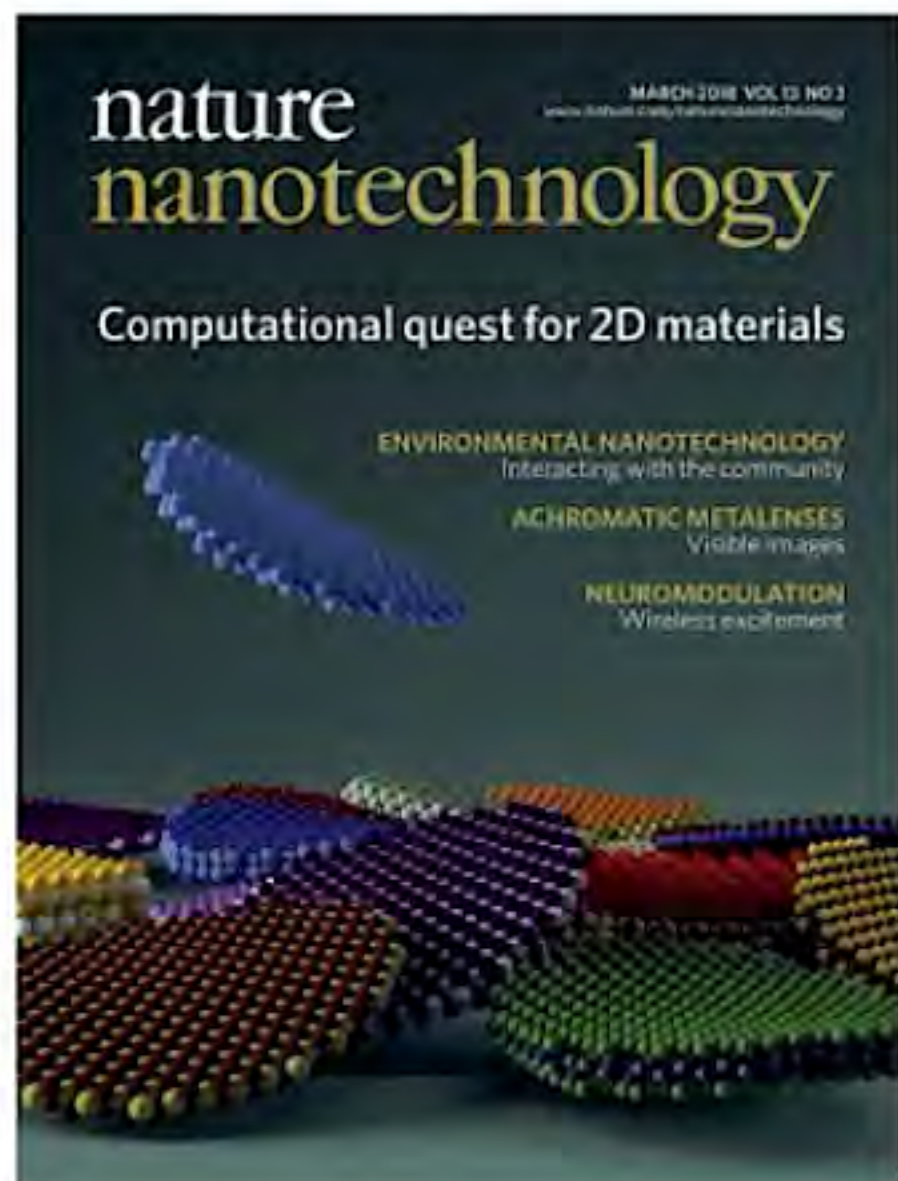
^b
**UNIVERSITÄT
 BERN**

PhD in Chemistry
 and Molecular Sciences ('21)
Prof Jean-Louis Reymond

EPFL

MANCHESTER
 1824
 The University of Manchester

Closing the loop...



Materials Science &
 Engineering

Virtual screening &
 simulation workflows
Prof Nicola Marzari

BSc ('14)/MSc ('16)



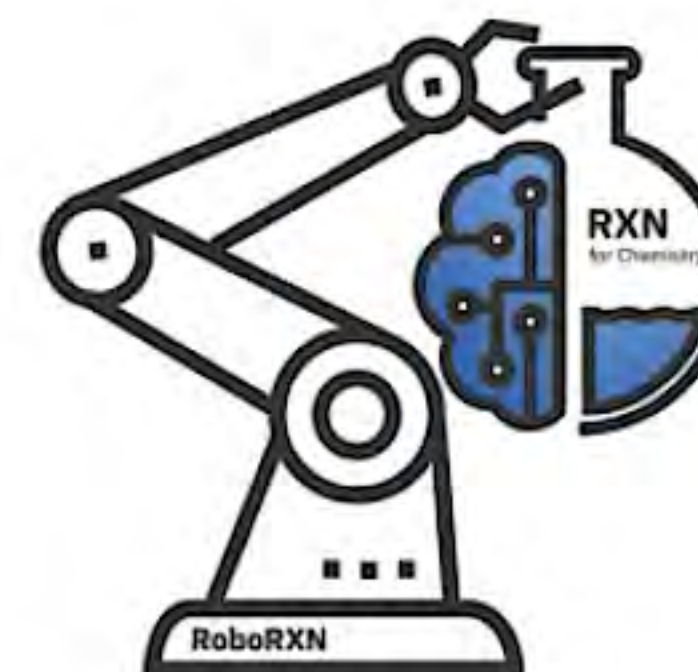
Materials Science and Technology

Lab work on ternary polymer
 blends for organic solar cells
 Prof Frank Nüesch

MPhil in Physics ('19)
 Dr Alpha Lee



**UNIVERSITY OF
 CAMBRIDGE**



Machine learning for chemical synthesis
 Intern/PhD/Postdoc
Dr Teodoro Laino



IBM Research Europe

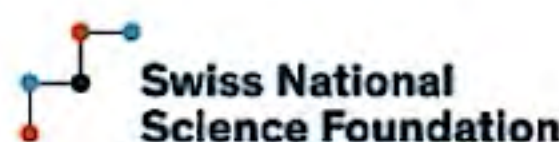
Mission:
**Accelerating
Discovery with
Digital
Chemistry**



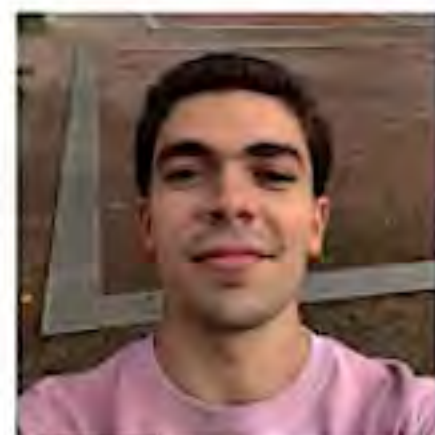
PhD students

Bojana Ranković
 Oliver Schilter (IBM Research)
 Andres CM Bran
 Junwu Chen
 Jeff Guo
 Victor Sabanza Gil (Lutembacher)
 Paulo Neves (Janssen)
 Rebecca Neeser (Correia, VantAI)
 Sarina Kopf (Nevado)
 Daniel Armstrong
 Joshua Sin (Roche)
 Sacha Raffaud
 Sandro Agostini (IBM Research)
 Théo Neukomm (Intel/Merck)

Funding:



IBM Research



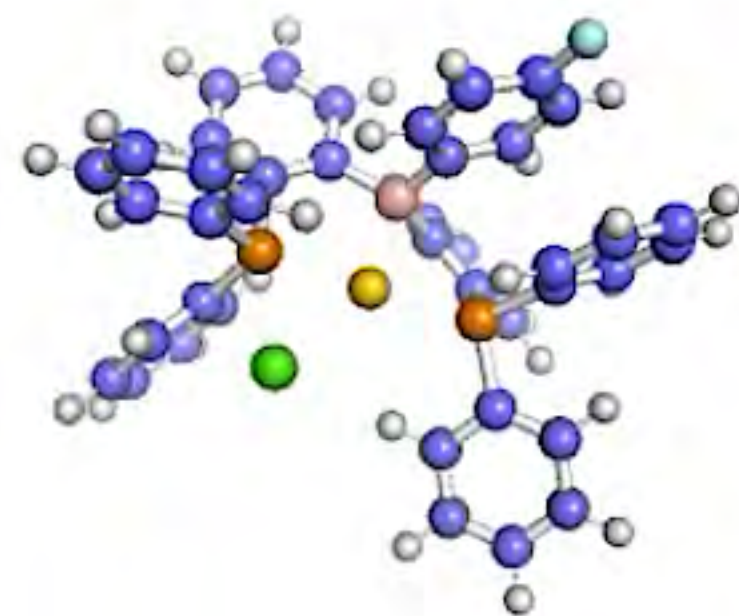
>10 nationalities — one team!
<https://schwallergroup.github.io>

A long-exposure photograph of the Milky Way galaxy arching across a dark night sky. The galaxy's core is visible as a bright, dense band of light in the upper right. Below the galaxy, a range of dark, jagged mountain peaks is silhouetted against a bright, orange-yellow glow on the horizon, likely from a low sun or moon. The overall scene is a vast, cosmic landscape.

Exploring the nearly
endless chemical
space

**10^{60} drug-like
molecules**

Cycle
What molecule to make?

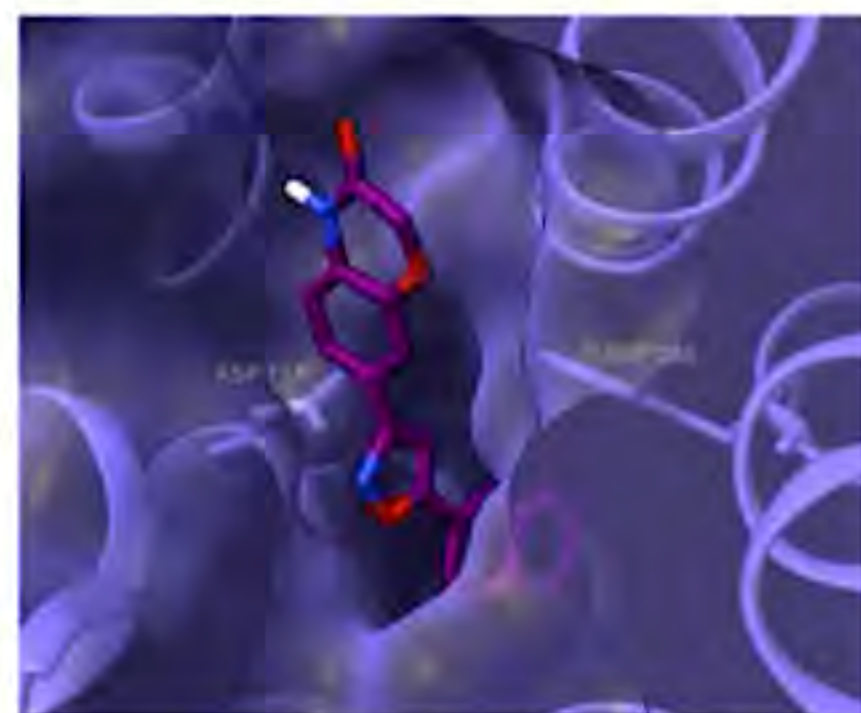


Catalysis



Saturn (Sample-efficient de novo molecular design)

Design

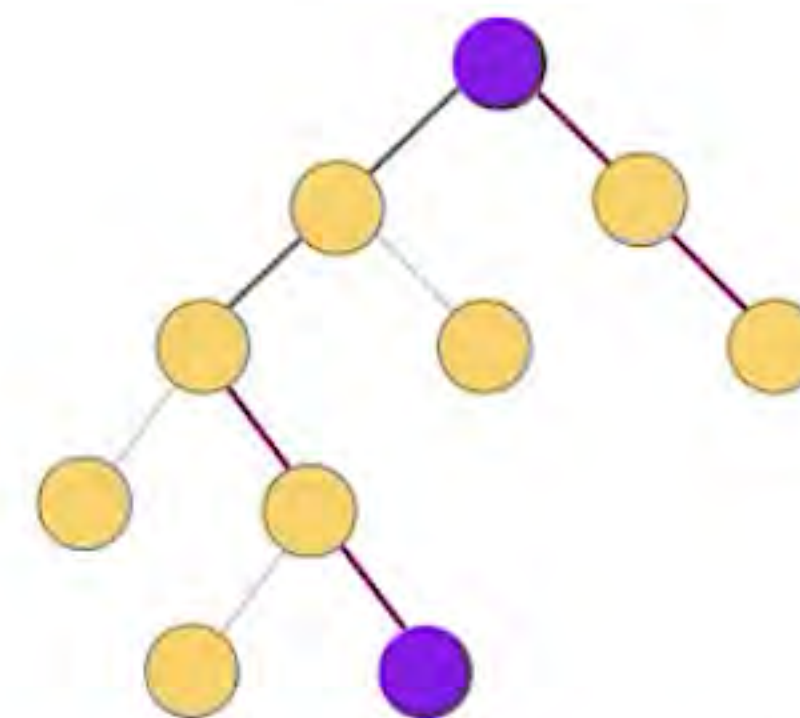


Drug discovery

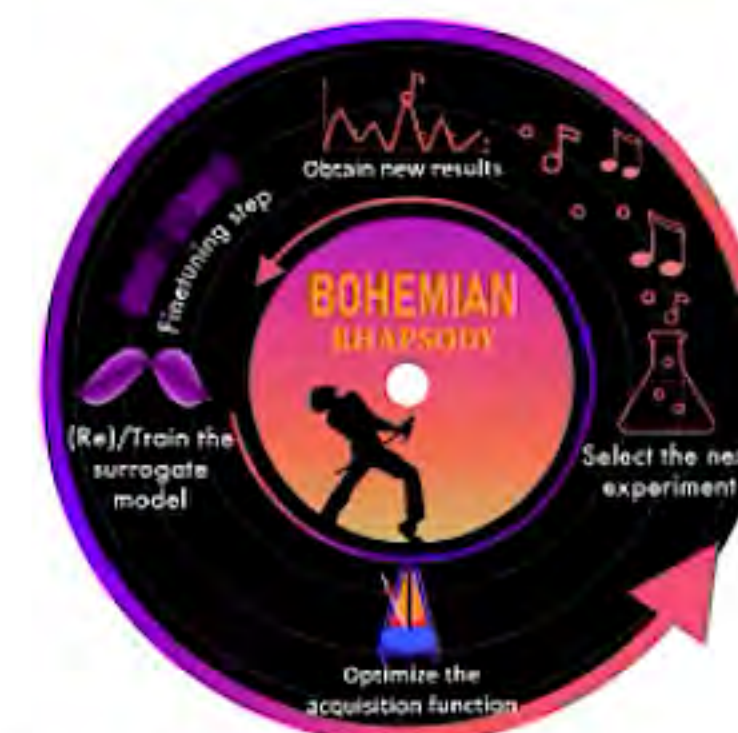
Experimental validation

How to make it?

Make



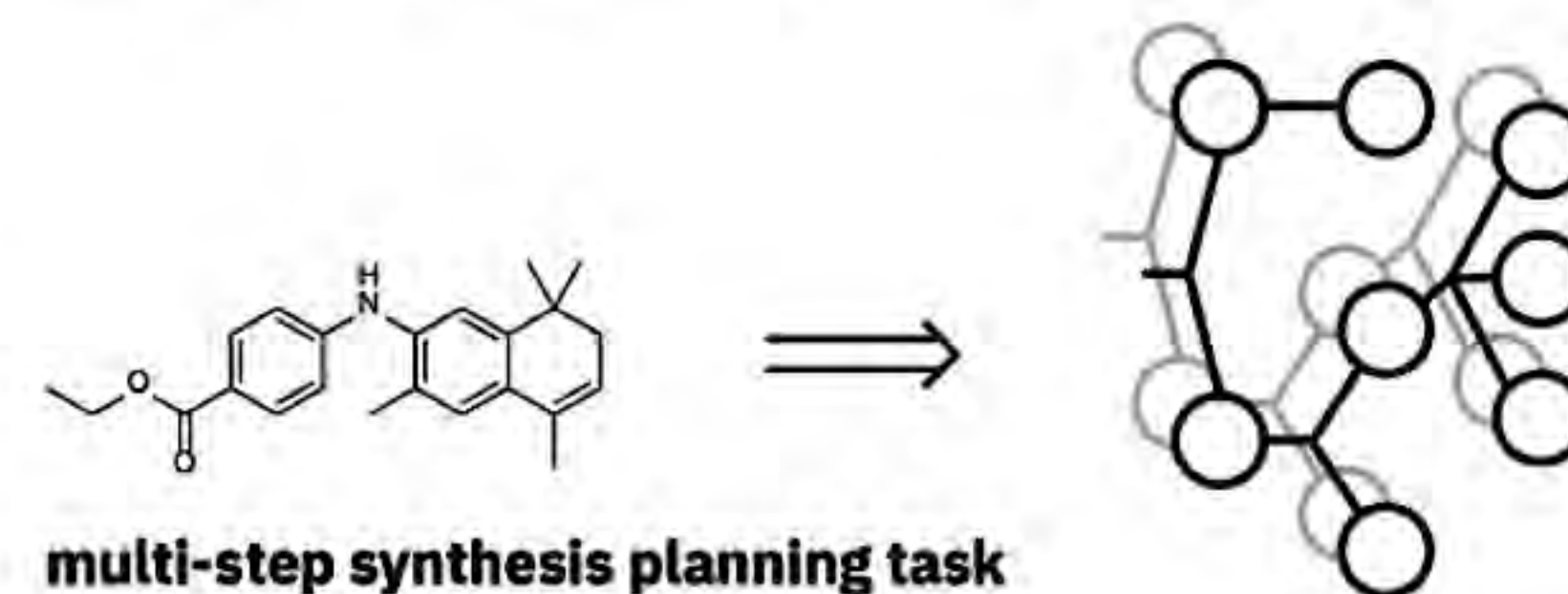
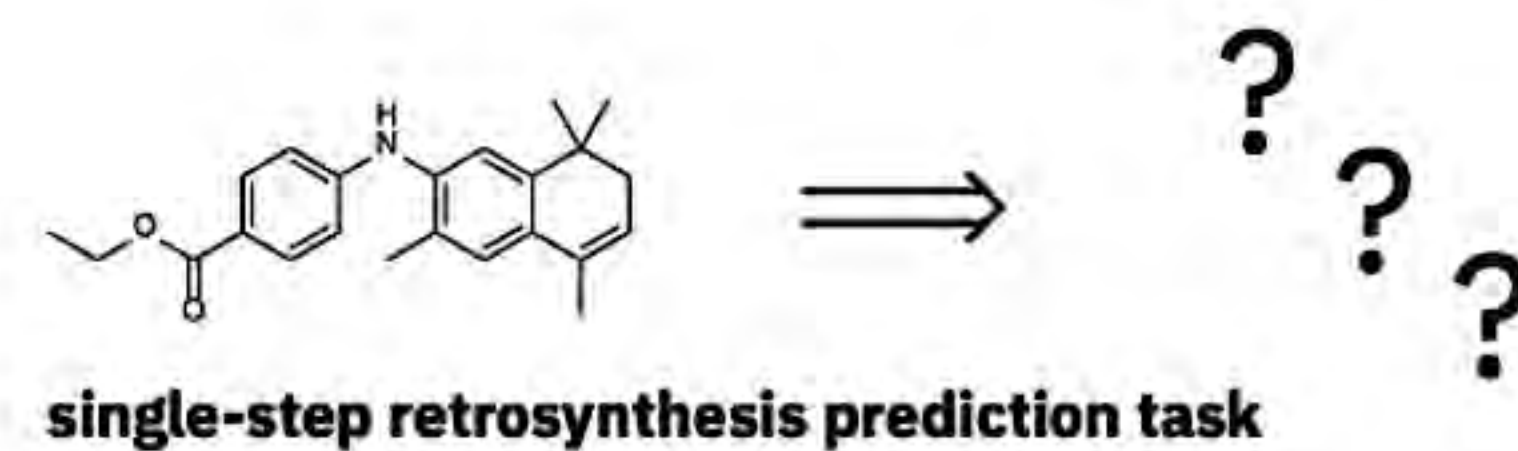
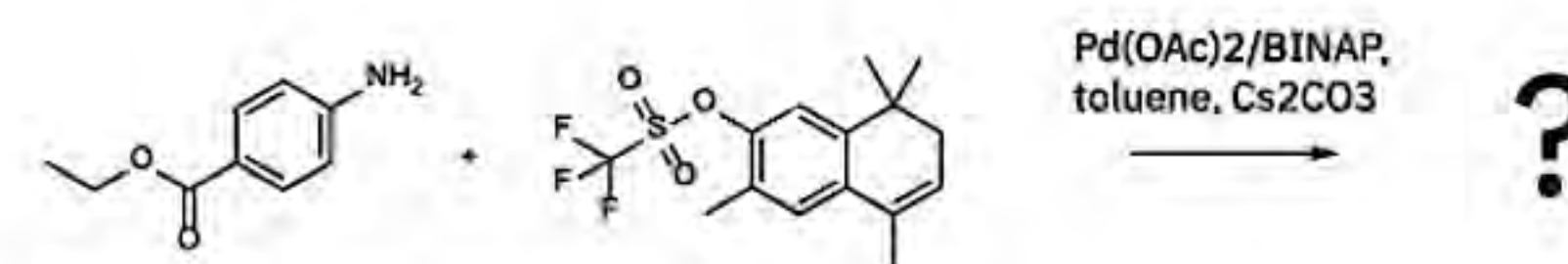
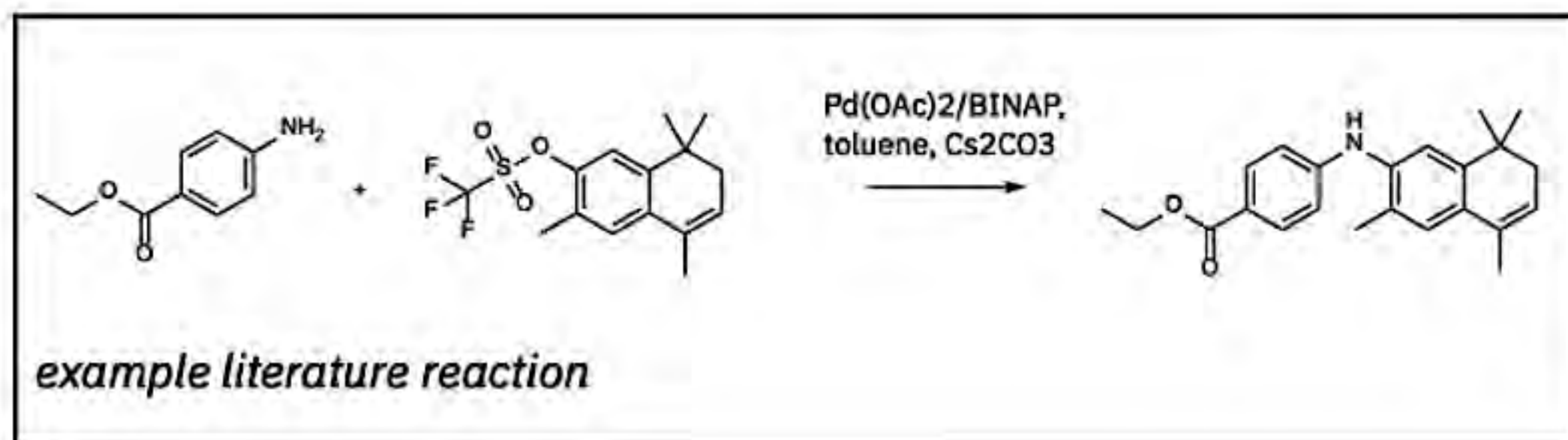
Synthesis planning

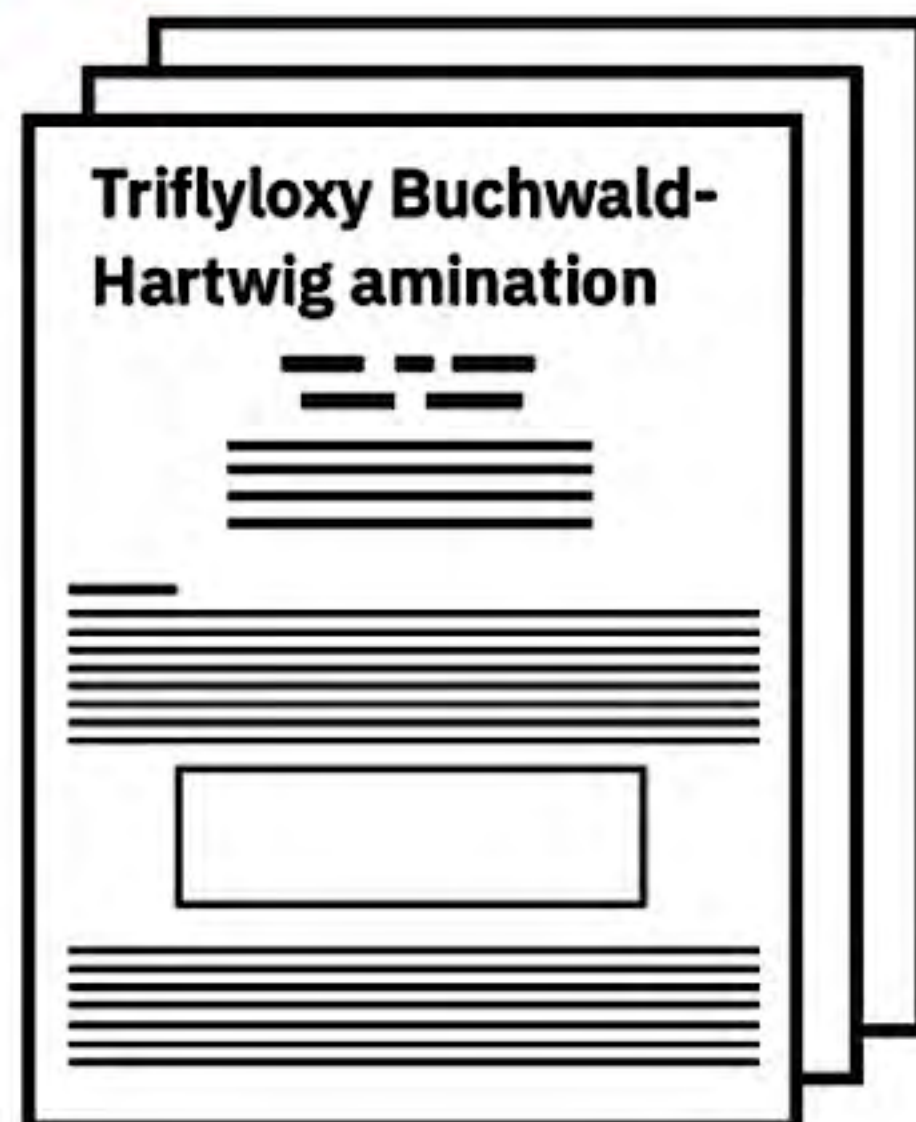


Synthesis optimization



ChemCrow (LLM agents for chemical research)



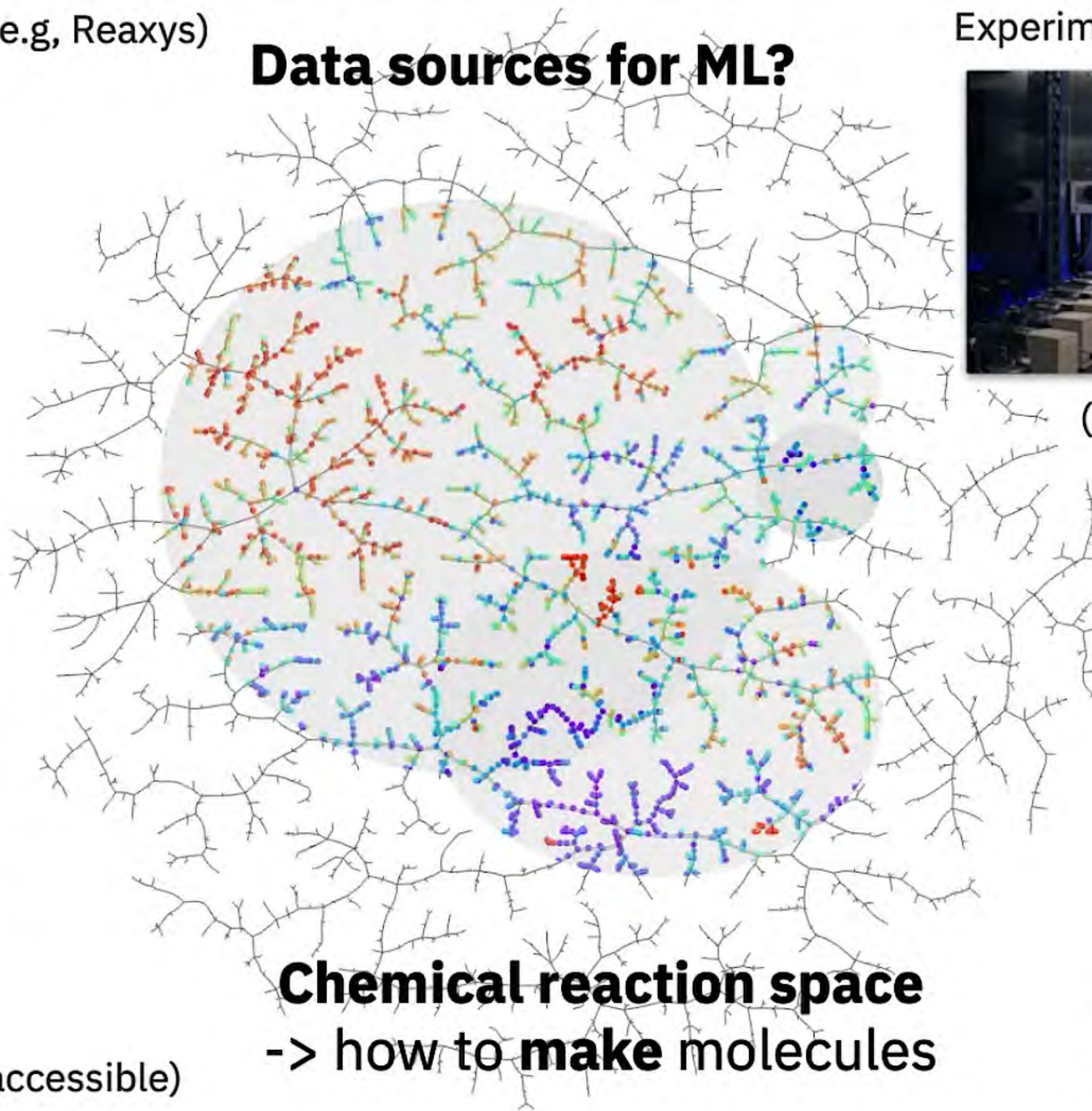


US20030166932A1: General Procedure H
A solution of trifluoromethanesulfonic acid 3,5,8,8-tetramethyl-7,8-dihydronaphthalen-2-yl ester (Compound 35, 0.41 g, 1.2 mmol), Pd(OAc)₂ (0.027 g, 0.12 mmol), BINAP (0.11 g, 0.18 mmol), Cs₂CO₃ (0.56 g, 1.72 mmol), ethyl 4-aminobenzoate (0.25 g, 1.5 mmol) and 5 mL of toluene was flushed with argon for 10 min, then stirred at 100° C. in a sealed tube for 48 h. After the reaction mixture had been cooled to room temperature, the solvent was removed, and the residue was purified by flash column (hexane:ethyl acetate=4:1) to give 0.34 g (80%) of the title compound as a yellowish solid.



Patents (broad, accessible)

Data sources for ML?



Chemical reaction space
-> how to **make** molecules

Experiments ELN/HTE (narrow)



(e.g. ORD, Kearnes et al.)

$$\hat{H}\Psi = E\Psi$$



Simulations (narrow)

US20030166932A1: General Procedure
 H A solution of trifluoromethanesulfonic acid 3,5,8,8-tetramethyl-7,8-dihydronaphthalen-2-yl ester (Compound 35, 0.41 g, 1.2 mmol), Pd(OAc)₂ (0.027 g, 0.12 mmol), BINAP (0.11 g, 0.18 mmol), Cs₂CO₃ (0.56 g, 1.72 mmol), ethyl 4-aminobenzoate (0.25 g, 1.5 mmol) and 5 mL of toluene was flushed with argon for 10 min, then stirred at 100° C. in a sealed tube for 48 h. ...

Daniel Lowe & Roger Sayle



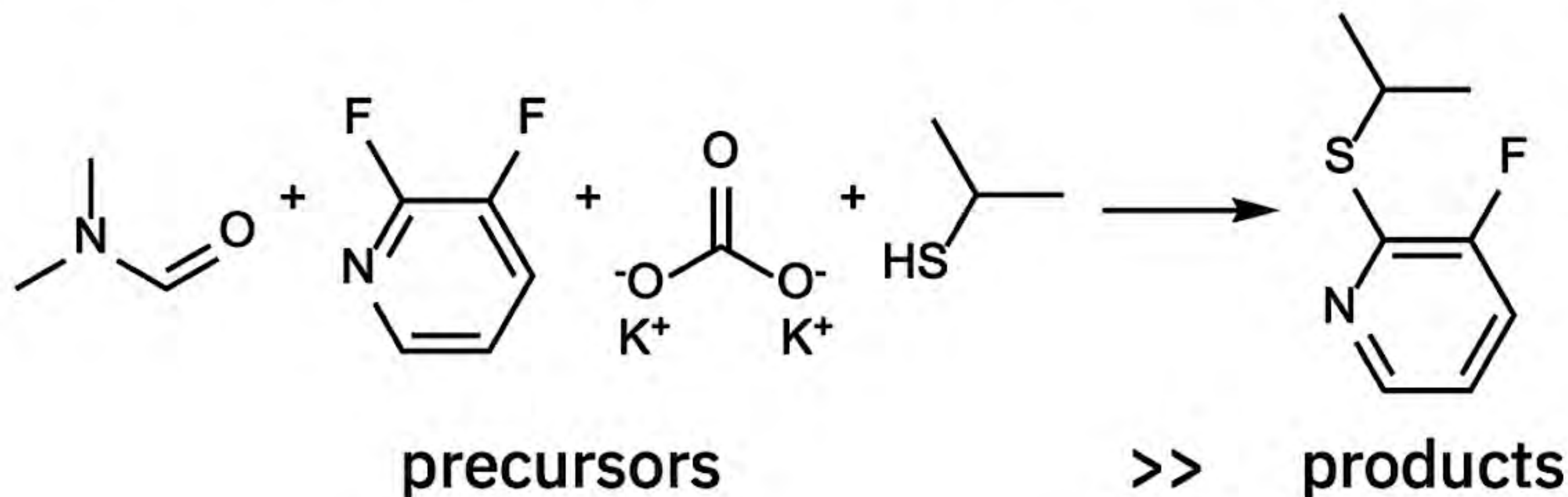
Text-mining

Millions of reaction equations and procedures.



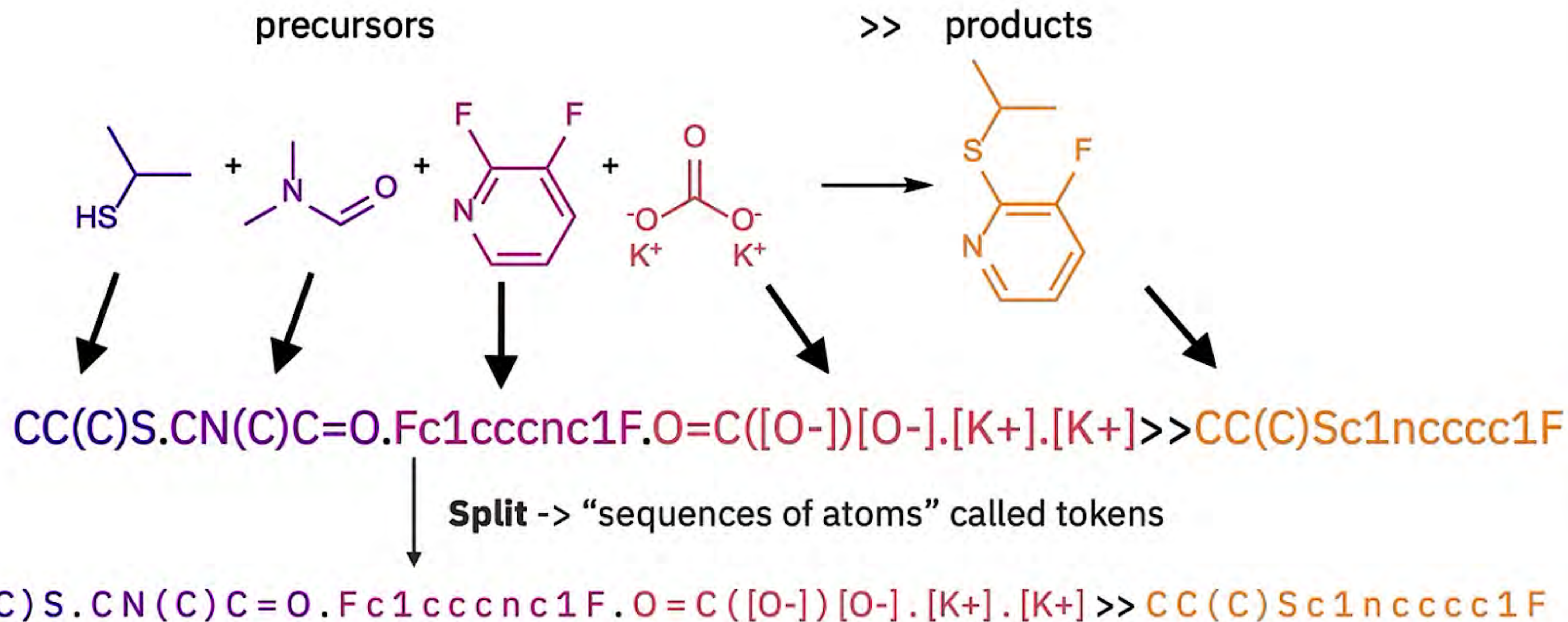
Reaction SMILES

CC(C)S.CN(C)C=O.Fc1ccnc1F.O=C([O-])[O-].[K+].[K+]>>CC(C)Sc1ncccc1F



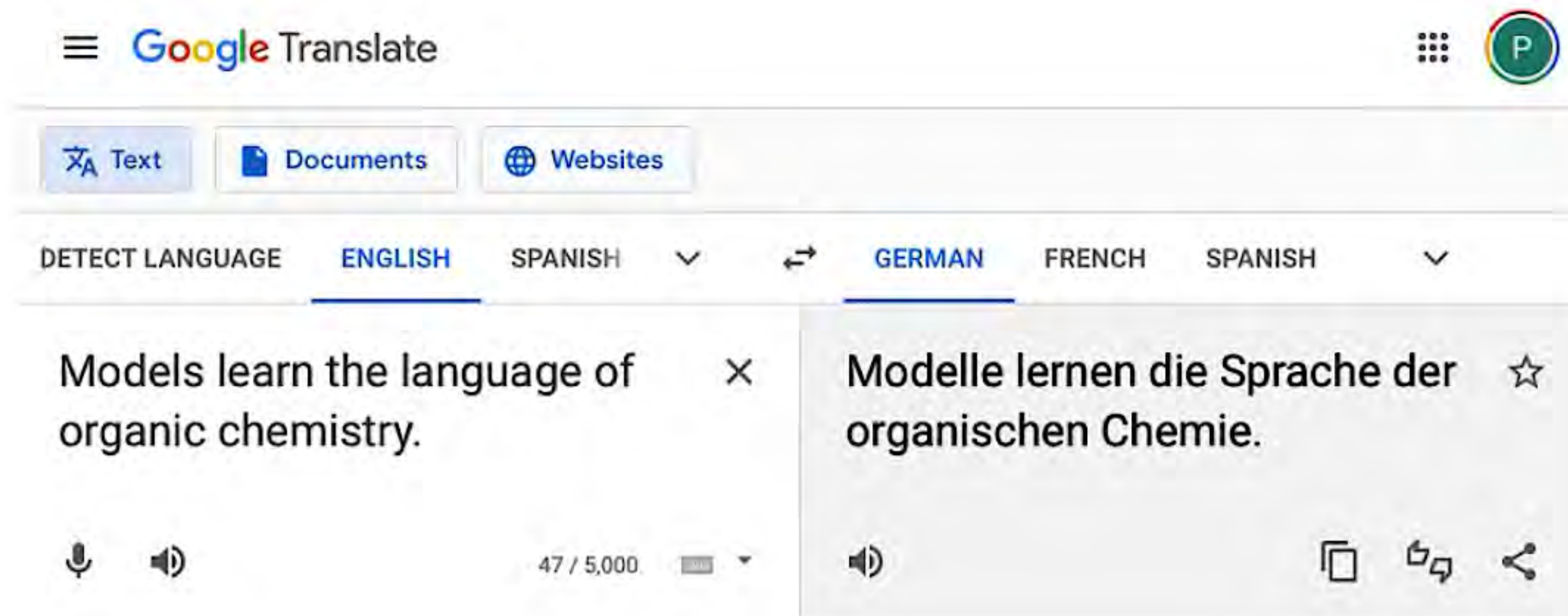
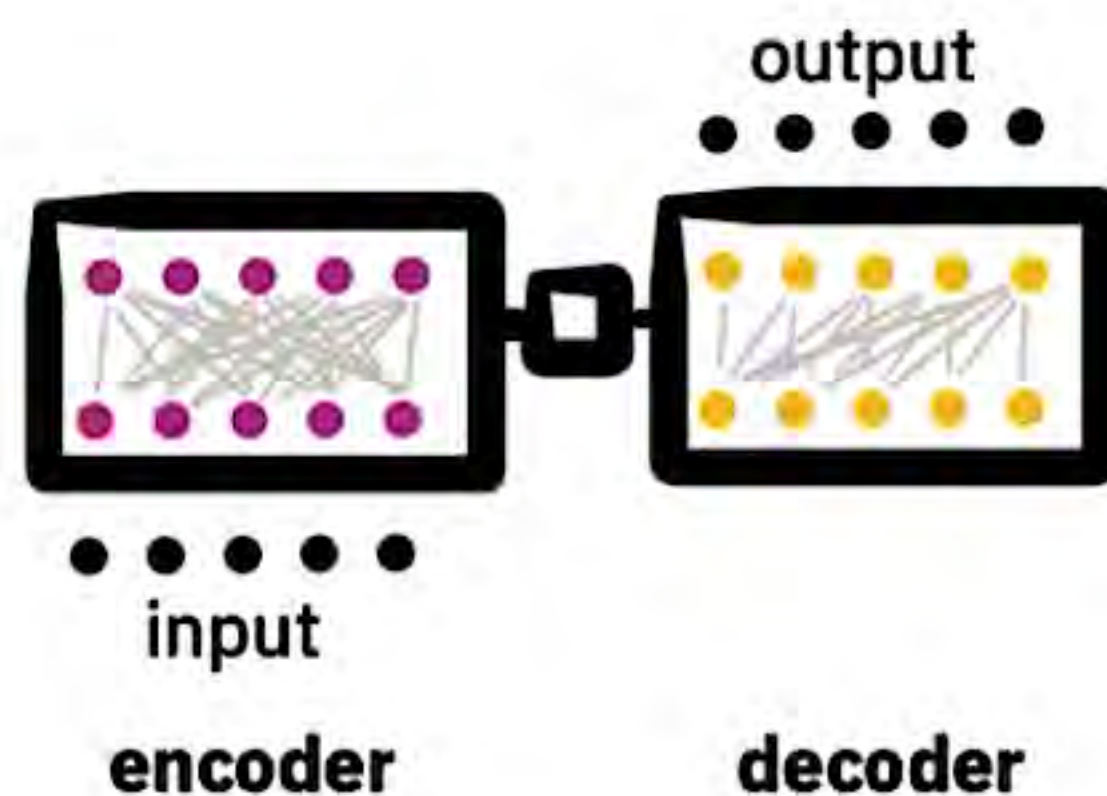
EPFL Atoms as *letters*, molecules as *words*

16

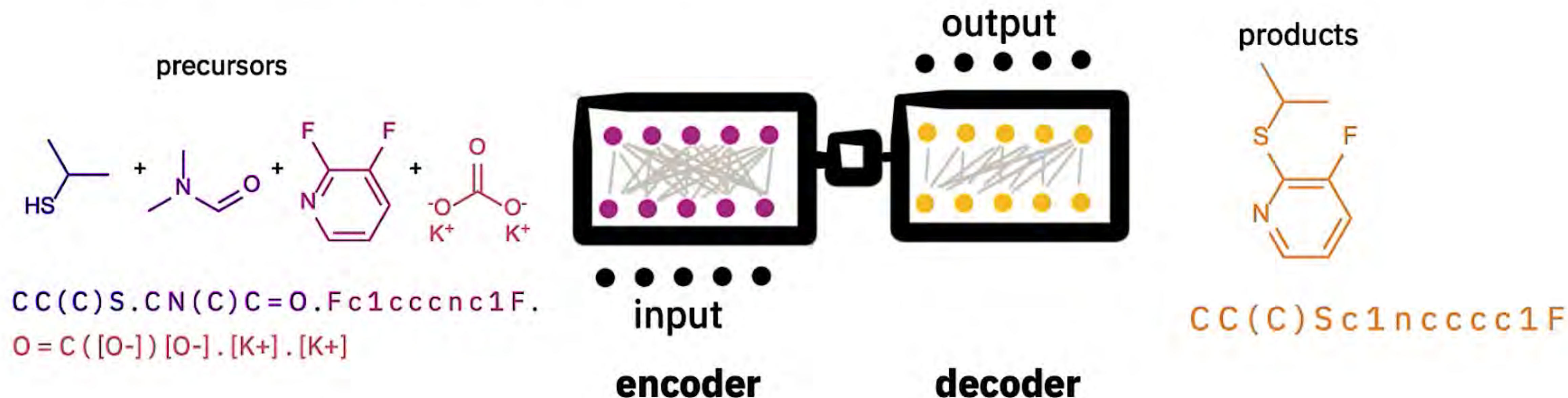


→ Borrow methods developed for human languages

Nam & Kim, arXiv:1612.09529; Liu et al., ACS Centr. Sci. 2017;
Schwaller et al., Chem. Sci, 2018



- Learns translation from examples



- **No rules** integrated / no chemical knowledge
- **Accurate predictions** on unseen reactions (top on benchmarks, back then)
- Better than rule and graph-based approaches

Methods	Top- <i>n</i> accuracy (%)			
	1	3	5	10
USPTO_480k_mixed				
MEGAN (Sacha et al., 2021)	86.3	92.8	94.0	95.4
Molecular Transformer (Schwaller et al., 2019)	88.6	93.5	94.2	94.9
Graph2SMILES (D-GCN) (<i>ours</i>)	90.3	94.0	94.6	95.2
Graph2SMILES (D-GAT) (<i>ours</i>)	90.3	94.0	94.8	95.3
Augmented Transformer (Tetko et al., 2020)	90.6	95.2	95.8	-
Chemformer (Irwin et al., 2021)	91.3	95.7	96.2	94.0

Transformer-decoder

**Same architecture as
Molecular Transformer**

PERMUTATION INVARIANT GRAPH-TO-SEQUENCE MODEL FOR TEMPLATE-FREE RETROSYNTHESIS AND REACTION PREDICTION

Graph2SMILES

Zhengkai Tu^{1,2} and Connor W. Coley^{1,3}

State-of-the-art augmented NLP transformer models for direct and single-step retrosynthesis

Igor V. Tetko , Pavel Karpov, Ruud Van Deursen & Guillaume Godin 

Augmented Transformer

Chemformer: a pre-trained transformer for computational chemistry

Ross Irwin¹, Spyridon Dimitriadis^{1,2}, Jiazhen He¹ and Esben Jannik Bjerrum^{3,1} 

Molecule Edit Graph Attention Network: Modeling Chemical Reactions as Sequences of Graph Edits

Mikołaj Sacha, Mikołaj Błaż, Piotr Byrski, Paweł Dąbrowski-Tumański, Mikołaj Chromiński, Rafał Łoska, Paweł Włodarczyk-Pruszyński, and Stanisław Jastrzębski*

MEGAN

Target molecule



Ages
6+
Easter Chick
30 pcs



Known (commercially available) building blocks



Szymkuć et al., ACIE 2016
Grzybowski et al., Chem 2018
Mikulak-Klucznik et al., Nature, 2020

- expert rule-based
- “wide & deep” search
- simulations, ML scoring

Segler & Waller
Nature, 2018

- template-based (reactants)
- Monte-Carlo tree search
- step-scoring NN

AiZynthFinder
Thakkar et al., Chem. Sci., 2020
Genheden et al., J. Cheminf., 2020

- template-based (reactants)
- Monte-Carlo tree search
- step-scoring NN
- interactive mode

And more...

Steps to
construct the
target

1



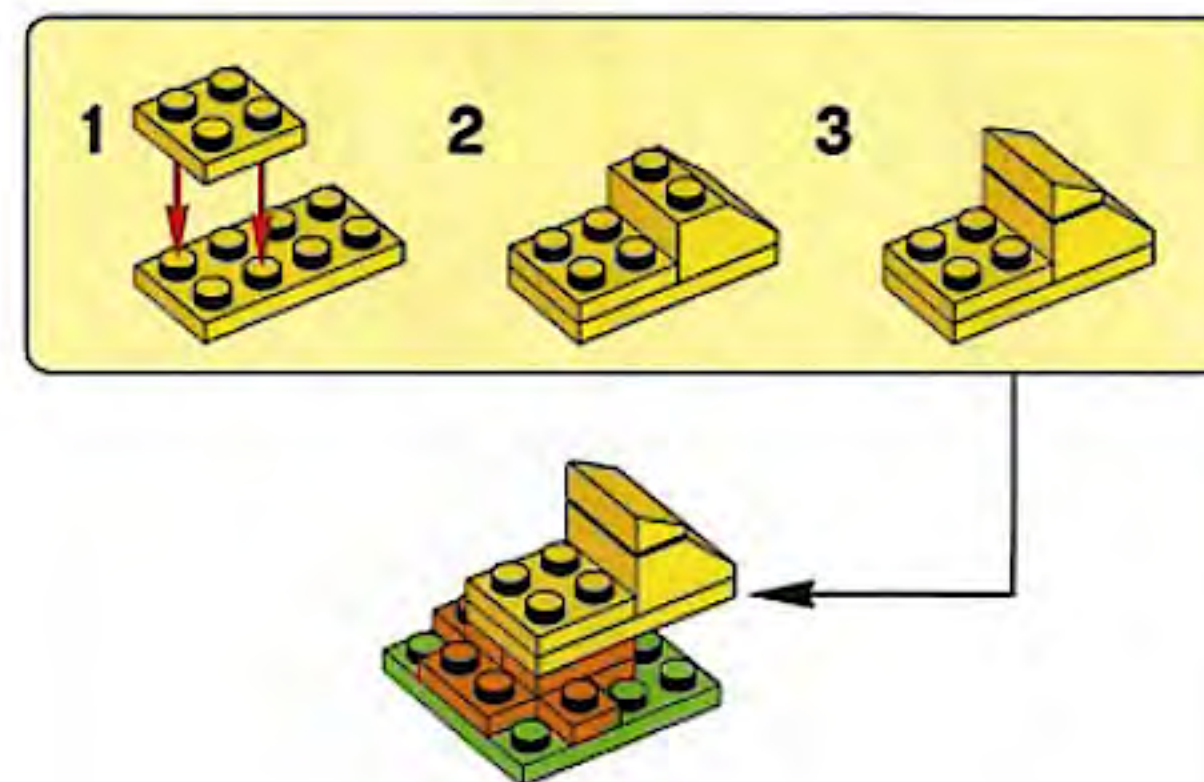
2



3



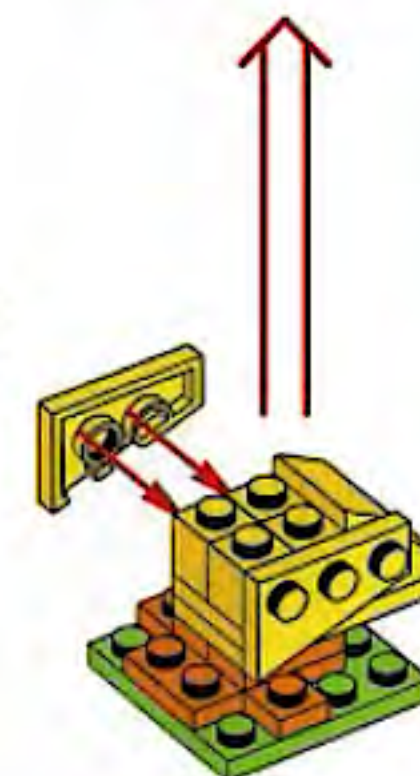
4



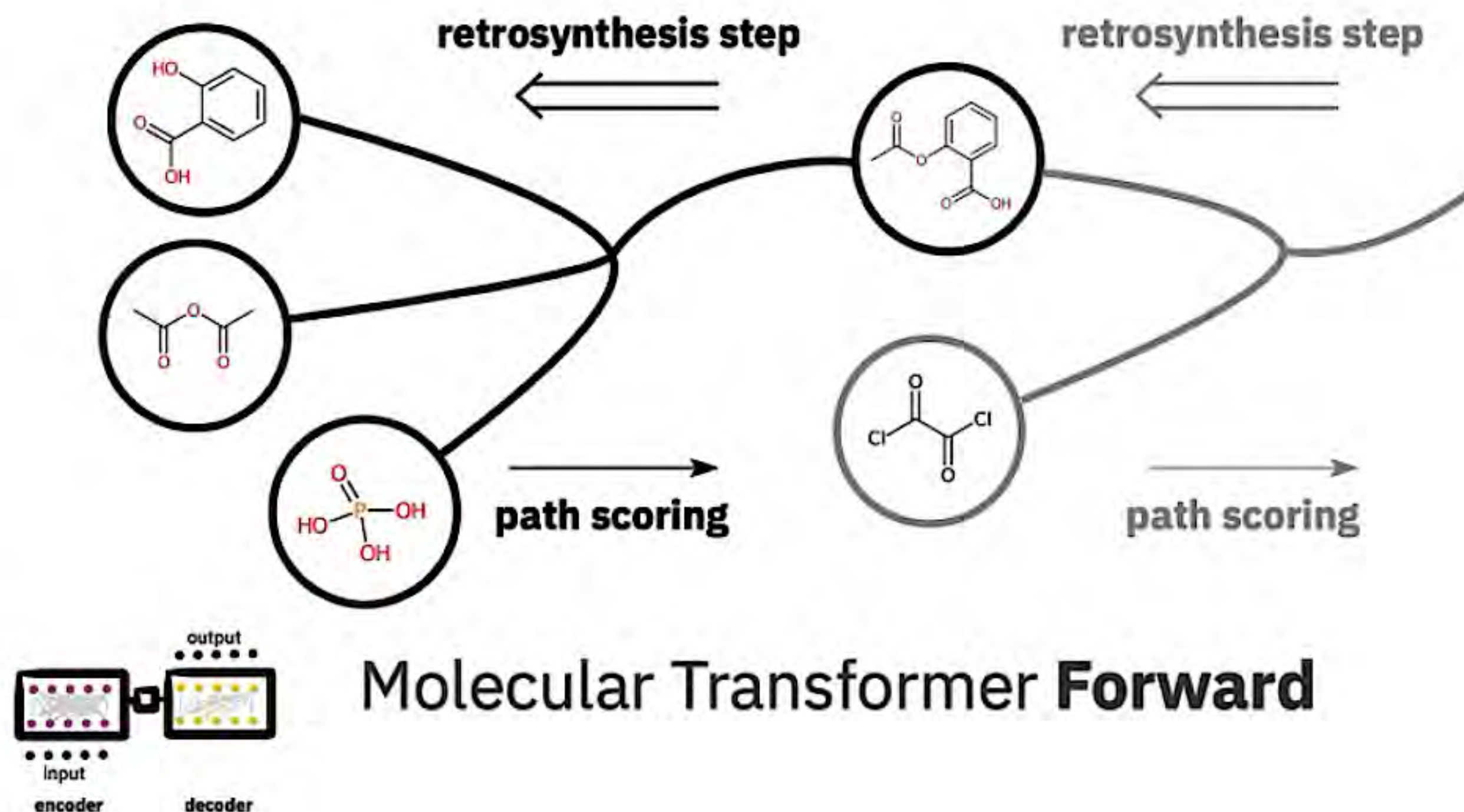
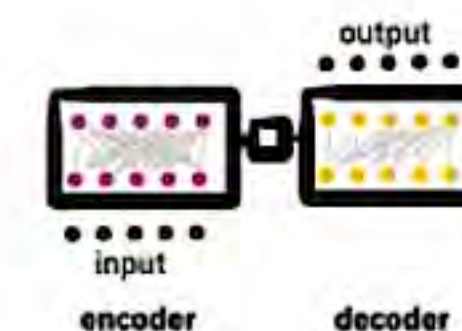
5



6



Lego analogy:
Amol Thakkar

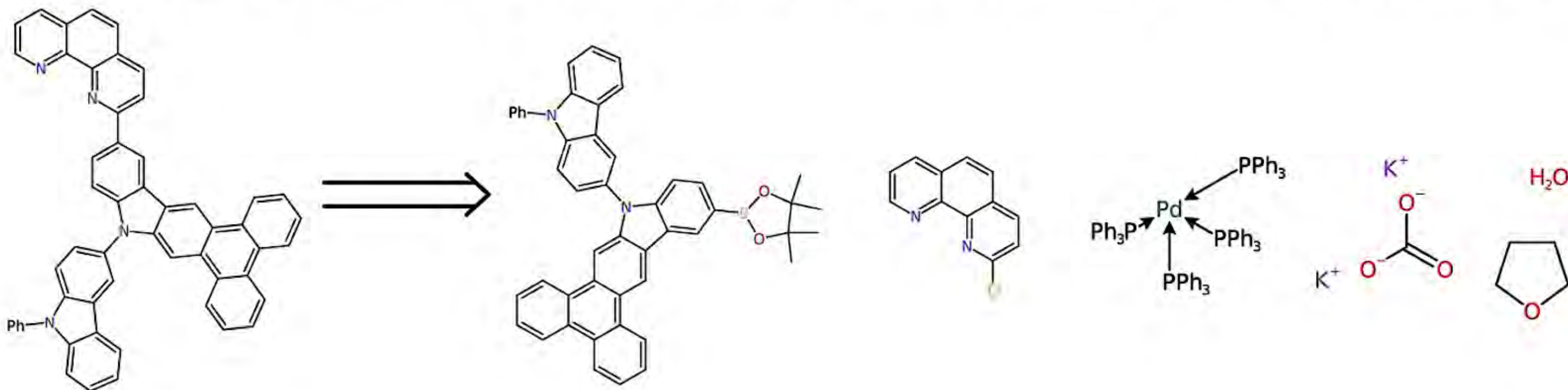
Molecular Transformer **Retro**

Predicting retrosynthetic pathways using a combined linguistic model and hyper-graph exploration strategy

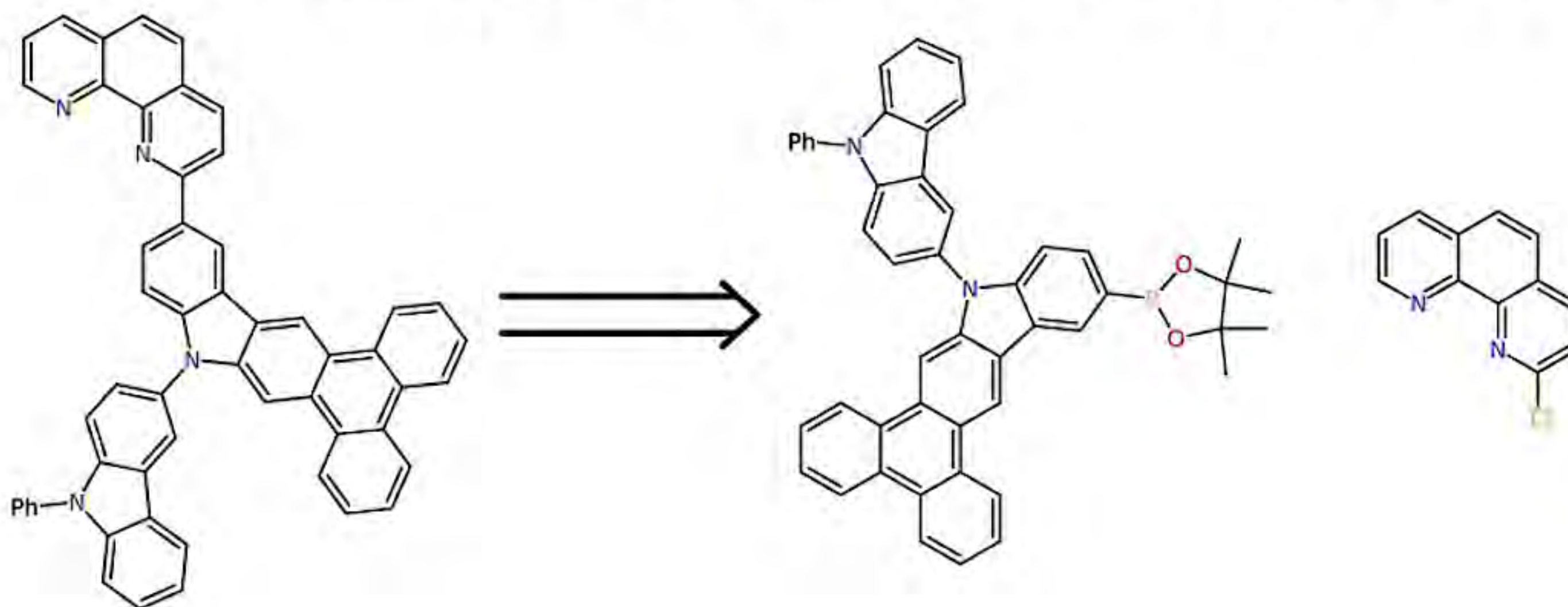
Philippe Schwaller, Riccardo Petraglia, Valerio Zullo, Vishnu H Nair, Rico Andreas Haeuselmann, Riccardo Pisoni, Costas Bekas, Anna Iuliano, Teodoro Laino

Chem. Sci., 2020, Advance Article

<https://doi.org/10.1039/C9SC05704H>



Reactants prediction (atom-mapping dependent)

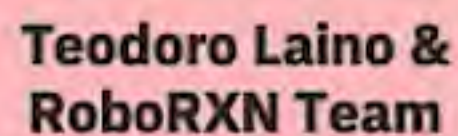


Rule-based approaches:

- Segler et al. Nature, 555, 604–61
- Coley et al. Science 365 (6453)
- Genheden et al. J Cheminf. 12 (1), 1-9
- Thakkar et al. Chem. Sci. 11 (1), 154-168

and other SMILES-based approaches:

- Liu et al. ACS Cent. Sci. 3 (10), 1103-1113
- Tetko et al. Nature Comm. 11 (1), 1-11



Automation

Remote AI-driven synthesis

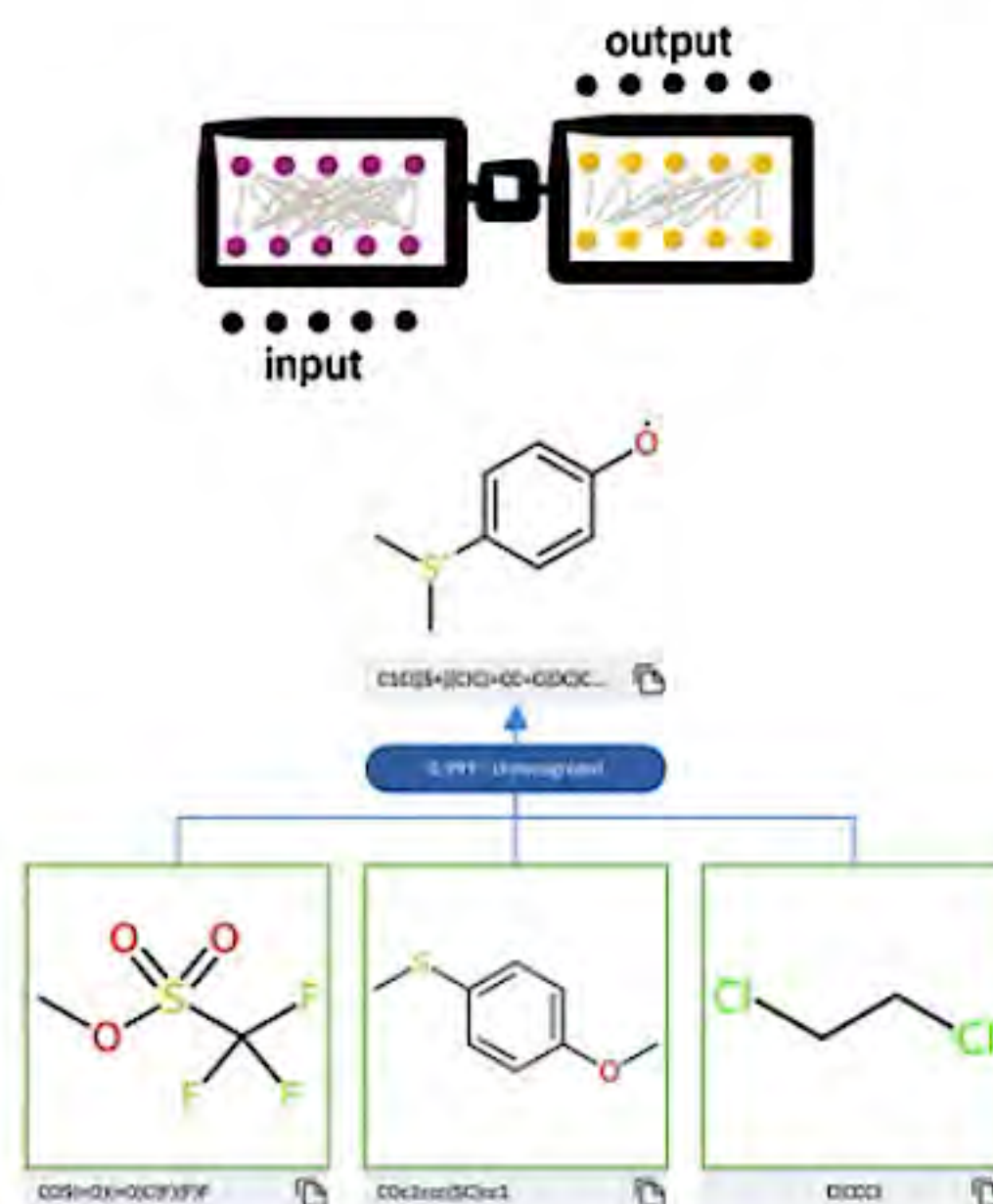
AI

Cloud-connected synthesis platform

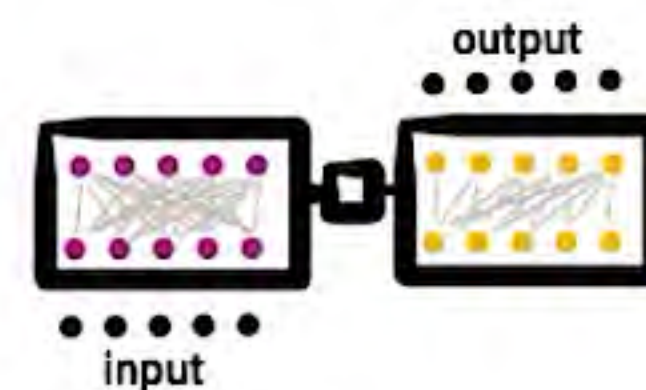


Chemists at home

- design of target molecule
- creative process

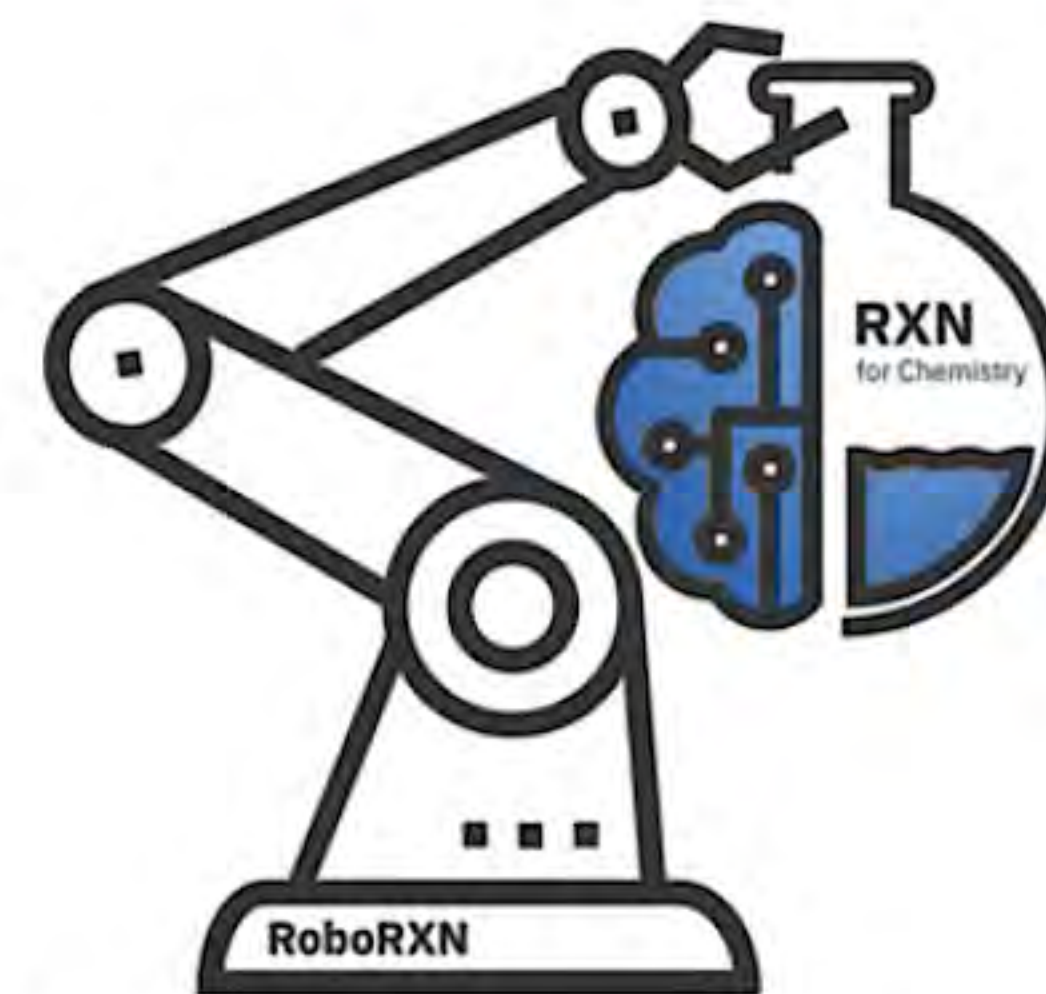


Synthesis planning



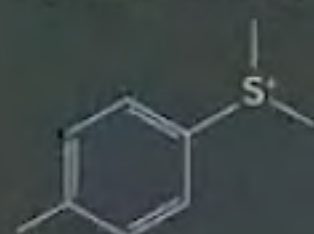
- Add: 1mmol 4-methoxythioanisole
- Add: 10 ml 1,2-dichloroethane
- Add: 1mmol methyl trifluoromethanesulfonate
- Stir: at 25°C for 1 day
- Filter: keep filtrate
- Concentrate

Recipe formulation



Execution on robot

Synthesizing new molecule



Started: Nov 30 2020, 6:49am PT

Live from IBM RoboRXN

Action 2

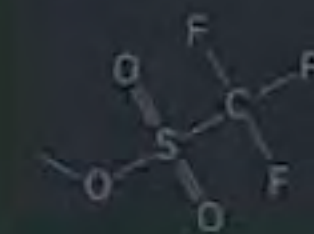
Overview

Adding $C_2H_3F_3O_3S$

In this action, the molecule methyl trifluoromethane sulfonate is added to Reactor 2.

Methyl trifluoromethane sulfonate
 $C_2H_3F_3O_3S$

20 30



Methyl trifluoromethane sulfonate is a brown liquid. Insoluble in water. This material is a very reactive methylating agent, also known as methyl triflate.

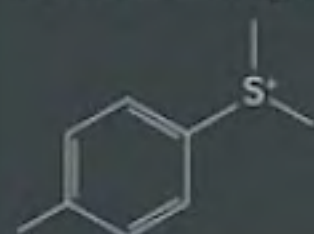
NOW

10 ml of reagent containing methyl trifluoromethane sulfonate is being moved from Vial 61 and added to Reactor 2.

Position of the robot arm
Moving to Vial 61



Synthesizing new molecule



Started: Nov 30 2020, 6:49am PT

Live from IBM RoboRXN

Action 2

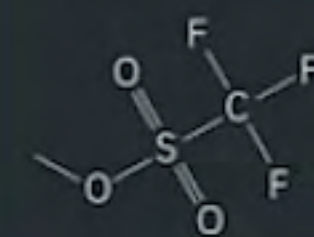
Overview

Adding $C_2H_3F_3O_3S$

In this action, the molecule methyl trifluoromethane sulfonate is added to Reactor 2.

Methyl trifluoromethane sulfonate
 $C_2H_3F_3O_3S$

2D 3D



Methyl trifluoromethane sulfonate is a brown liquid. Insoluble in water. This material is a very reactive methylating agent, also known as methyl triflate.

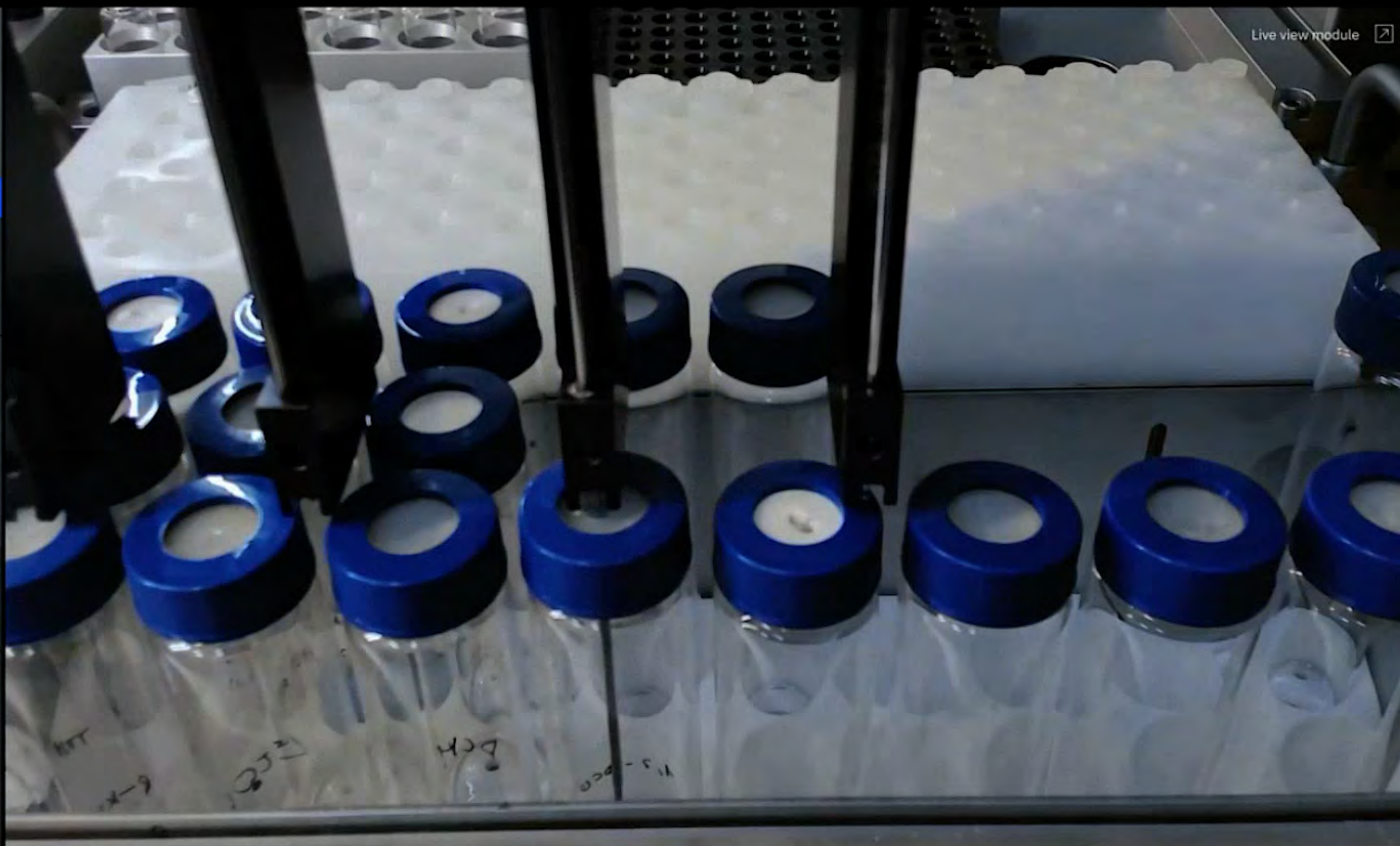
NOW

10 ml of reagent containing methyl trifluoromethane sulfonate is being moved from Vial 61 and added to Reactor 2.

Position of the robot arm
Extracting from vial 61



Live view module



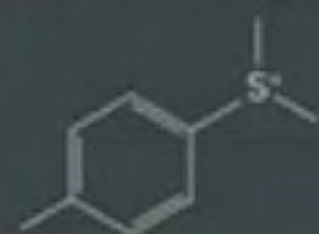
2 Adding $C_2H_3F_3O_3S$

00:06:07



LIVE

Synthesizing new molecule



Started: Nov 30 2020, 6:49am PT

Live from IBM RoboRXN

Action 2

Overview

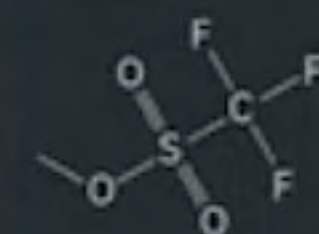
Adding $C_2H_3F_3O_3S$

In this action, the molecule methyl trifluoromethane sulfonate is added to [Reactor 2](#).

Methyl trifluoromethane

$C_2H_3F_3O_3S$

20 30



Methyl trifluoromethane sulfonate is a brown liquid. Insoluble in water. This material is a very reactive methylating agent, also known as methyl triflate.

NOW

10 ml of [reagent](#) containing methyl trifluoromethane sulfonate is being moved from [Vial 61](#) and added to [Reactor 2](#).

Position of the robot arm

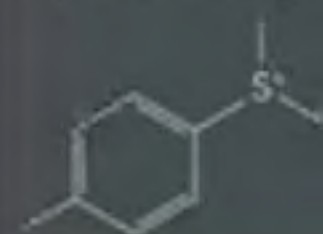
Inserting into reactor 2



Live view module



Synthesizing new molecule



Started: Nov 30 2020, 8:49am PT

Live from IBM RoboRXN

Action 2

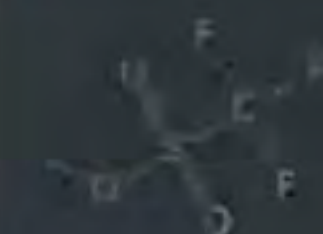
Overview

Adding $C_2H_5F_3O_2S$

In this action, the molecule methyl trifluoromethane sulfonate is added to [Reactor 2](#).

Methyl trifluoromethane sulfonate
 $C_2H_5F_3O_2S$

300 30



Methyl trifluoromethane sulfonate is a brown liquid. Insoluble in water. This material is a very reactive methylating agent, also known as methyl triflate.

NOW

10 ml of [reagent](#) containing methyl trifluoromethane sulfonate is being moved from [Vial 61](#) and added to [Reactor 2](#).

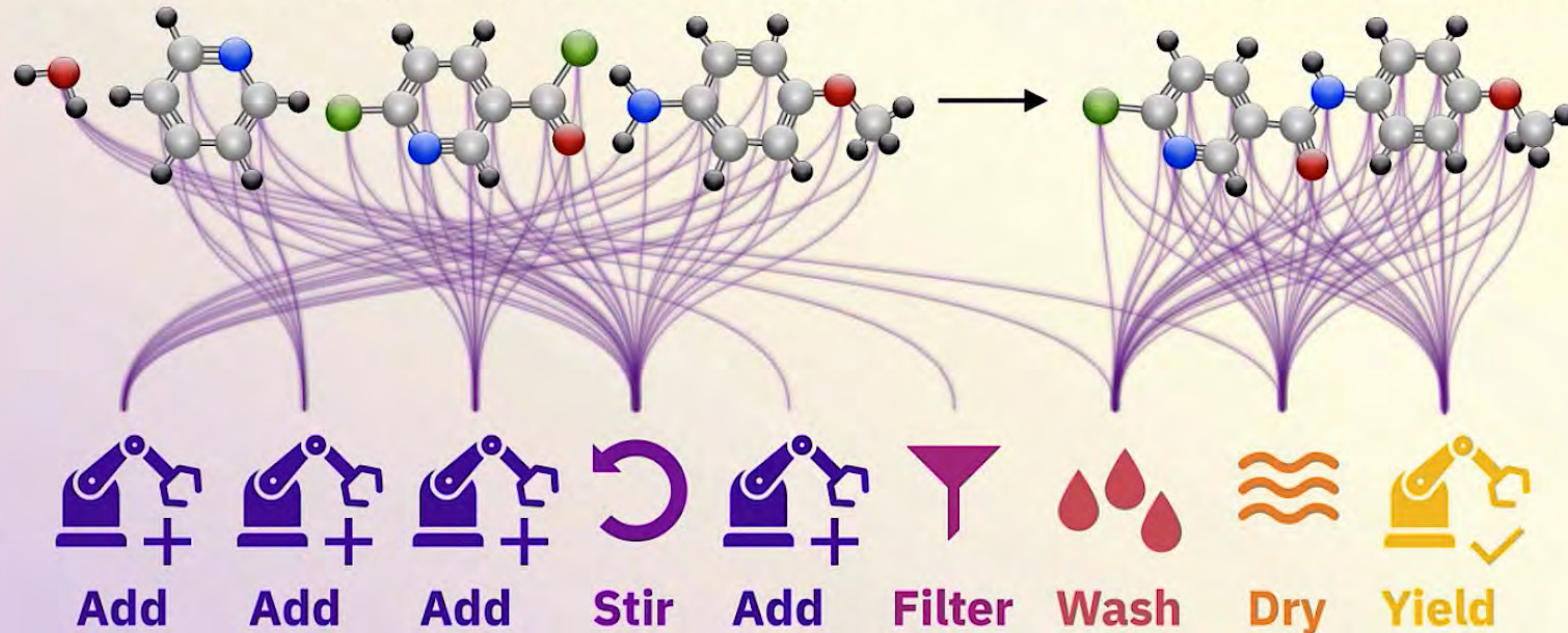
Position of the robot arm
Moving to Vial 61



Live view module



O . c1ccncc1 . O=C(Cl)c1ccc(Cl)nc1 . COc1ccc(N)cc1 >> COc1ccc(NC(=O)c2ccc(Cl)nc2)cc1



SMILES-2-ACTIONS

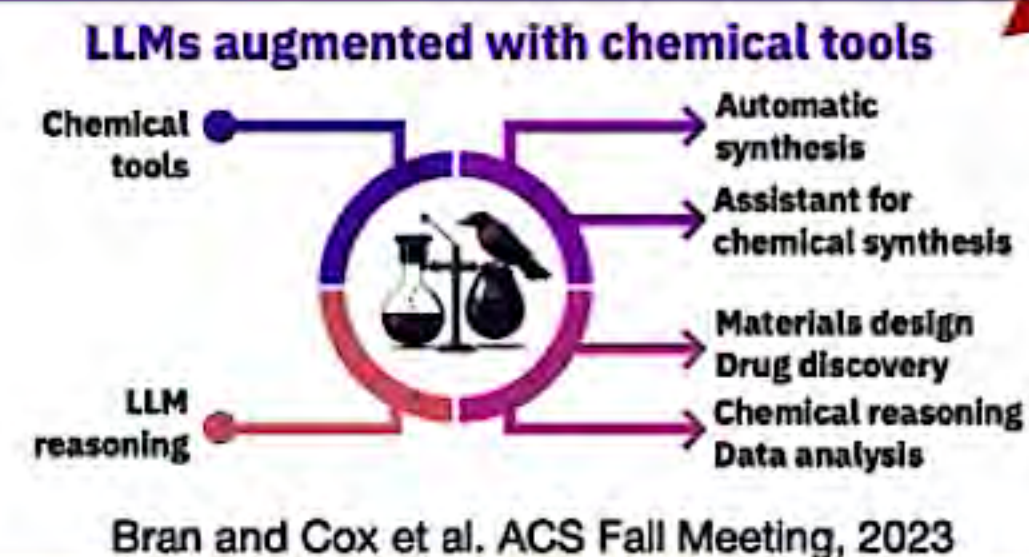
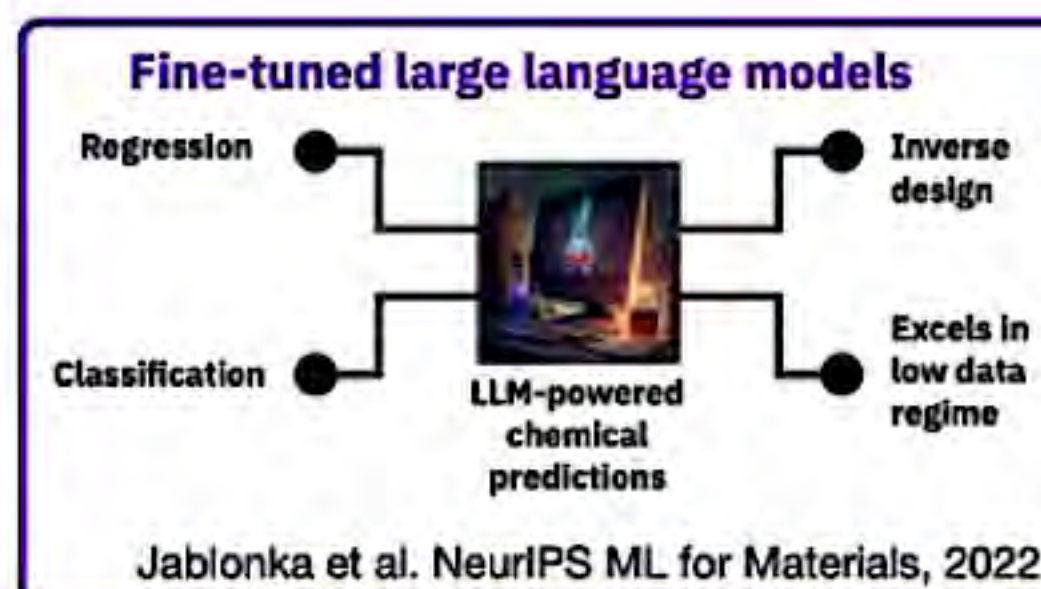
Vaucher, A.C., Schwaller, P., Geluykens, J. *et al.* Inferring experimental procedures from text-based representations of chemical reactions. *Nat Commun* **12**, 2573 (2021).

Vaucher, A.C., Zipoli, F., Geluykens, J. *et al.* Automated extraction of chemical synthesis actions from experimental procedures. *Nat Commun* **11**, 3601 (2020).



Alain Vaucher

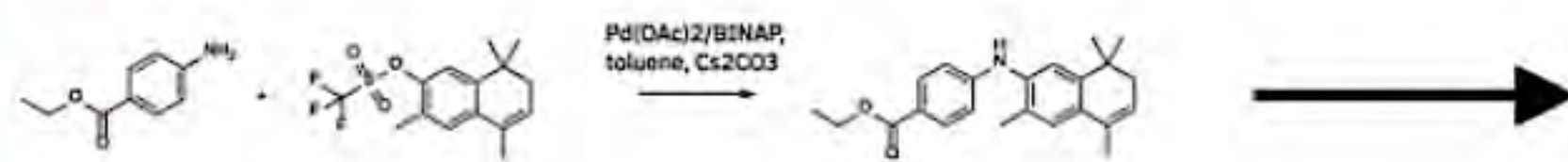
Other examples:
- Darwin
General task solvers (open models)



Other examples:
- Clairify (Skreta et al.)
- Coscientist (Boiko et al.)

Multiple modalities

Reaction to synthesis procedure

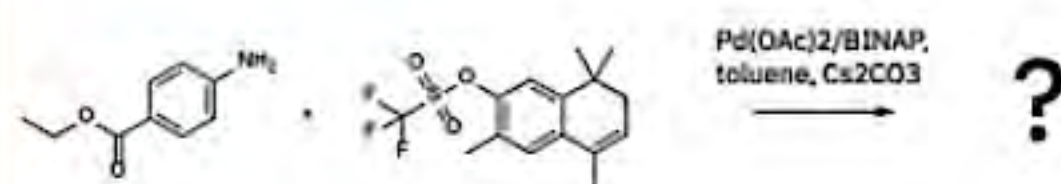


Vaucher et al. *Nature Comm.* 2020
Vaucher et al. *Nature Comm.* 2021

1. MAKESOLUTION with trifluoromethanesulfonic acid 3,5,8,8-tetramethyl-7,8-dihydronaphthalen-2-yl ester (0.41 g, 1.2 mmol) and Pd(OAc)₂ (0.027 g, 0.12 mmol) and BINAP (0.11 g, 0.18 mmol) and Cs₂CO₃ (0.56 g, 1.72 mmol) and ethyl 4-aminobenzoate (0.25 g, 1.5 mmol) and toluene (5 mL)
2. ADD SLN
3. STIR for 48 hours at 100° C
4. SETTEMPERATURE room temperature
5. CONCENTRATE
6. PURIFY
7. YIELD product

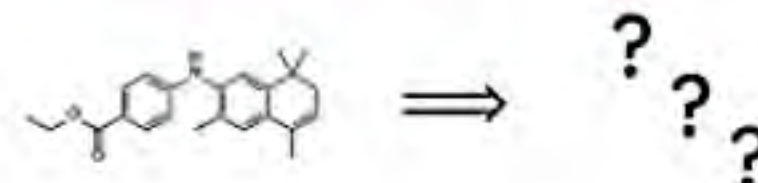
Single modality

Reaction prediction



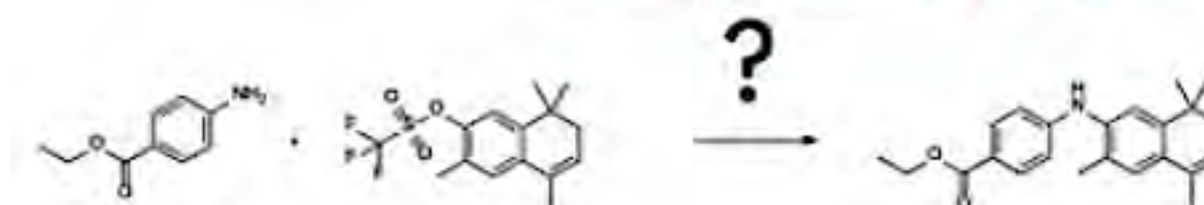
Schwaller et al. *ACS Cent. Sci.* 2019
Pesciullesi et al. *Nature Comm.* 2020

Retrosynthetic planning



Schwaller et al. *Chem. Sci.*, 2020
Thakkar et al. *ACS Cent. Sci.* 2023

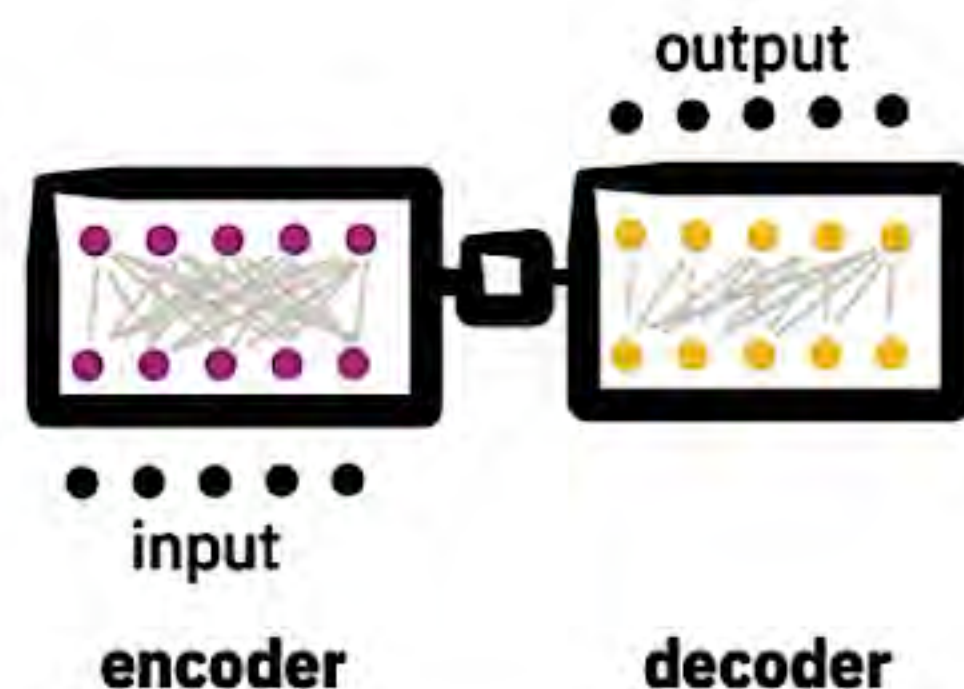
Reaction condition prediction



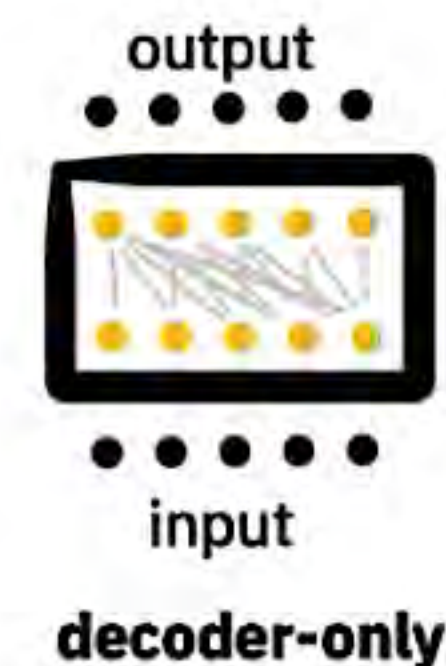
Gao et al. *ACS Cent. Sci.* 2018
Schilter et al. *ACS Fall Meeting* 2023

So far, all I showed you had max ~20M parameters, trained on 1 consumer GPU in ~2 days.

Attention is all you need
(Vaswani et al., 2017)



GPT
(Radford et al., 2018)



- Decoder-only models are trained to **predict the next “word”** given all previous words. → GPT3, ChatGPT, GPT4
- Philippe should
- Philippe should **prepare**
- Philippe should prepare **slides**
- ...
- Philippe should prepare slides for the SPREE **seminar**.



Not many tools are used

- Too many of them (100+)
- They live in isolation
- Hard to set-up
- Non-intuitive to use

There's a gap between computational and experimental fields!



Aim: Bridge the gap

- ✓ Continue making useful tools
- ✓ Make them usable
-> also for non-experts

- Large language models (GPT4/ChatGPT) are **bad at chemistry**
- There are **excellent chemistry tools** (but in isolation)

So can what do we do? → ChemCrow

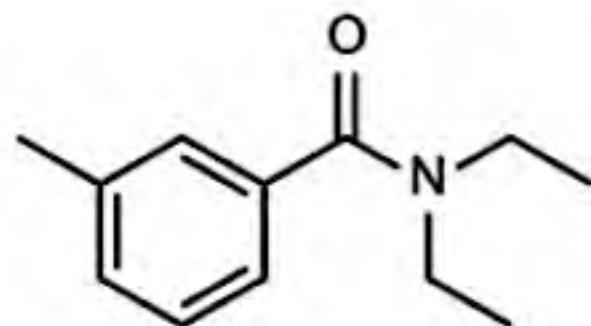
What **tools** for ChemCrow?



[Andres M. Bran](#), [Sam Cox](#), [Oliver Schilter](#), [Carlo Baldassari](#), [Andrew D. White](#) ✉ & [Philippe Schwaller](#) ✉

[Nature Machine Intelligence](#) **6**, 525–535 (2024) | [Cite this article](#)

Molecule tools



- SMILES to Weight
- SMILES to Price
- SMILES to CAS
- Similarity
- Modify Mol
- Func Groups
- Patent Check
- Name to SMILES
- Safety Assessment
- Explosive Check

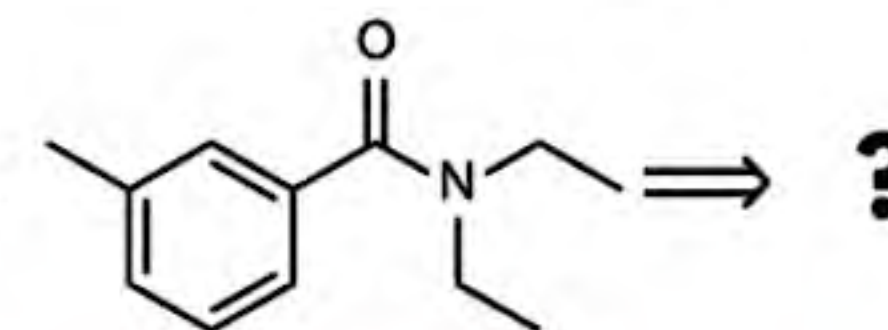


General tools

- Literature Search
- Web Search
- Code interpreter
- Human expert



- RXN to Name
- RXN Predict
- Synth Plan



Safety tools

- RDKit: Open-source cheminformatics. <https://www.rdkit.org>
- PubChem Compound Database <https://pubchem.ncbi.nlm.nih.gov/>
- Synspace: <https://github.com/whitead/synspace>
- Molbloom: <https://github.com/whitead/molbloom>
- Chem-space: <https://chem-space.com/>
- paper-qa: <https://github.com/whitead/paper-qa/tree/main>
- IBM RXN for Chemistry: <https://github.com/rxn4chemistry/rxn4chemistry/>
- ClinTox: Substance Toxicity Dataset: <https://www.clintox.org/>
- GHS Classification: <https://pubchem.ncbi.nlm.nih.gov/ghs/>

Reaction tools

Expert-designed chemistry tools

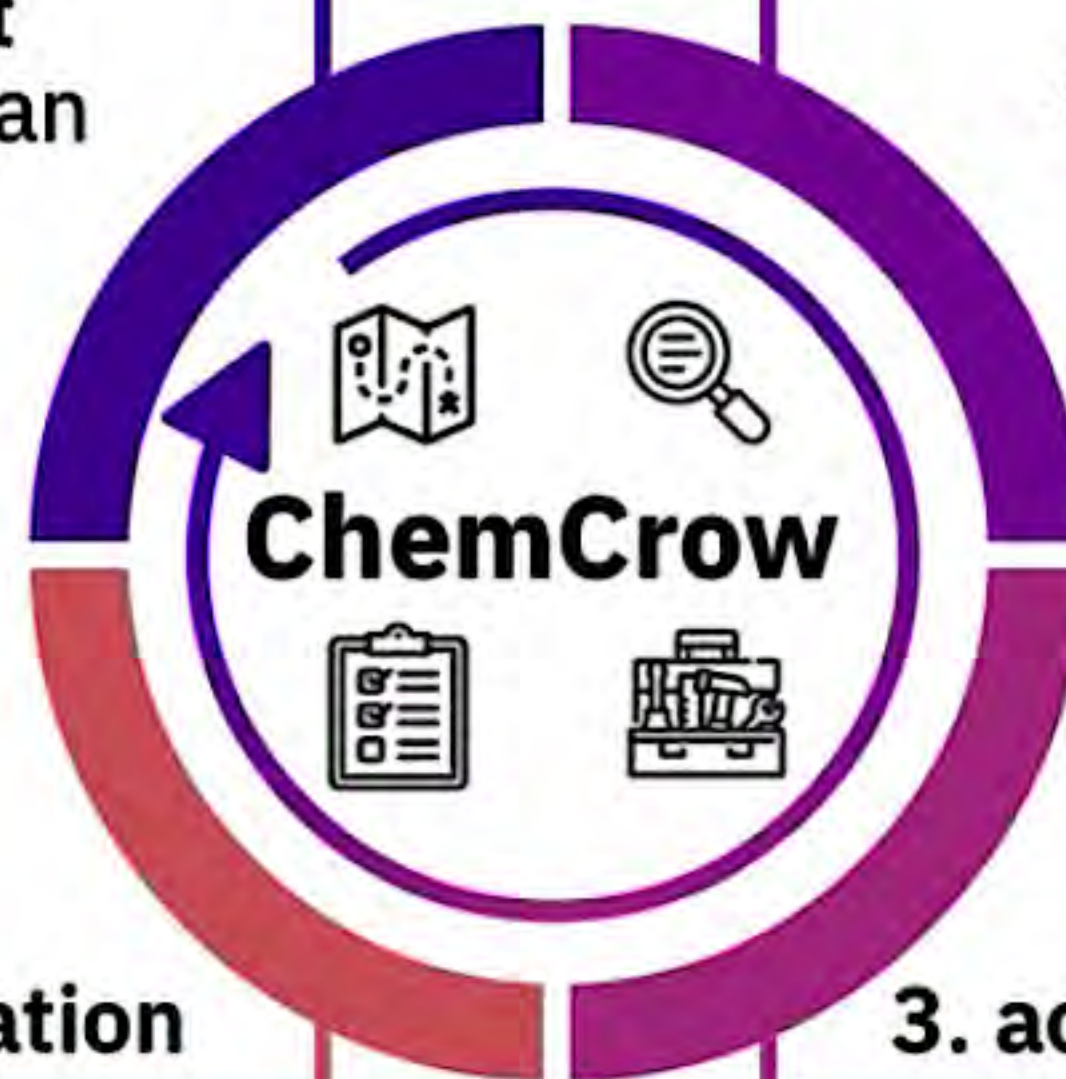


Example prompt:
Plan and execute
the synthesis of an
insect repellent.

User-defined scientific tasks

1. thought
reason, plan

2. action
select tool



analyse
4. observation

use tool
3. action input



- Yao, S.; React: Synergizing reasoning and acting in language models. *arXiv preprint arXiv:2210.03629* **2022**,
- Karpas, E. et al. MRKL Systems: A modular, neuro-symbolic architecture that combines large language models, external knowledge sources and discrete reasoning. *arXiv preprint arXiv:2205.00445* **2022**,

Task: Plan and execute the synthesis of an insect repellent

Thought: I need to know which insect repellent to synthesize. I will do a literature search to find a common insect repellent.

Action: LitSearch

Action input: common insect repellent

Observation: DEET, picaridin, ...

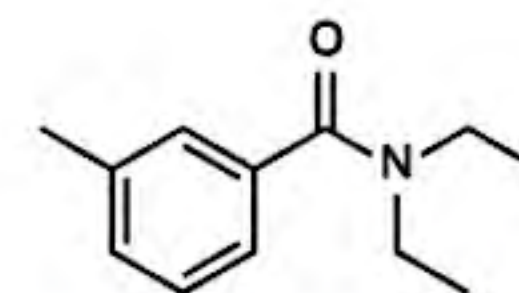


Thought: I will choose DEET as the insect repellent to synthesize. First I need to get the SMILES of DEET.

Action: Name2SMILES

Action input: DEET

Observation: CCN(CC)C(=O)c1ccccc(C)c1



Thought: Now that I have the SMILES of DEET, I will plan the synthesis.

Action: RXNPlanner

Action input: CCN(CC)C(=O)c1cccc(C)c1

Observation:

To make CCN(CC)C(=O)c1cccc(C)c1, follow these steps:

1. Add 0.3 mL of diethylamine to the reaction vessel.
2. Add 15 mL of dichloromethane to the reaction vessel.
3. Add 3.2 mmol of 3-methylbenzoyl chloride to the reaction vessel at 0°C.
4. Stir the mixture at 25°C for 1 hour [...]

RXN4Chemistry:
Recipe prediction

Thought: I have a synthesis plan for DEET. Now I will execute the synthesis.

Action: RXNExecute

Action input: CCN(CC)C(=O)c1cccc(C)c1

Observation: Synthesis procedure completed!

RoboRXN

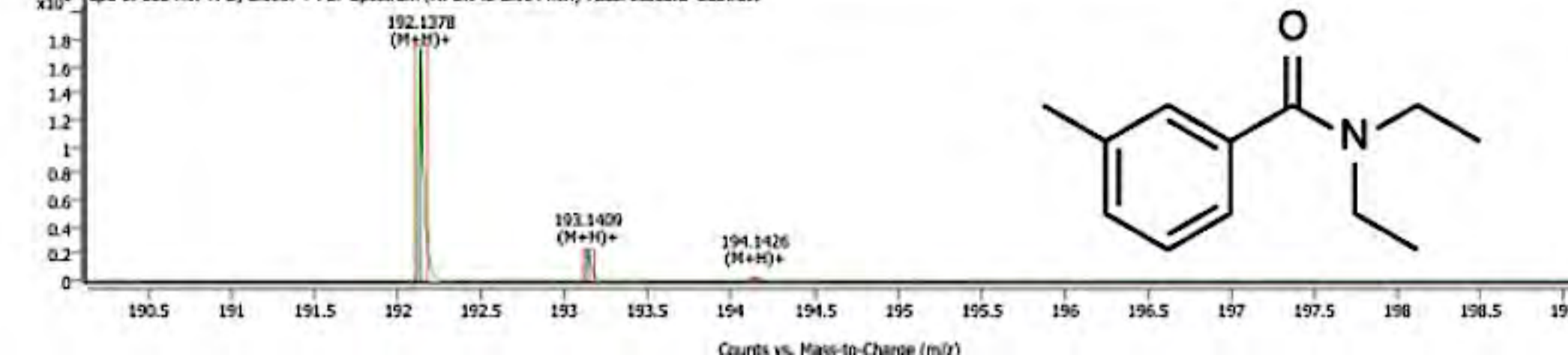


Compound Summary

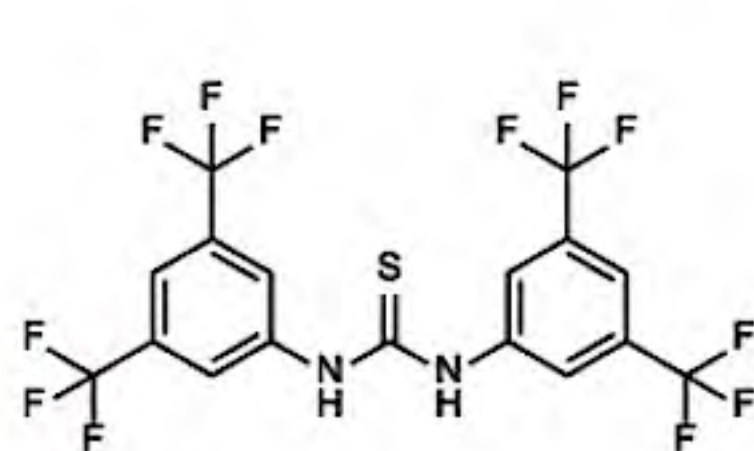
Cpd	Name	Formula	CAS	RT	Mass	Mass (Tgt)
1		C12 H17 N O		2.695	191.1305	191.1310
2		C4 H11 N		3.405	73.0896	73.0891

Compound Spectra (overlaid)

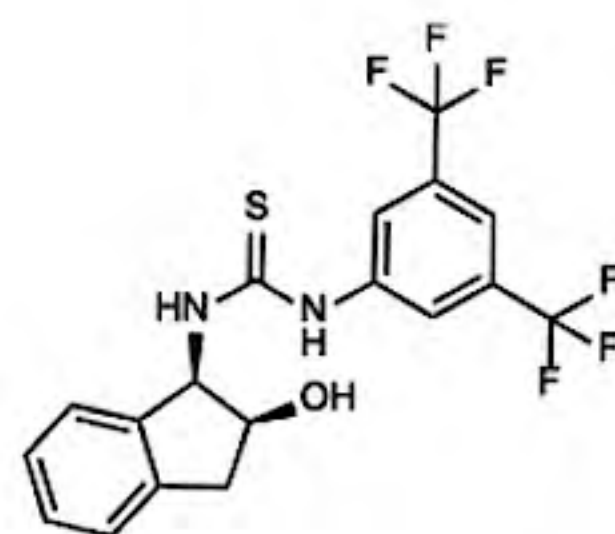
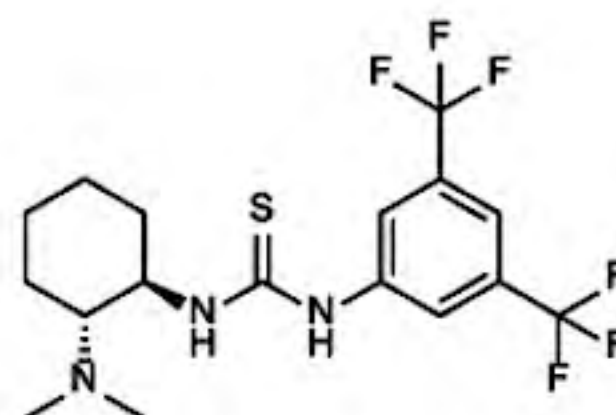
Cpd 1: C12 H17 N O; 2.695: + FBP Spectrum (rt: 2.943-2.954 min) 7adc76cbb2.d Subtract



- Find and synthesize a thiourea organocatalyst which accelerates a Diels-Alder reaction.



Schreiner's

Ricci's
Organocatalysts

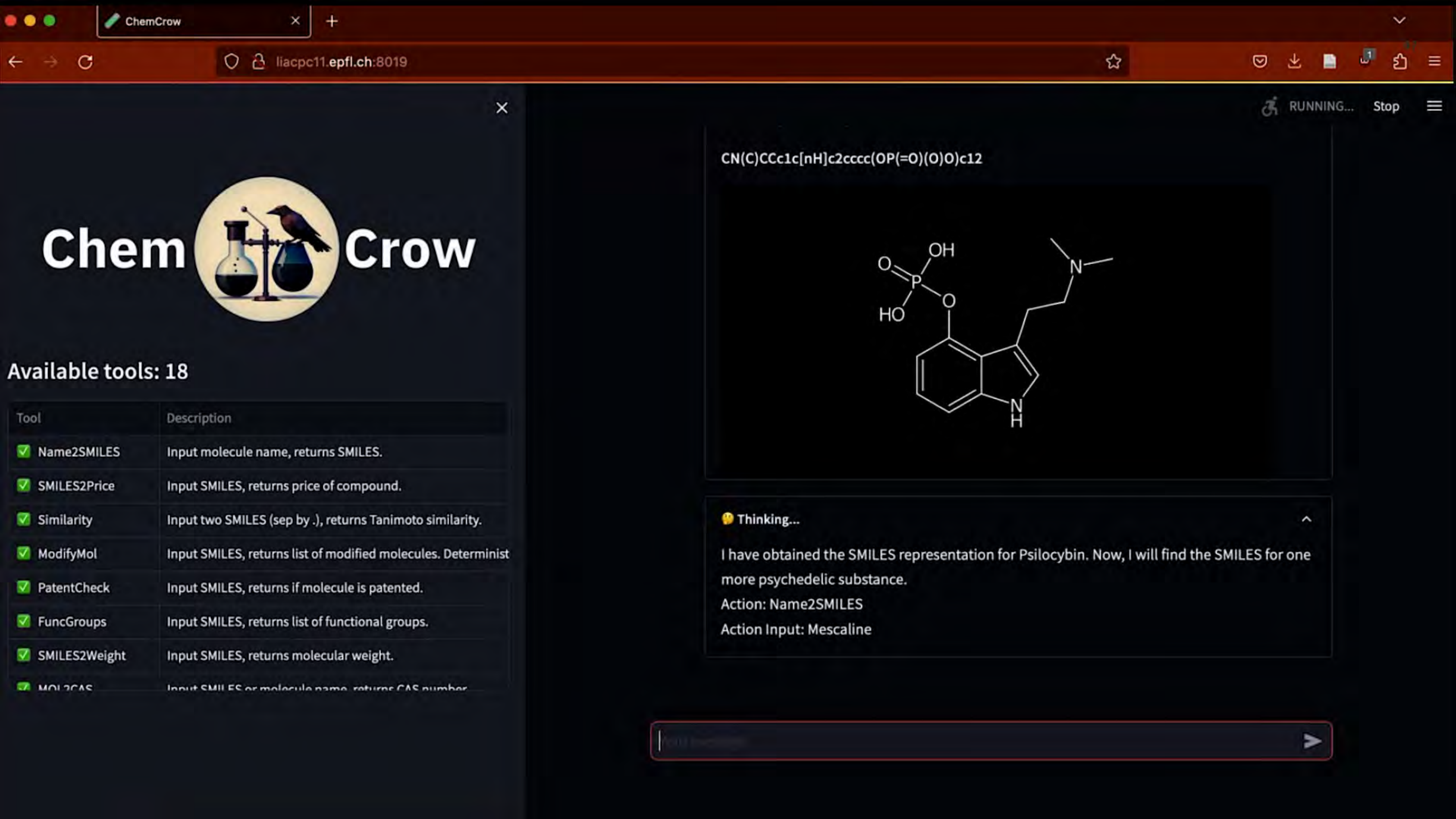
Takemoto's



ETH zürich

EPFL





ChemCrow

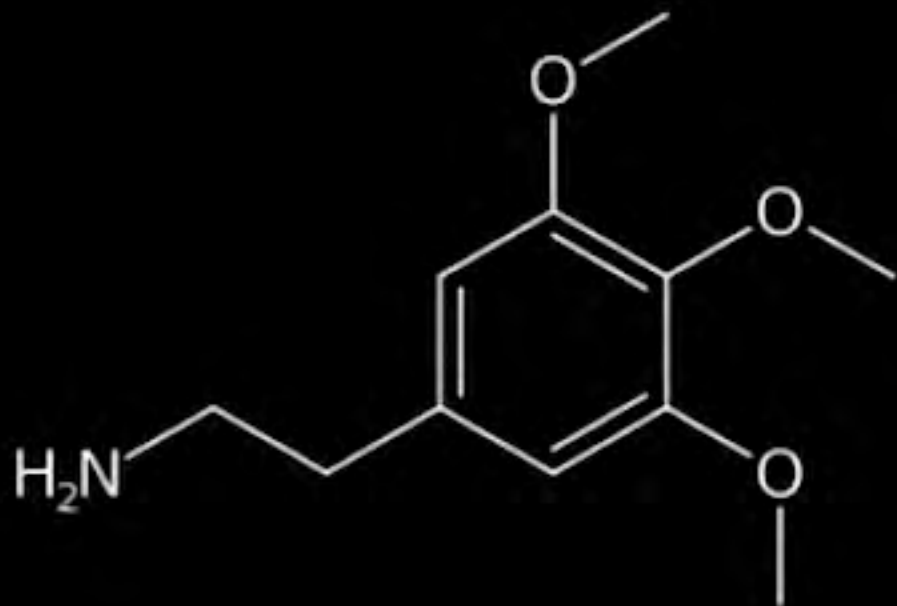
liacpc11.epfl.ch:8019

ChemCrow

Available tools: 18

Tool	Description
✓ Name2SMILES	Input molecule name, returns SMILES.
✓ SMILES2Price	Input SMILES, returns price of compound.
✓ Similarity	Input two SMILES (sep by .), returns Tanimoto similarity.
✓ ModifyMol	Input SMILES, returns list of modified molecules. Determinist
✓ PatentCheck	Input SMILES, returns if molecule is patented.
✓ FuncGroups	Input SMILES, returns list of functional groups.
✓ SMILES2Weight	Input SMILES, returns molecular weight.
✓ MOL2CAS	Input SMILES or molecule name, returns CAS number.

COc1cc(CCN)cc(OC)c1OC



🔔 Name2SMILES: Acetic chloride

I have obtained the SMILES representation for Mescaline. Now, I will find the SMILES for acetic chloride to predict the reaction.
Action: Name2SMILES
Action Input: Acetic chloride

ChemCrow

liacpc11.epfl.ch:8019

🔍 ⬇️ 📄 📌 1 🏠 ☰

ChemCrow

Available tools: 18

Tool	Description
✓ Name2SMILES	Input molecule name, returns SMILES.
✓ SMILES2Price	Input SMILES, returns price of compound.
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✓ ModifyMol	Input SMILES, returns list of modified molecules. Determinist
✓ PatentCheck	Input SMILES, returns if molecule is patented.
✓ FuncGroups	Input SMILES, returns list of functional groups.
✓ SMILES2Weight	Input SMILES, returns molecular weight.
✓ MOL2CAS	Input SMILES or molecule name, returns CAS number.

⌂ RUNNING... Stop ☰

CC(=O)Cl

🔔 RXNPredict: CCN(CC)C(=O)[C@@H]1C=C2c3ccc4[nH]cc(c34)C[C@H]2N(C)C1.CC(=O)...

Now that I have the SMILES for LSD, Psilocybin, Mescaline, and Acetic chloride, I can predict the reaction products using the RXNPredict tool. I will do this for each psychedelic substance separately.

Action: RXNPredict

Action Input: CCN(CC)C(=O)[C@@H]1C=C2c3ccc4[nH]cc(c34)C[C@H]2N(C)C1.CC(=O)Cl

➡

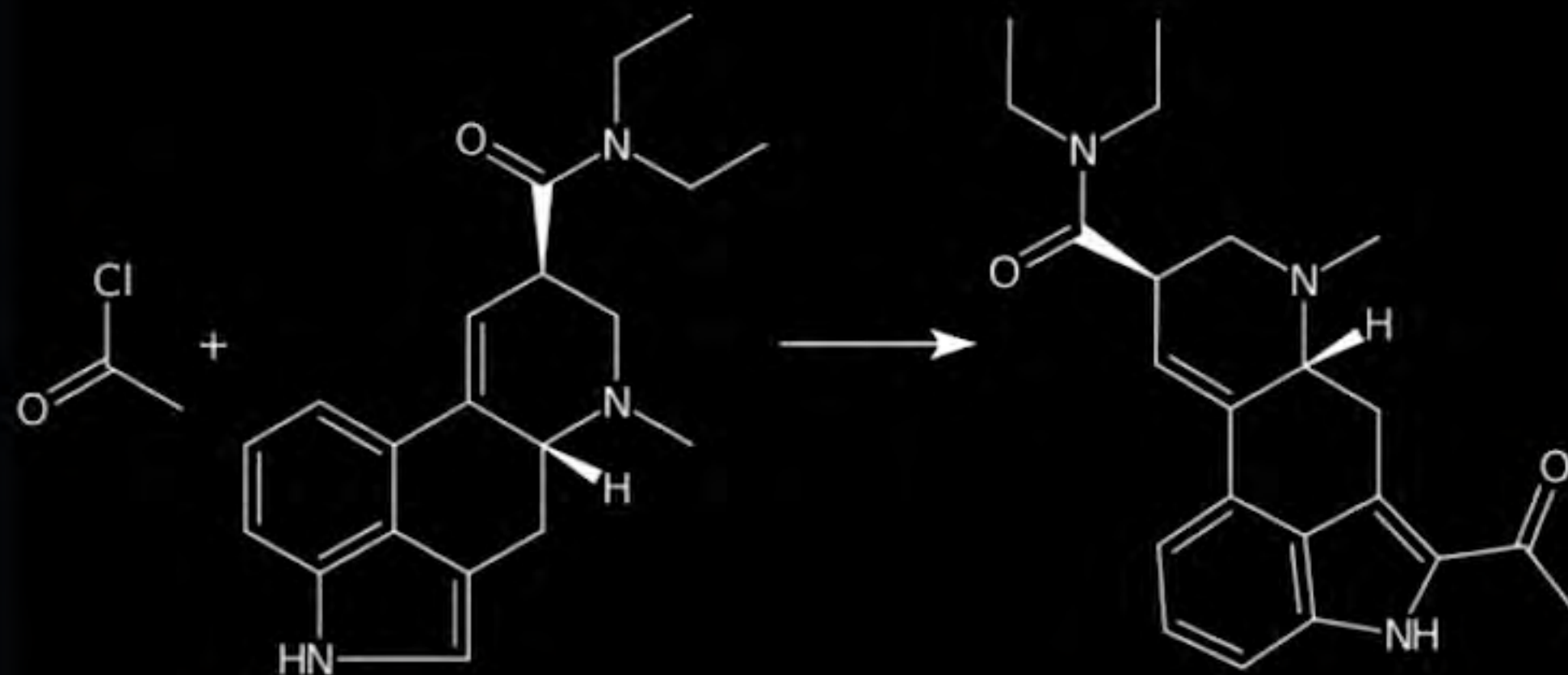
ChemCrow



Available tools: 18

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CCN(CC)C(=O)[C@@H]1C=C2c3ccc4[nH]c(C(C)=O)c(c34)C[C@H]2N(C)C1

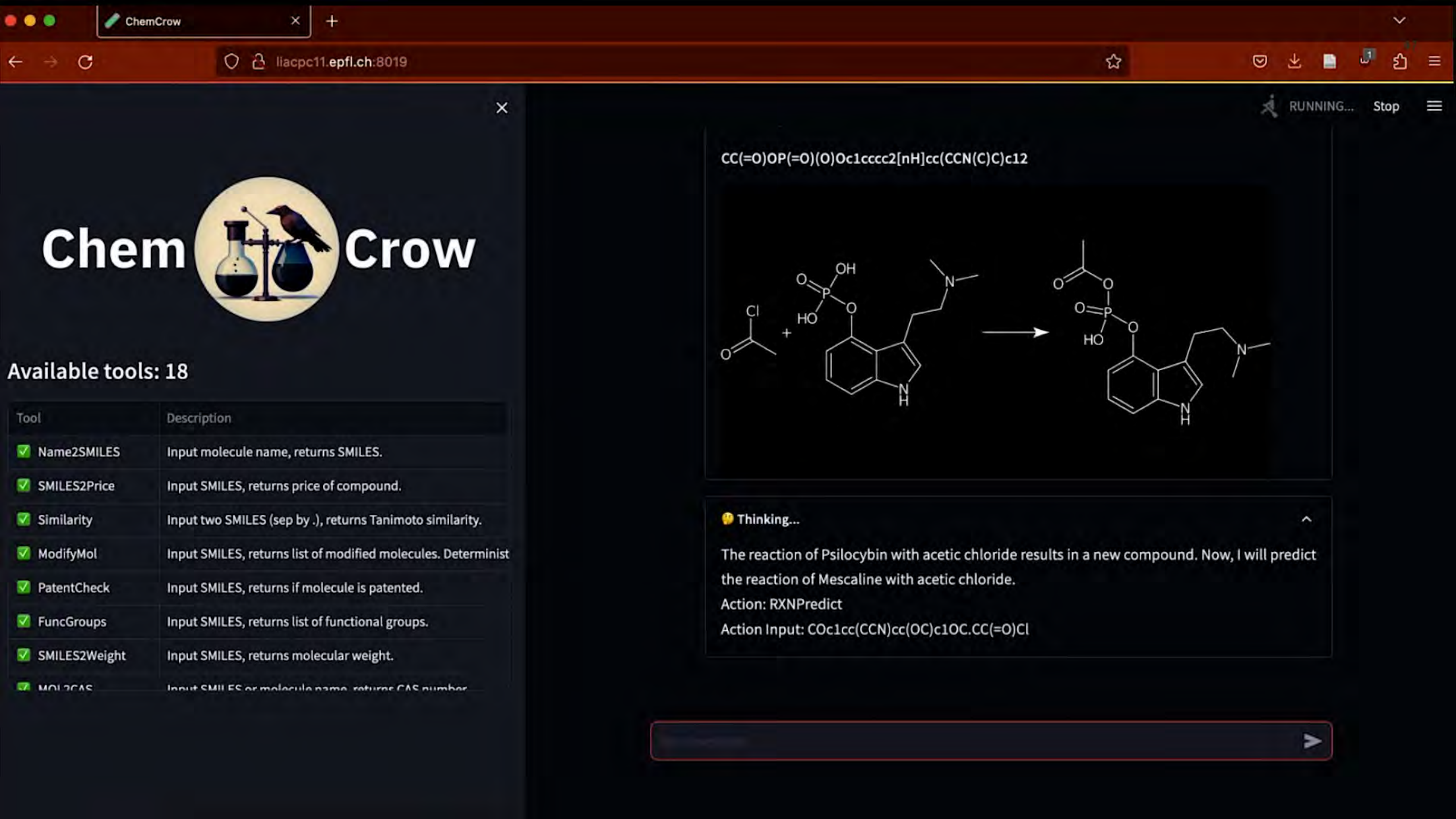


🚩 RXNPredict: CN(C)CCc1c[nH]c2cccc(OP(=O)(O)O)c12.CC(=O)Cl

The reaction of LSD with acetic chloride results in a new compound. Now, I will predict the reaction of Psilocybin with acetic chloride.

Action: RXNPredict

Action Input: CN(C)CCc1c[nH]c2cccc(OP(=O)(O)O)c12.CC(=O)Cl



ChemCrow is modular.

Toolset and LLMs can be changed.

More use cases in the paper.

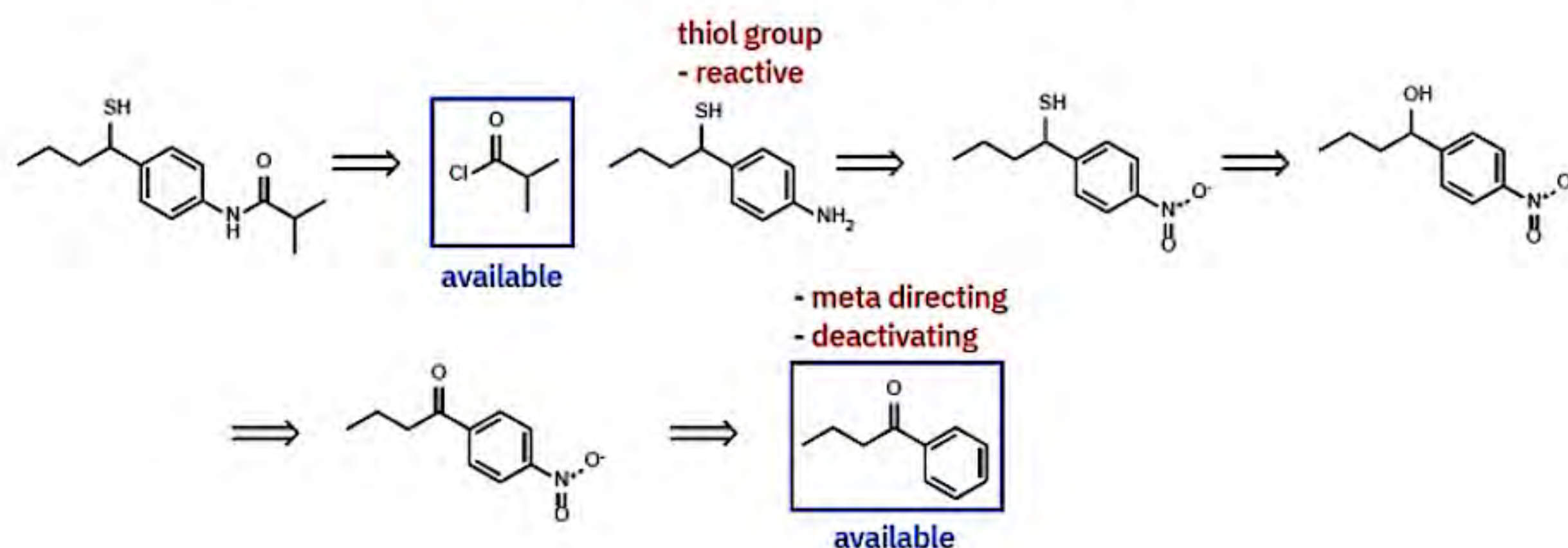


Code: <https://github.com/ur-whitelab/chemcrow-public>

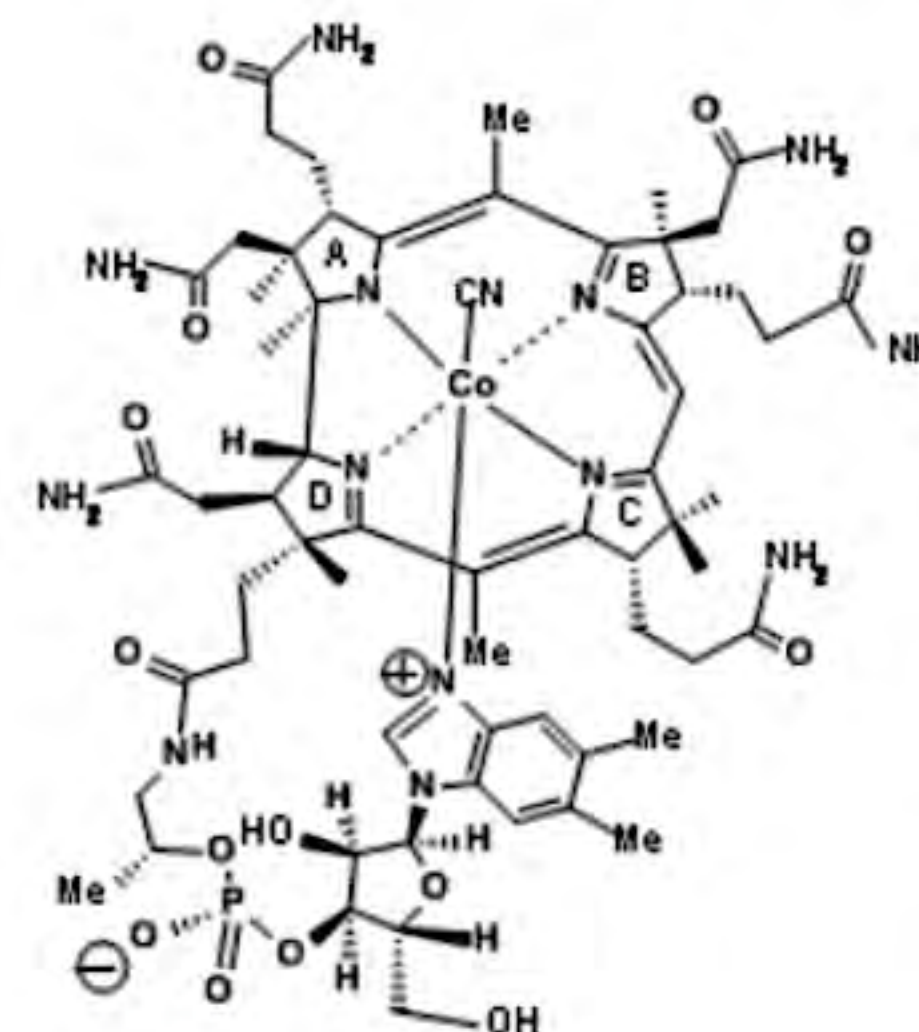
- Supply chain/robotics challenges
- Weak synthesis planning models



Daniel Armstrong & Zlatko Jončev

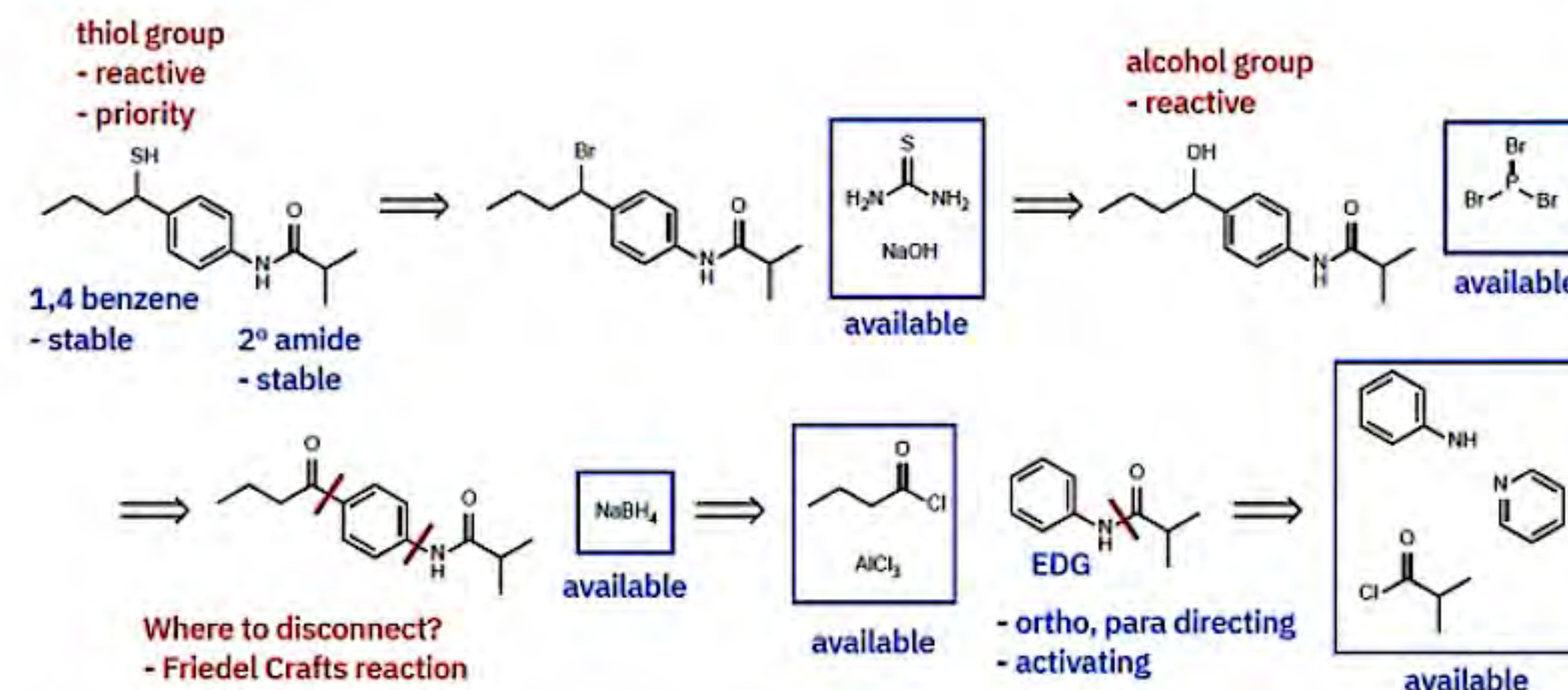


Typical output of unnamed system



Complexity that an
chemist would like to target:
Vitamin B12

- 12 year
- ~90 postdocs, 12 PhD students
- Eschenmoser, Woodward



More strategic synthesis plan (including reagents)

Bayesian optimization for additive screening and yield improvements – beyond one-hot encoding

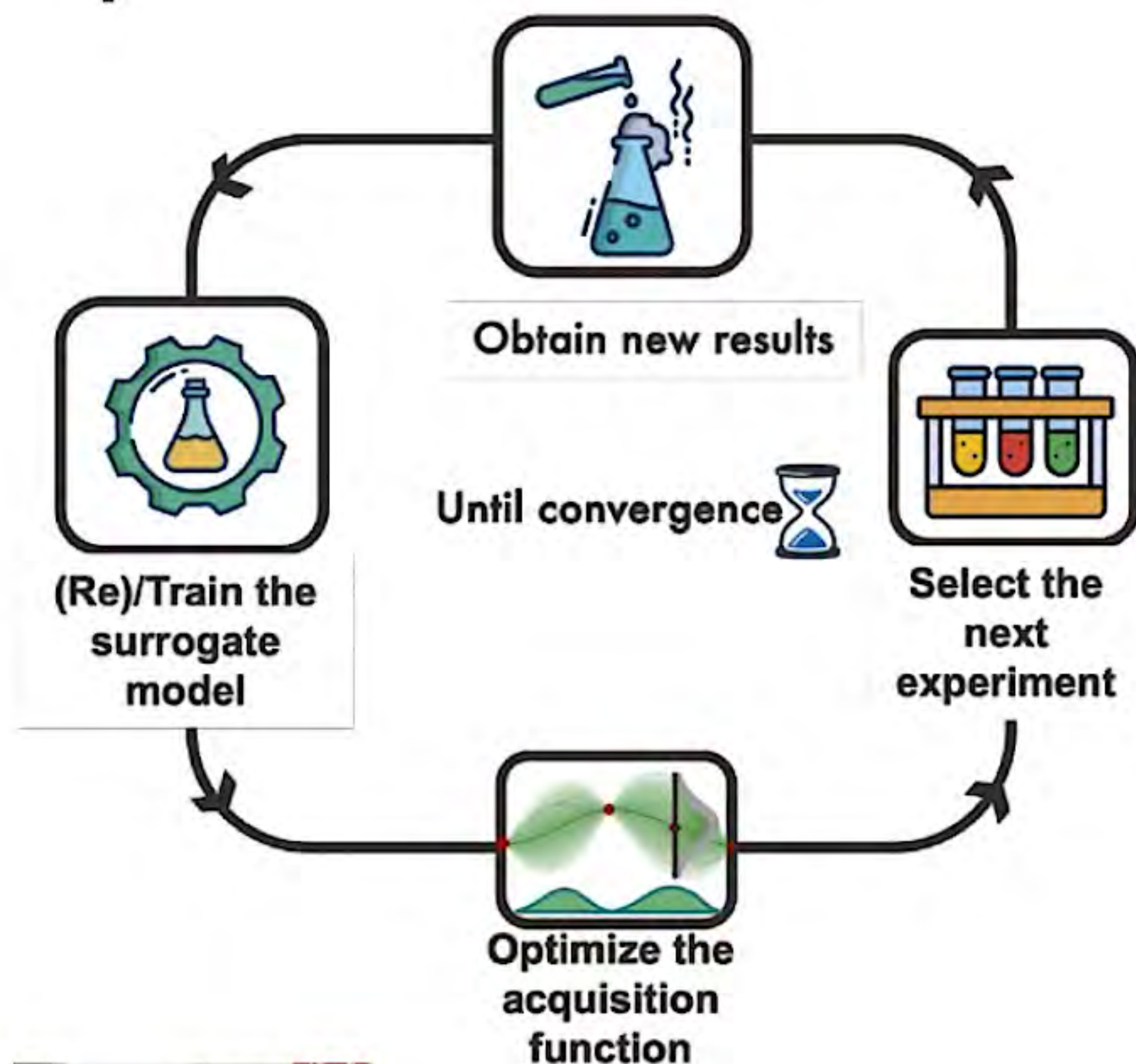
Digital Discovery



ROYAL SOCIETY OF CHEMISTRY

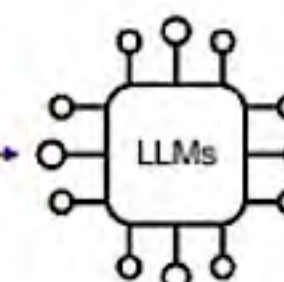
PAPER
Bojana Ranković, William Schreiner et al.
Bayesian optimization for additive screening and yield improvements – beyond one-hot encoding

BoChemian: Large Language Model embeddings for Bayesian optimization of chemical reactions



■ BOjana Ranković

```
Reaction Setup
The following solution was
prepared in DMSO:
- Ligand: {ligand_smile}
- Aryl halide:
{aryl_halide_smile}
- Additive: {additive_smile}
- Base: {base_smile}
```

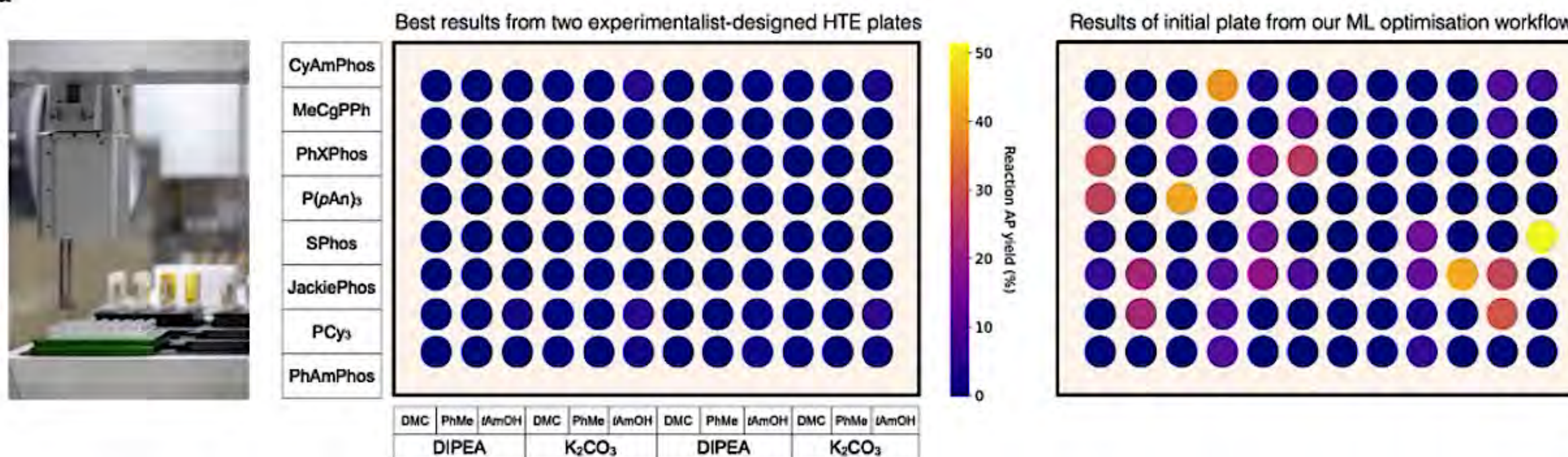


Embedding vector

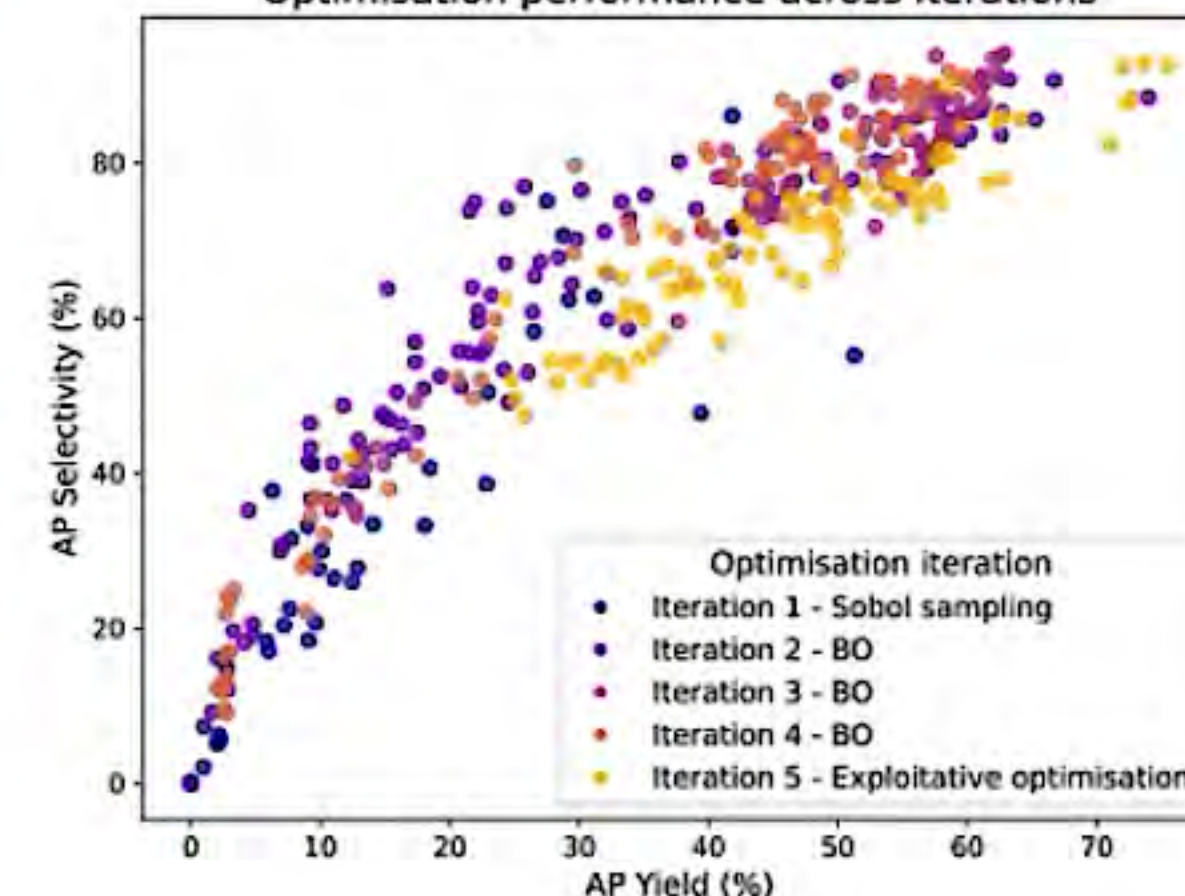




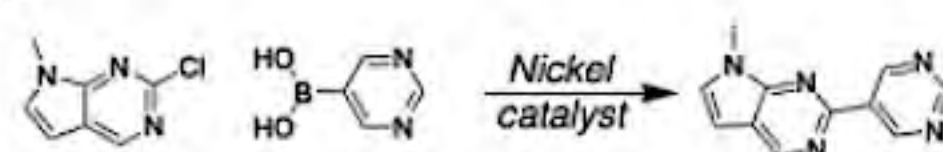
a



Optimisation performance across iterations

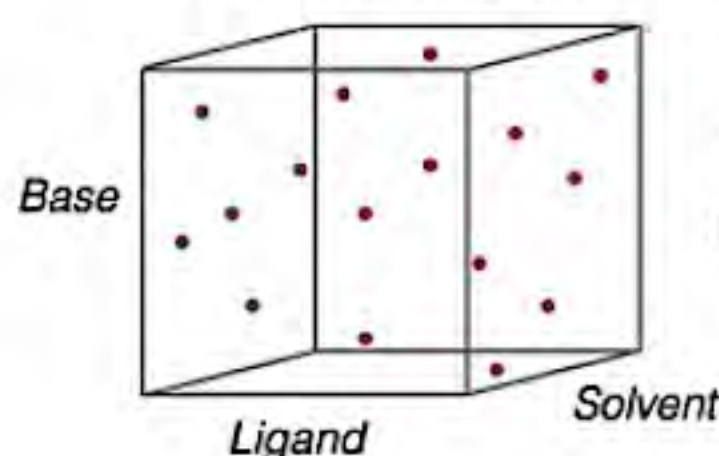


b



Ligand	Base	Solvent	...	Objectives
L1	B1	S1	...	Yield
...	...	...	...	...

Large, chemist-defined reaction condition space of many possible combinations

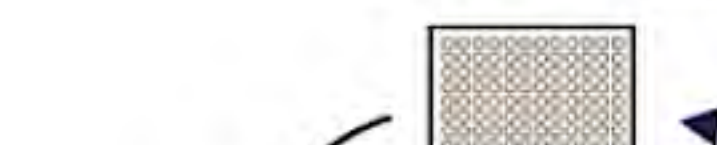


Initial quasi-random Sobol guided experiment selection

Experiments • diversely cover reaction condition space

Rapid suggestion and execution of batches of 96 parallel experiments at multiple temperatures

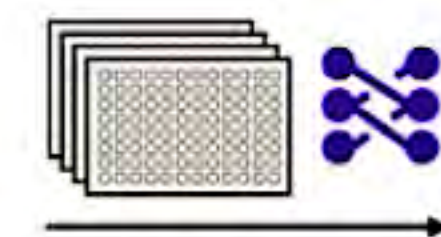
ML model suggests next 96 well HTE plate



Highly-parallel reaction condition optimisation

Automated HTE Robotic Experimentation

Scalable noise robust ML pipeline



Exploitative iteration with knowledge gained from multiple 96-well plates

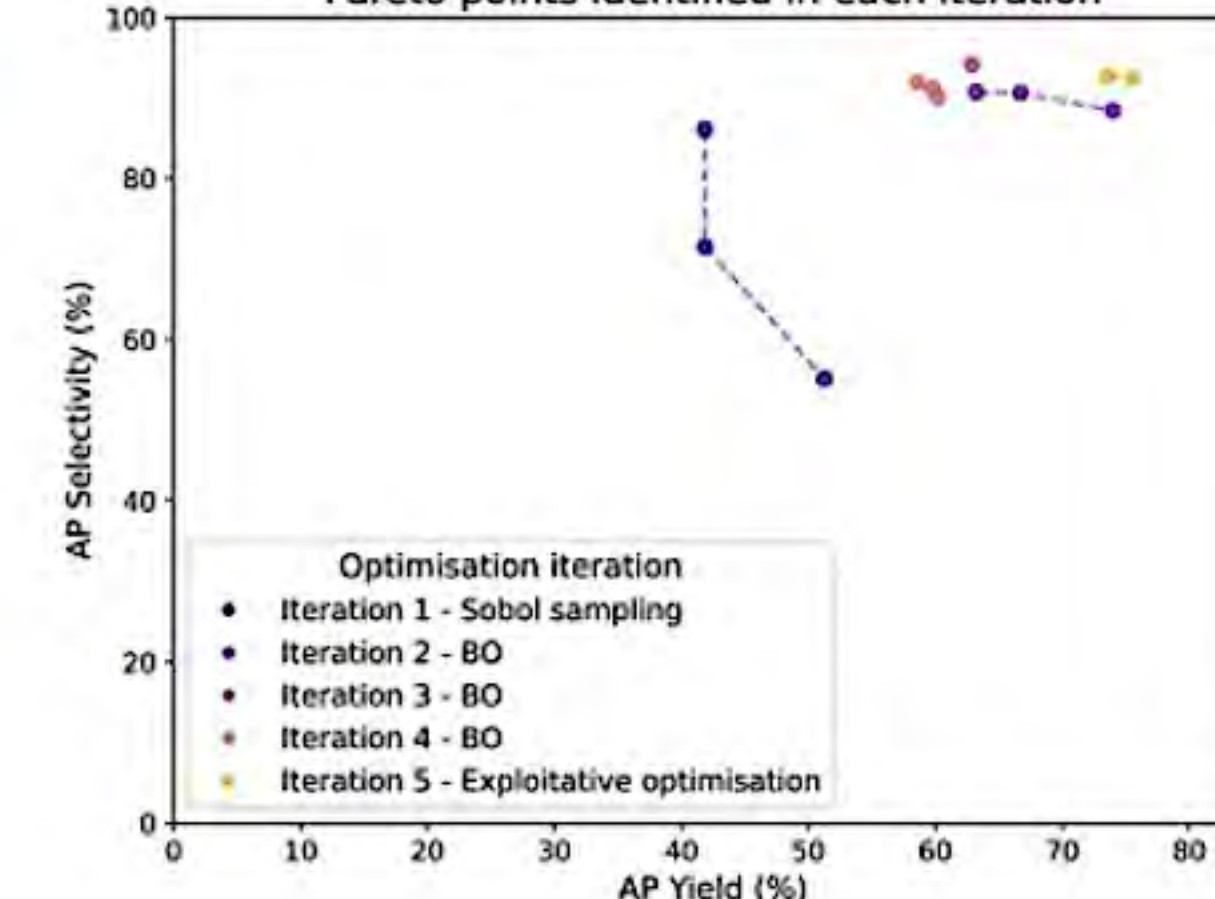
- Optimised selectivity and yield
- Unexpected reactivity insights

Feature analysis uncovers reactivity trends



Reaction objectives further improved

Pareto points identified in each iteration

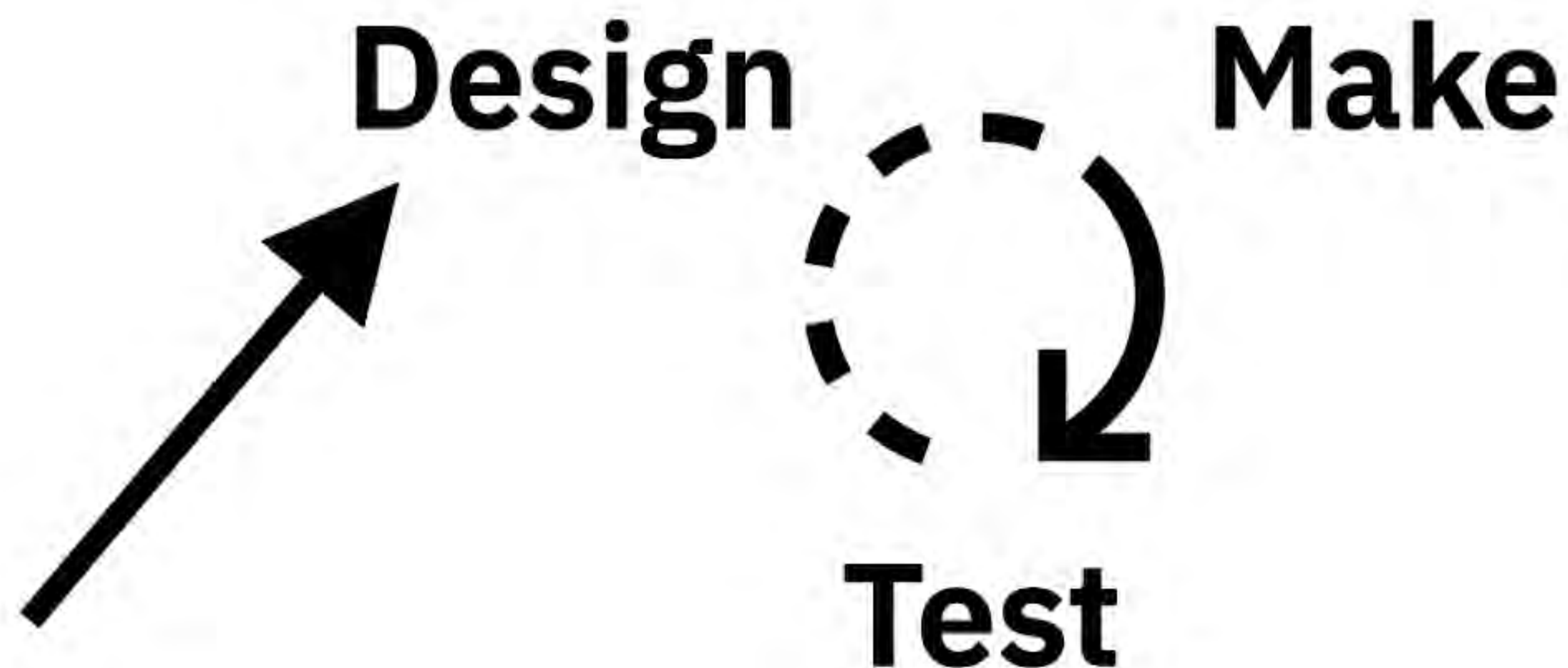


Highly Parallel Optimisation of Nickel-Catalysed Suzuki Reactions through Automation and Machine Intelligence

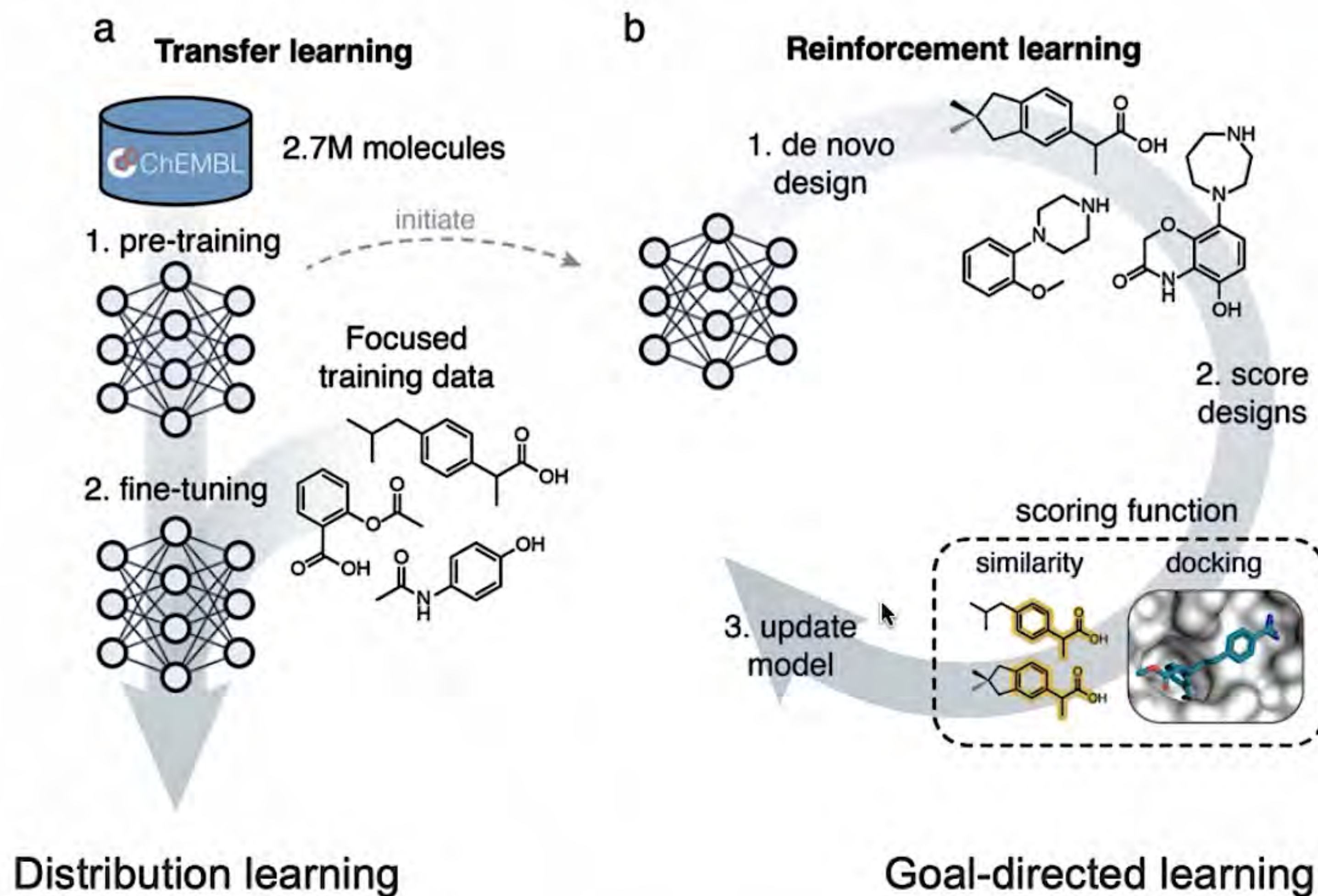
What molecule to make?

How to make it?

Reaction prediction
Synthesis planning



Experimental validation





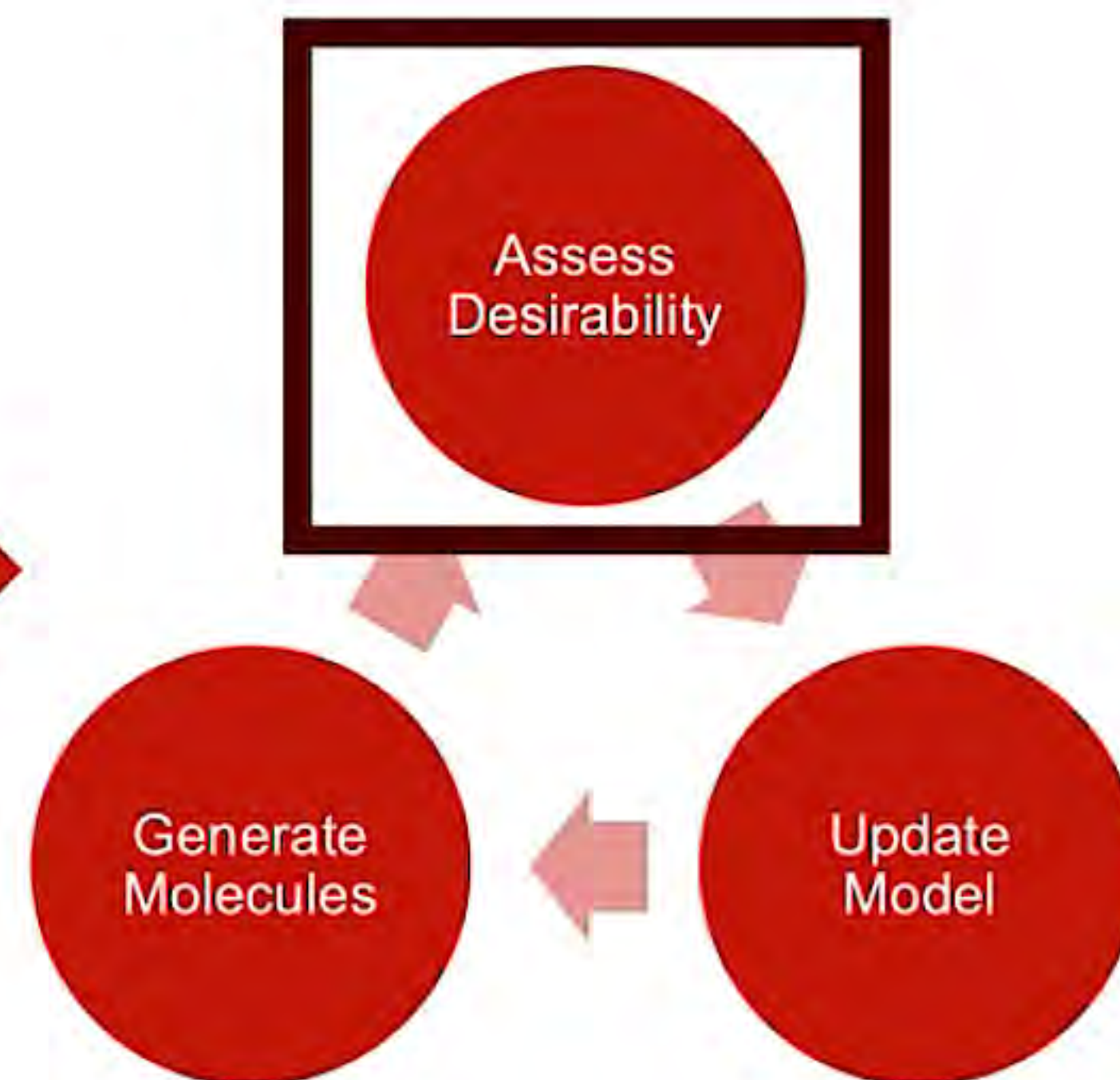
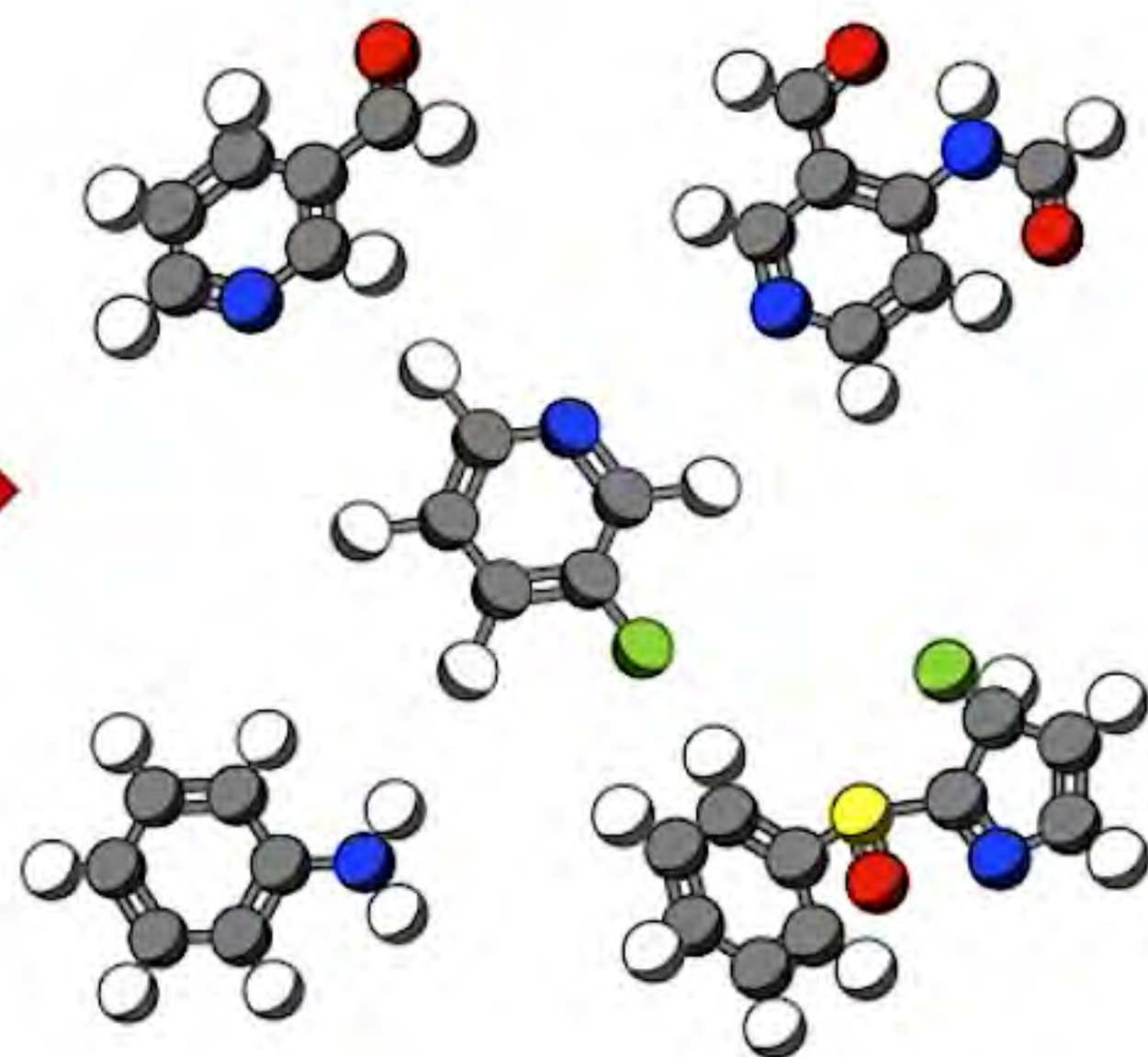
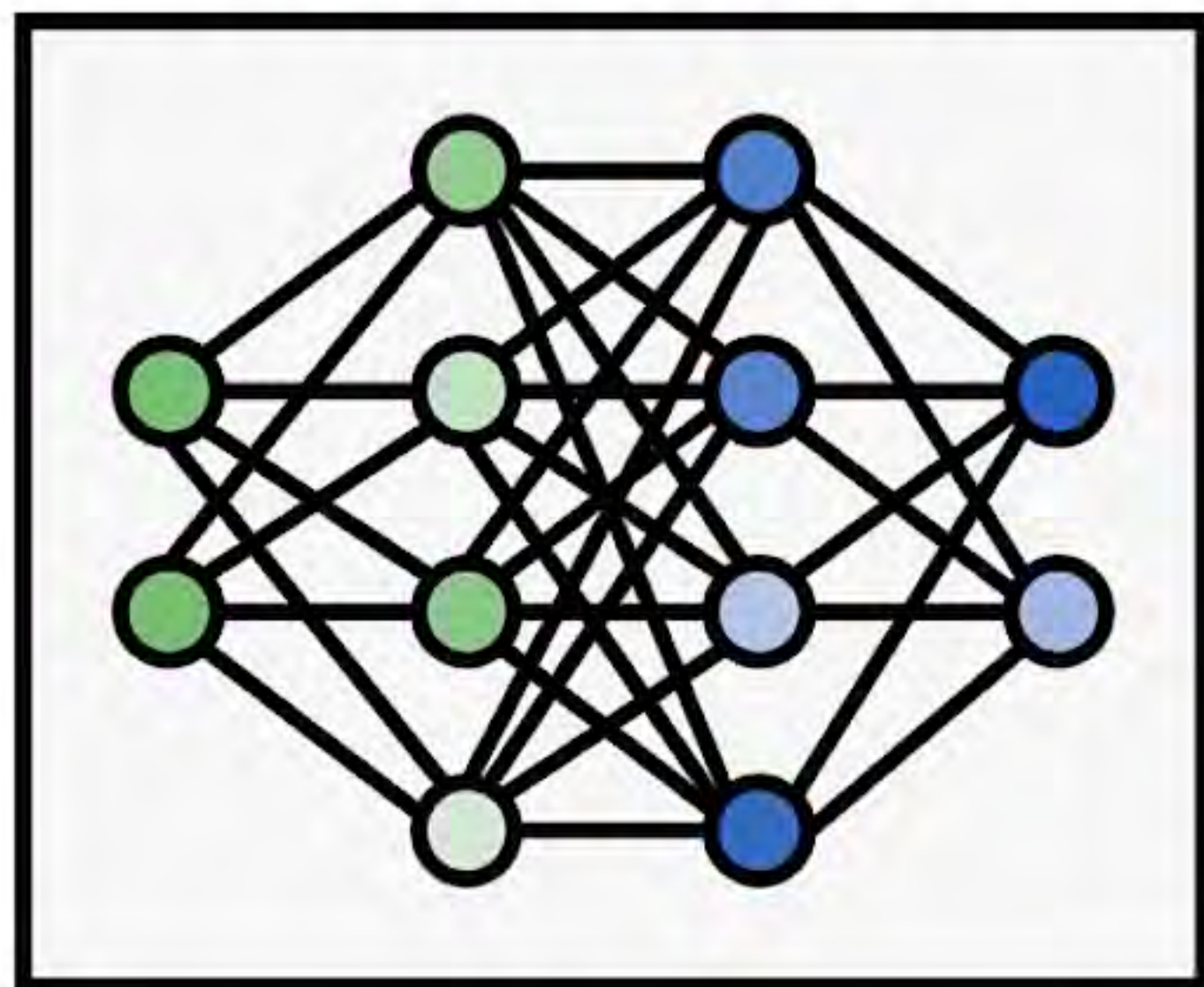
Objective

Design molecules satisfying a desired property profile

Key Consideration

Is the *in silico* predictor even correlated with the desired end-point?

Generative Model





Oracle: Computational prediction or simulation

Sample Efficiency: How few oracle evaluations are required to optimize the objective?

Increasing predictive accuracy *but* also computational cost

Drug Discovery

Predictive Models



*Can be accurate but may have narrow domain of applicability

Molecular Docking



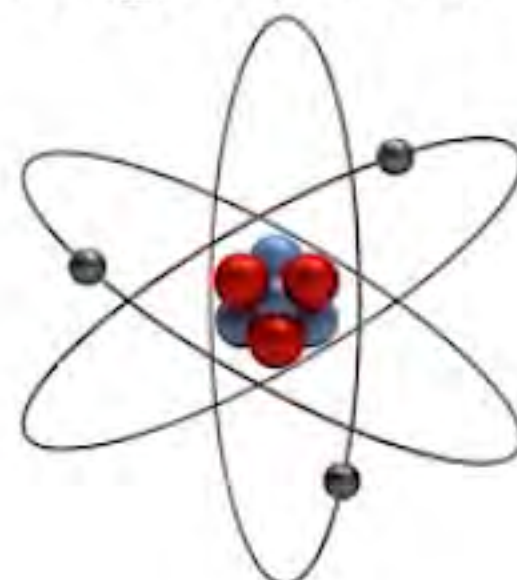
Materials and Catalyst Design



Semi-empirical QM



Single-point DFT



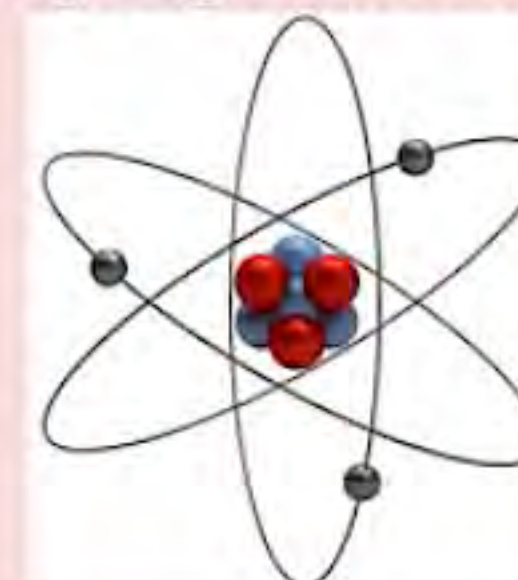
MMPB(GB)SA



Free Energy Calculations



DFT – Varying Functionals / MD





Augmented Memory

- Combines data augmentation with experience replay
- New state-of-the-art on the PMO benchmark



Sample Efficiency Matters: A Benchmark for Practical Molecular Optimization

Wenhao Gao^{1*}, Tianfan Fu^{2*}, Jimeng Sun^{3,4}, Connor W. Coley^{1,5}

¹Department of Chemical Engineering, Massachusetts Institute of Technology,

²Department of Computational Science and Engineering, Georgia Institute of Technology,

³Department of Computer Science, University of Illinois at Urbana-Champaign,

⁴Carle Illinois College of Medicine, University of Illinois at Urbana-Champaign,

⁵Department of Electrical Engineering and Computer Science, Massachusetts Institute of Technology,

*Equal Contributions

{whgao,ccoley}@mit.edu, tfu42@gatech.edu, jimeng@illinois.edu

- Benchmarks 25 models across 23 optimization tasks
(imposing a **fixed** oracle budget -> **10k**)



PMO (Sample efficiency) Benchmark

Model	Rank (/30)	Score
Augmented Memory	1	15.002
REINVENT [4]	2	14.016
SynNet [8] [ICLR '22]	12	11.498
DoG-Gen [9] [NeurIPS '20]	13	11.456
DST [10] [ICLR '22]	15	10.989
MARS [11] [ICLR '21]	16	10.989
MIMOSA [12] [AAAI '21]	17	10.651
DoG-AE [9] [NeurIPS '20]	20	9.790
GFlowNet [13] [NeurIPS '21]	21	9.131
GA+D [14] [ICLR '20]	22	8.964
GFlowNet-AL [13] [NeurIPS '21]	27	8.406
JT-VAE [15] [ICML '18]	28	8.358

The benchmark includes all recent de novo design algorithms.

All within 1 point

Big ML conferences models

10k oracle calls

AUC Top-10



Published at the GEM workshop, ICLR 2024

DyMol: DYNAMIC MANY-OBJECTIVE MOLECULAR OPTIMIZATION WITH OBJECTIVE DECOMPOSITION AND PROGRESSIVE OPTIMIZATION

Dong-Hee Shin[†], Young-Han Son[†], Deok-Joong Lee, Ji-Wung Han & Tae-Eui Kam^{*}

Department of Artificial Intelligence, Korea University

{dongheeshin, yhson135, deokjoong, danielhan, kamte}@korea.ac.kr

Method	Four objectives		Five objectives		Six objectives	
	HV(↑)	R2(↓)	HV(↑)	R2(↓)	HV(↑)	R2(↓)
Random ZINC	0.065±0.013	4.809±0.243	0.005±0.003	8.867±0.202	0.001±0.000	14.456±0.344
SMILES-VAE	0.073±0.023	4.839±0.435	0.004±0.001	9.226±0.512	0.001±0.000	15.109±0.548
GFlowNet	0.063±0.011	4.952±0.196	0.011±0.004	9.011±0.314	0.004±0.001	14.462±0.422
MIMOSA	0.082±0.028	4.905±0.410	0.016±0.012	9.195±0.761	0.005±0.005	15.454±1.701
LaMBO	0.123±0.006	4.496±0.148	0.009±0.001	9.302±0.159	Out-of-memory	Out-of-memory
HN-GFN	0.120±0.000	4.013±0.060	0.004±9.861	9.861±0.096	0.002±0.000	15.349±0.102
MOEA/D	0.176±0.123	4.615±1.193	0.094±0.052	8.105±1.561	0.025±0.018	15.054±2.028
NSGA-III	0.234±0.107	3.477±0.837	0.071±0.047	7.130±1.349	0.016±0.020	14.366±2.393
GraphGA	0.254±0.069	3.379±0.666	0.100±0.057	7.676±1.312	0.051±0.030	11.814±1.787
GPBO	0.275±0.091	3.311±0.757	0.091±0.031	7.670±0.761	0.026±0.024	12.840±1.811
REINVENT BO	0.309±0.021	2.795±0.103	0.071±0.014	7.537±0.620	0.033±0.019	10.929±1.019
REINVENT	0.338±0.030	2.770±0.116	0.099±0.054	7.578±1.187	0.062±0.028	10.032±0.922
AugMem	0.395±0.038	2.496±0.192	0.090±0.043	8.011±0.955	0.071±0.072	12.472±3.758
DyMol (Ours)	0.422±0.023	2.297±0.095	0.247±0.087	4.943±0.990	0.143±0.056	8.842±1.632

- DyMol = REINVENT + “progressive optimization”
- Augmented Memory is 2nd most sample-efficient

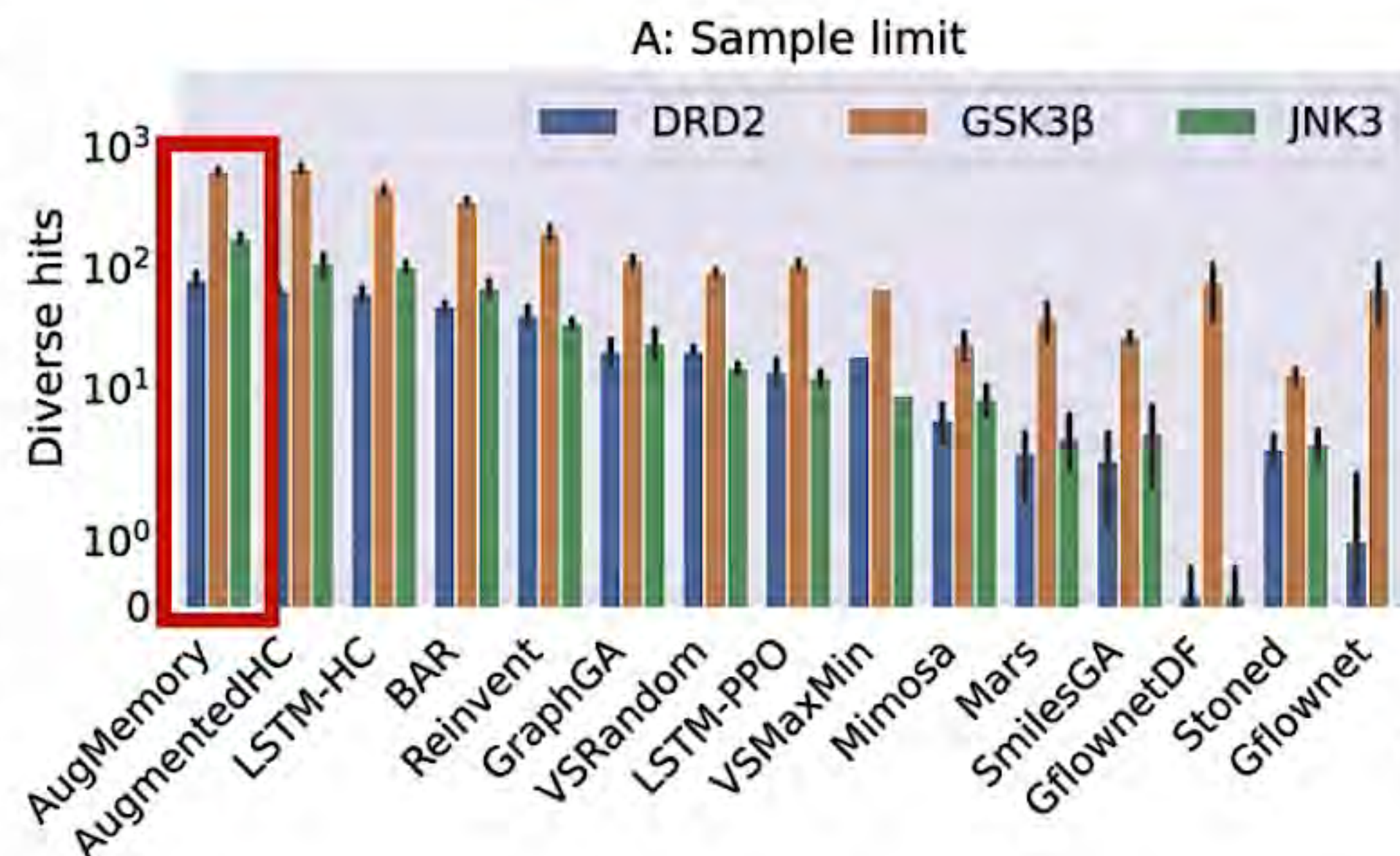
Published at the GEM workshop, ICLR 2024

DIVERSE HITS IN DE NOVO MOLECULE DESIGN: A DIVERSITY-BASED COMPARISON OF GOAL-DIRECTED GENERATORS

Philipp Renz

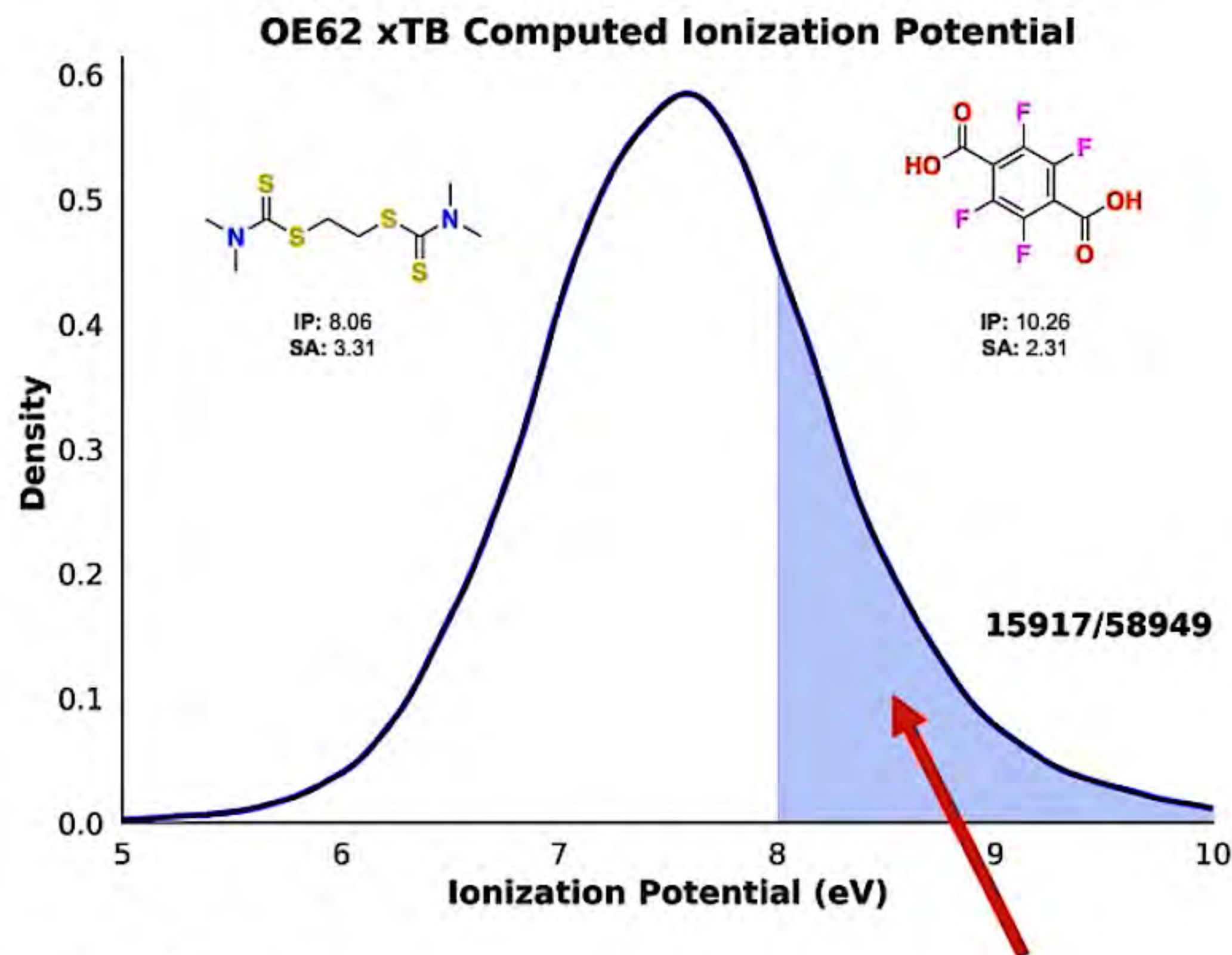
Sohvi Luukkonen

Günter Klambauer



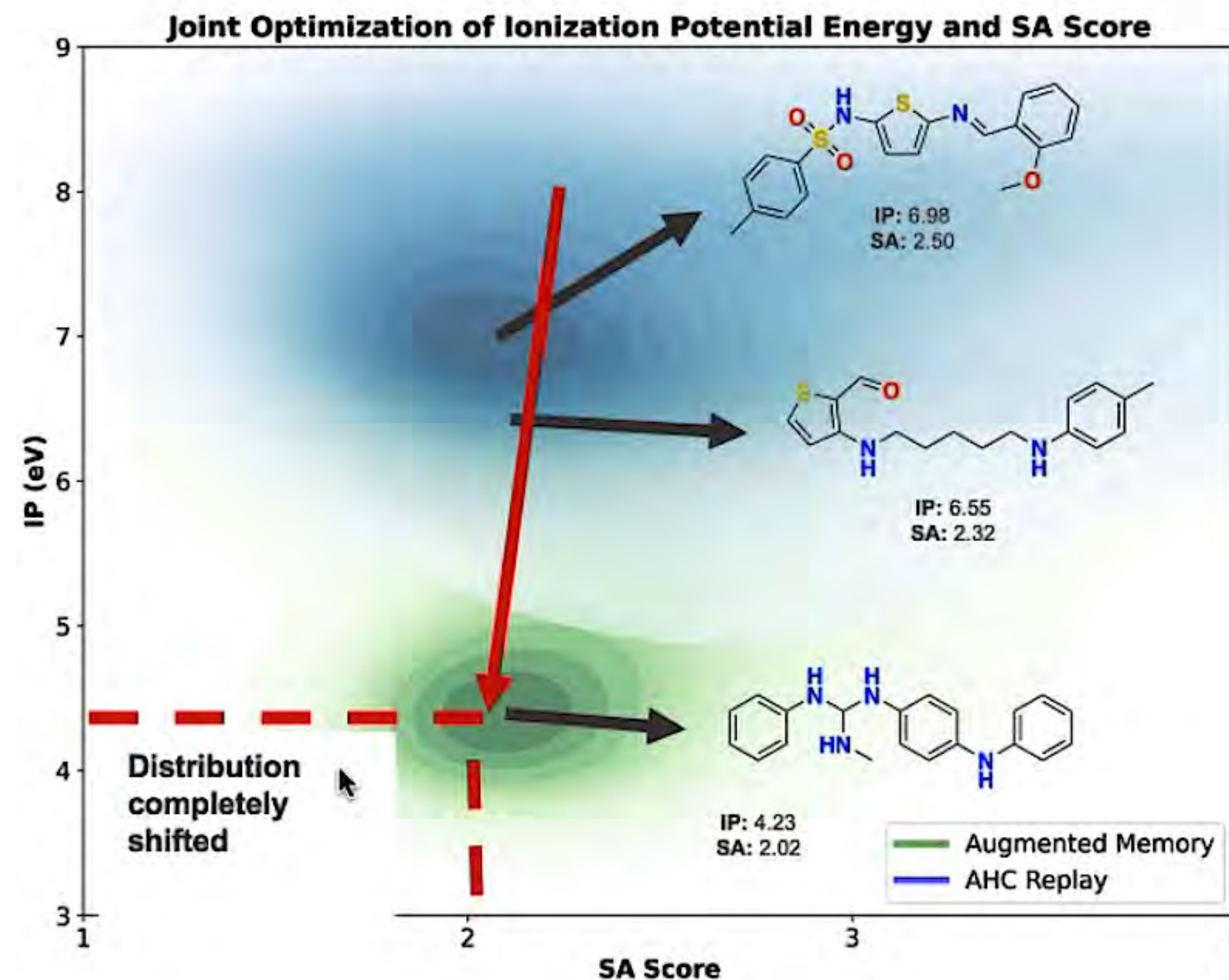
Augmented Memory is most sample-efficient

Semiconductor design – shifting the distribution away from the training set

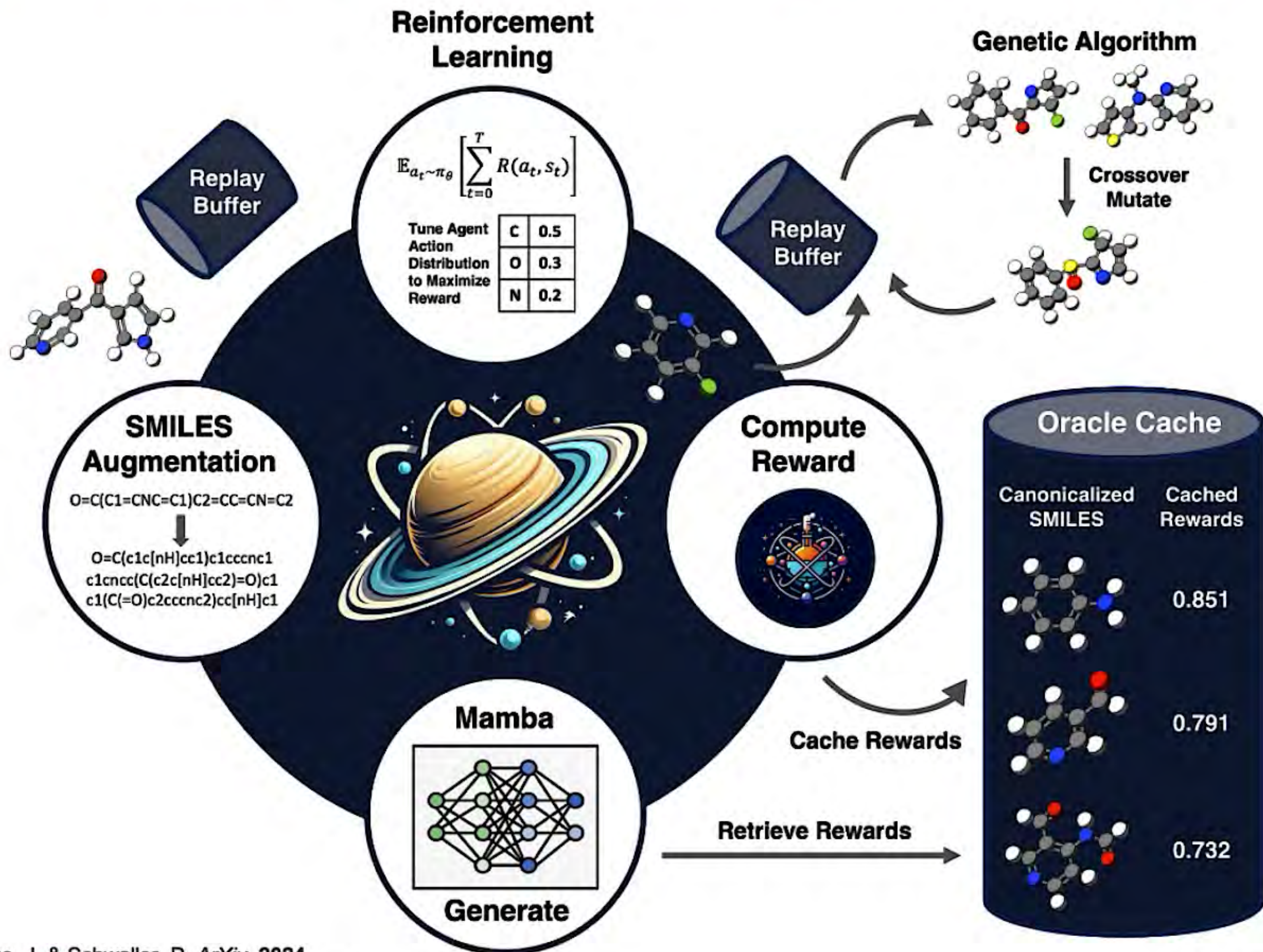


Optimization Objective

1. Minimize IP
2. Minimize SA



EPFL Saturn – Sample-efficient generative design framework



- Sample efficiency ***much*** better than *baseline*
- **Augmented Memory**
- State-of-the-art on drug discovery optimization tasks



Jeff Guo

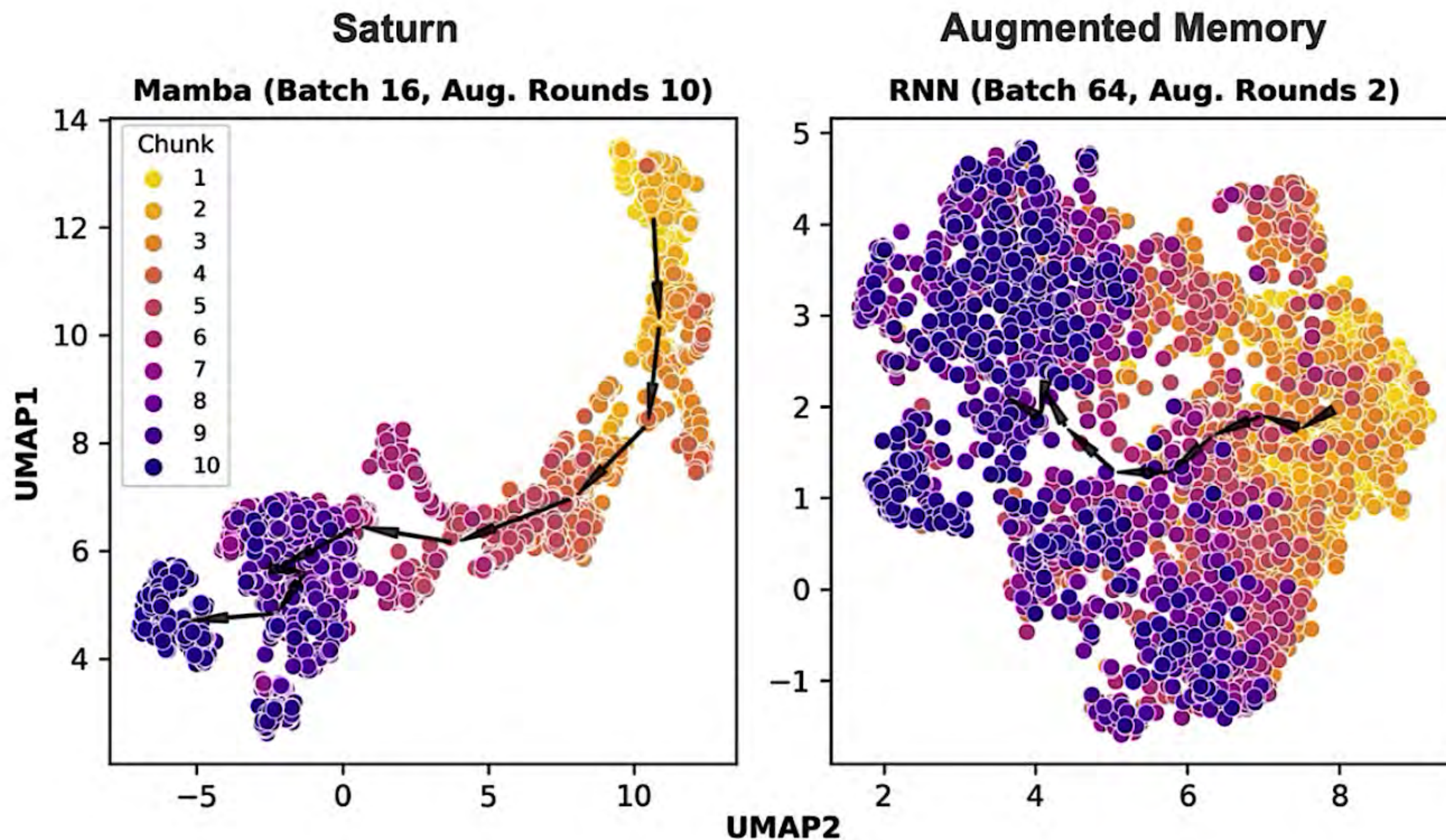


Target	Model	Yield (↑)	IntDiv1 (↑)	Scaffolds (↑)	OB 1 (↓)	OB 10 (↓)	OB 100 (↓)
DRD2	Augmented Memory	22 ± 7	0.774 ± 0.019	22 ± 7	143 ± 75(10)	733 ± 120(10)	Failed
	Saturn	369 ± 62	0.671 ± 0.050	310 ± 70	93 ± 53(10)	391 ± 56(10)	663 ± 55(10)
AChE	Augmented Memory	173 ± 19	0.843 ± 0.009	170 ± 18	57 ± 2(10)	189 ± 52(10)	776 ± 58(10)
	Saturn	480 ± 79	0.757 ± 0.020	400 ± 96	32 ± 24(10)	185 ± 82(10)	508 ± 80(10)
MK2	Augmented Memory	0.2 ± 0.4	—	0.2 ± 0.4	836 ± 186(2)	Failed	Failed
	Saturn	14.9 ± 14.1	0.454 ± 0.212	14.1 ± 13.2	677 ± 186(9)	861 ± 108(6)	Failed

Multi-objective docking optimization (docking score, MW < 500 Da, maximize QED), reward threshold 0.8, **1000 oracle calls**.

With Saturn (compared to Augmented Memory)

- Diversity decreases slightly
- **Yield**, number of molecules above reward threshold, **increases**
- **Oracle Burden (OB)**, number to get to X molecules above threshold, **decreases**





Problem Statement

Generated molecules need to be synthesizable for ***tangible real-world impact***

Synthesizability-constrained Generation

Generate molecules starting from available building blocks stitched together by ***known***, feasible chemical transformations (in 2024, Swanson et al., Nat. Mach. Intell. / Cretu et al., *arXiv* / Koziarski, et al., *arXiv*)

Why not directly optimize for synthesizability?

Inference time can make this ***impractical*** due to generative models' ***insufficient*** optimization efficiency

→ ***Sample efficiency problem***



Problem Statement

Generated molecules

tangible results

Synthesizability

Generate molecules

blocks stitch

transformation

Intell. / Creative

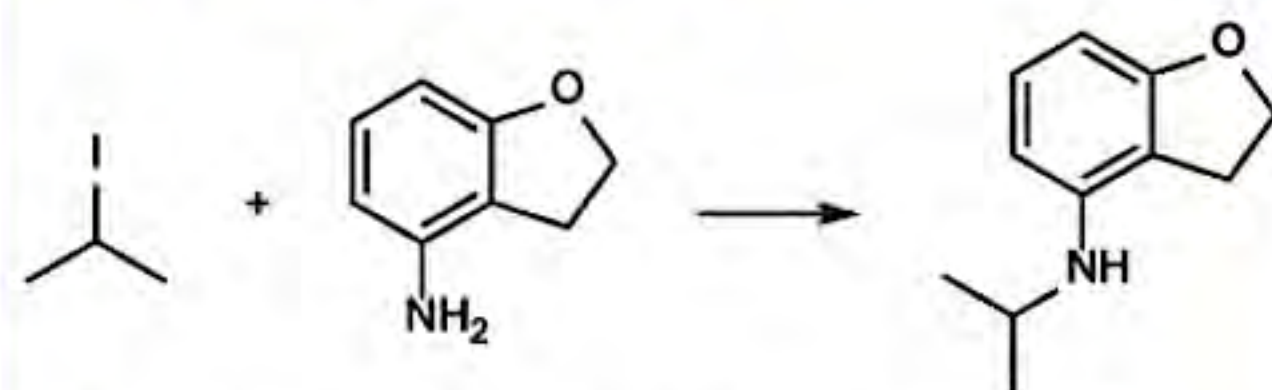
Why not directly optimize for synthesizability?

practical due to optimization

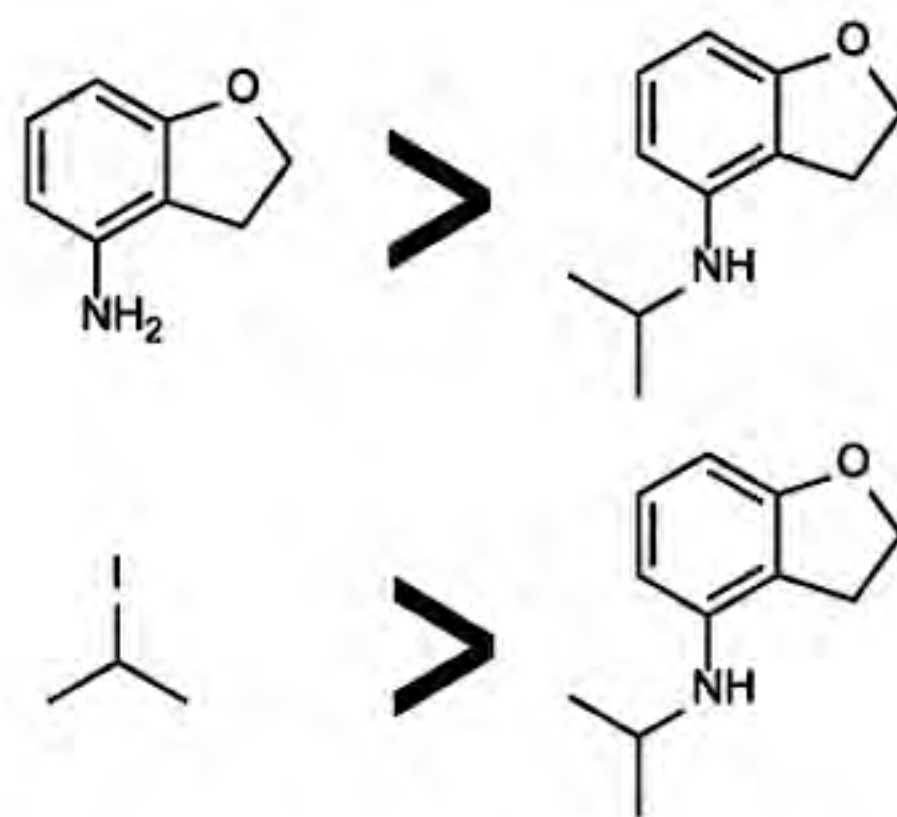
Method	Modes (Yield)	Mol. weight ($\downarrow$)	QED ($\uparrow$)	SA score ($\downarrow$)	AiZynth ($\uparrow$)	Oracle calls (Wall time)
Previous work		top-500	top-500	top-500	top-100	
GraphGA ⁶⁸	NR	521.0 $\pm$ 31.8	0.32 $\pm$ 0.07	4.14 $\pm$ 0.51	0.00	400,000 (NR)
SyntheMol ²⁷	NR	458.2 $\pm$ 60.7	0.45 $\pm$ 0.16	2.86 $\pm$ 0.56	0.56	100,000 ^a (72h)
FGFN ⁶⁹	NR	548.6 $\pm$ 42.9	0.22 $\pm$ 0.03	2.94 $\pm$ 0.54	0.25	400,000 (NR)
RGFN ²⁹	NR	526.2 $\pm$ 37.6	0.23 $\pm$ 0.04	2.83 $\pm$ 0.22	0.65	400,000 (72h)
$R_{All\ MPO}$ (ours)	4 objectives (Docking, QED, SA, AiZynth)					
Saturn-ChEMBL (10)	4 $\pm$ 1 (5 $\pm$ 3)	367.7 $\pm$ 15.7	0.70 $\pm$ 0.13	2.11 $\pm$ 0.19	0.91 $\pm$ 0.11	1,000 (2.9h $\pm$ 34m)
Saturn-GA-ChEMBL (10)	7 $\pm$ 6 (10 $\pm$ 9)	373.3 $\pm$ 20.9	0.67 $\pm$ 0.09	2.08 $\pm$ 0.23	0.82 $\pm$ 0.17	1,000 (2.1h $\pm$ 24m)
Saturn-ZINC (9)	6 $\pm$ 3 (8 $\pm$ 10)	368.7 $\pm$ 27.6	0.79 $\pm$ 0.08	2.15 $\pm$ 0.22	0.87 $\pm$ 0.19	1,000 (2.1h $\pm$ 29m)
Saturn-GA-ZINC (10)	7 $\pm$ 4 (10 $\pm$ 7)	382.8 $\pm$ 27.9	0.71 $\pm$ 0.08	2.10 $\pm$ 0.15	0.85 $\pm$ 0.17	1,000 (2.0h $\pm$ 26m)

Saturn: 1/400th the oracle budget and within 2 hours is sufficient to find at least *some* desirable molecules (across 10 replicates)

1. Pretraining

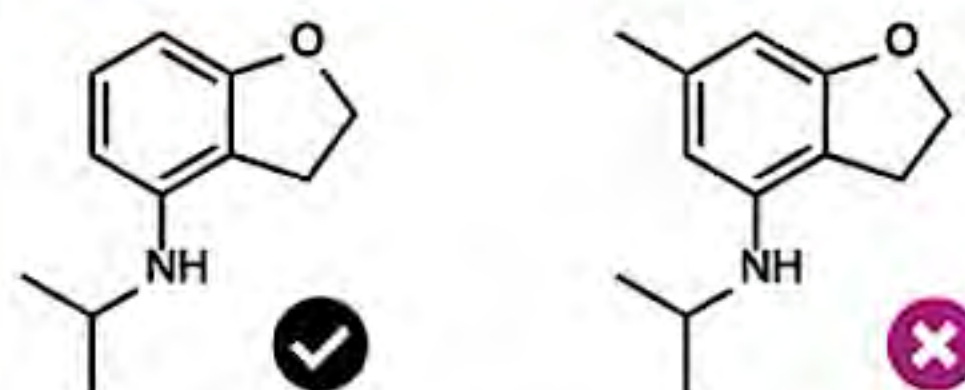


RXNs from USPTO/CHORISO



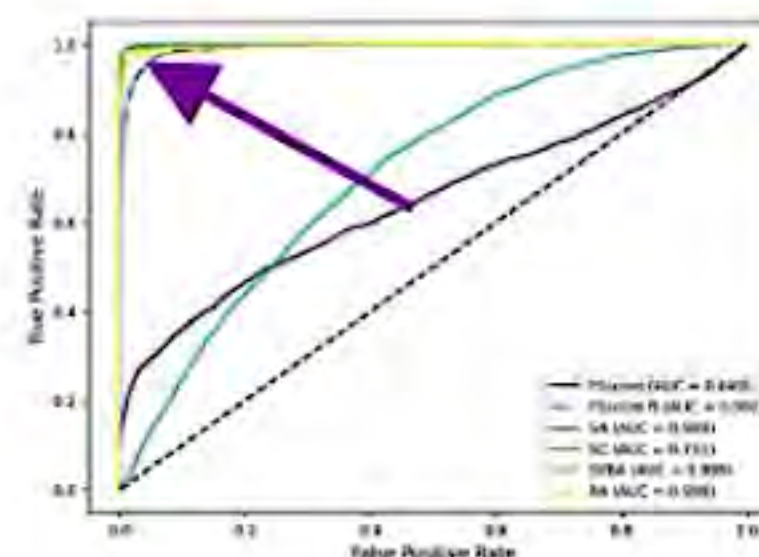
GNN for pair-wise ranking
Inspired by (SCScore, Coley)

2. Fine-tuning



Human preference labelling

- 50 pairs (AUC 0.65 to 0.99)
- 20 pairs (AUC 0.49 to 0.67)

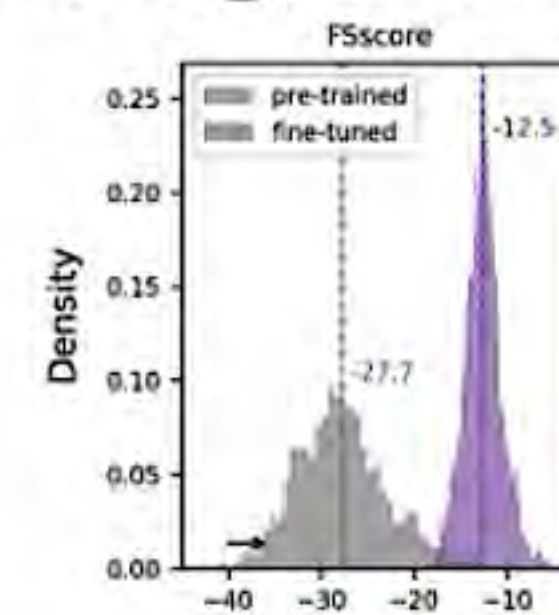


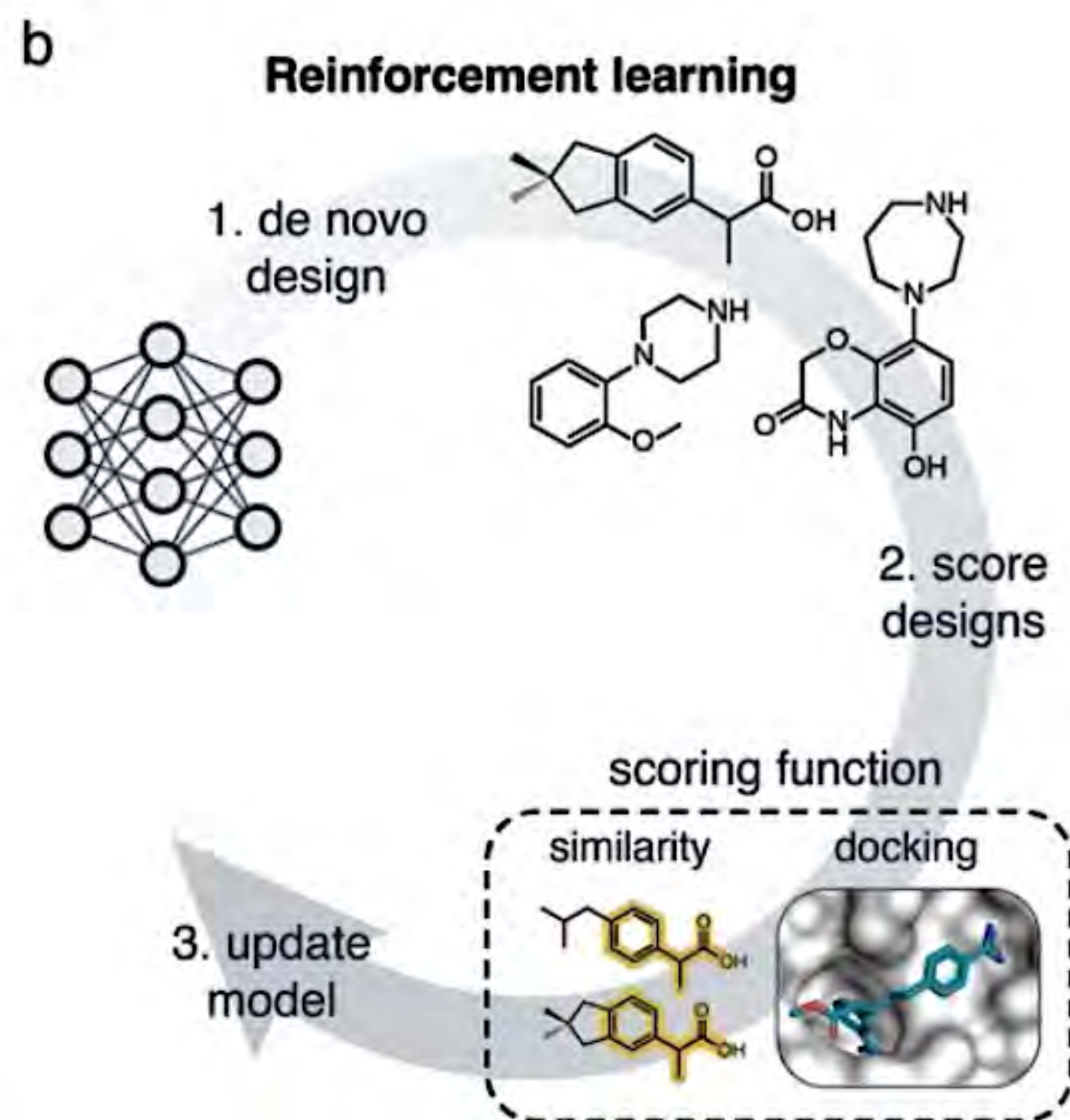
Inspired by MolSkill (Microsoft/
Novartis)

Case studies

Diverse applications

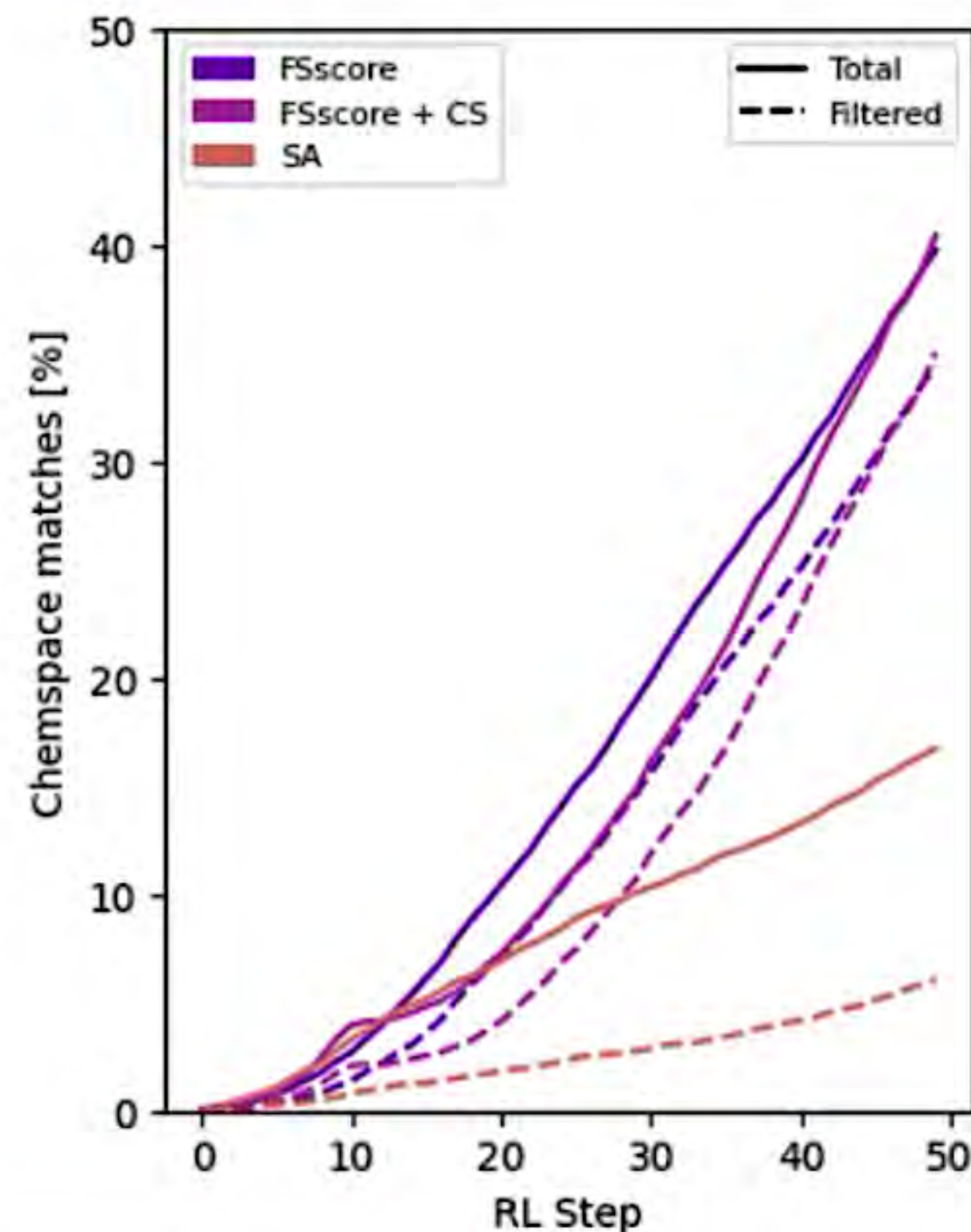
- Chiral molecules
- Two “hard to synthesize” vs “easy to synthesize” test
- PROTAC degrader molecules (labelled by human expert)
- Natural products (COCONUT) vs known drugs
- DRD2 docking molecules from de novo generation agent





Delivering Discovery Solutions®

Take closest
make-on-demand molecule
for pair to fine-tune scorer.



40% commercially available molecules
(ChemSpace) without human labelling.

nature machine intelligence

Review article

Machine learning-ai molecular design

Received: 19 July 2023

Accepted: 24 April 2024

Published online: 18 June 2024

Check for updates

Yuanqi Du^{1,2}
Charles Harris^{1,2}
Philippe Schwallier^{1,2}

Machine learning-aided drug discovery by single architecture of medicinal chemistry that can act as a central hub to have addressed into a series of comprehensive understanding using machine learning such as the and optimization of practical applications experiment efforts. Finally and empirically highlight future realistic drug

Alex Zhavoronkov • 1st
Founder and CEO at Insilico Medicine
1w •

Anyone interested in starting or ramping up in generative chemistry should read this paper and share it with their friends, students, and unrelated medicinal and computational chemists to help them get into the field. This is probably the most concise and useful review of generative AI in small molecule chemistry, which takes 20-30 min for experts to read but provides very deep insights (11 pages, mostly figures). If you want to see what published generative models and platforms worked experimentally for which targets and with what hit rates, you can print it out and get a cheat sheet. Of course, the most commercially tractable examples are rarely published but from what I can see - pretty much every published example was captured and scrutinized. This group reconstructed the timeline and did a gargantuan amount of work that an army of modern LLM agents would not be able to complete with a high degree of accuracy since many of the numbers are in image format and require additional processing.

I was planning to write a review like this one day, but this group beat me to this and did a way better job as I saw some of the models I was not aware of. [Yuanqi Du](#), [Arian Jamasb](#), [Jeff Guo](#), [Tianfan Fu](#), [Charlie Harris](#), [Yingheng W. Chen](#), [Ru Duan](#), [Pietro Lio](#), [Philippe Schwallier](#), [Tom Blundell](#)

Review article

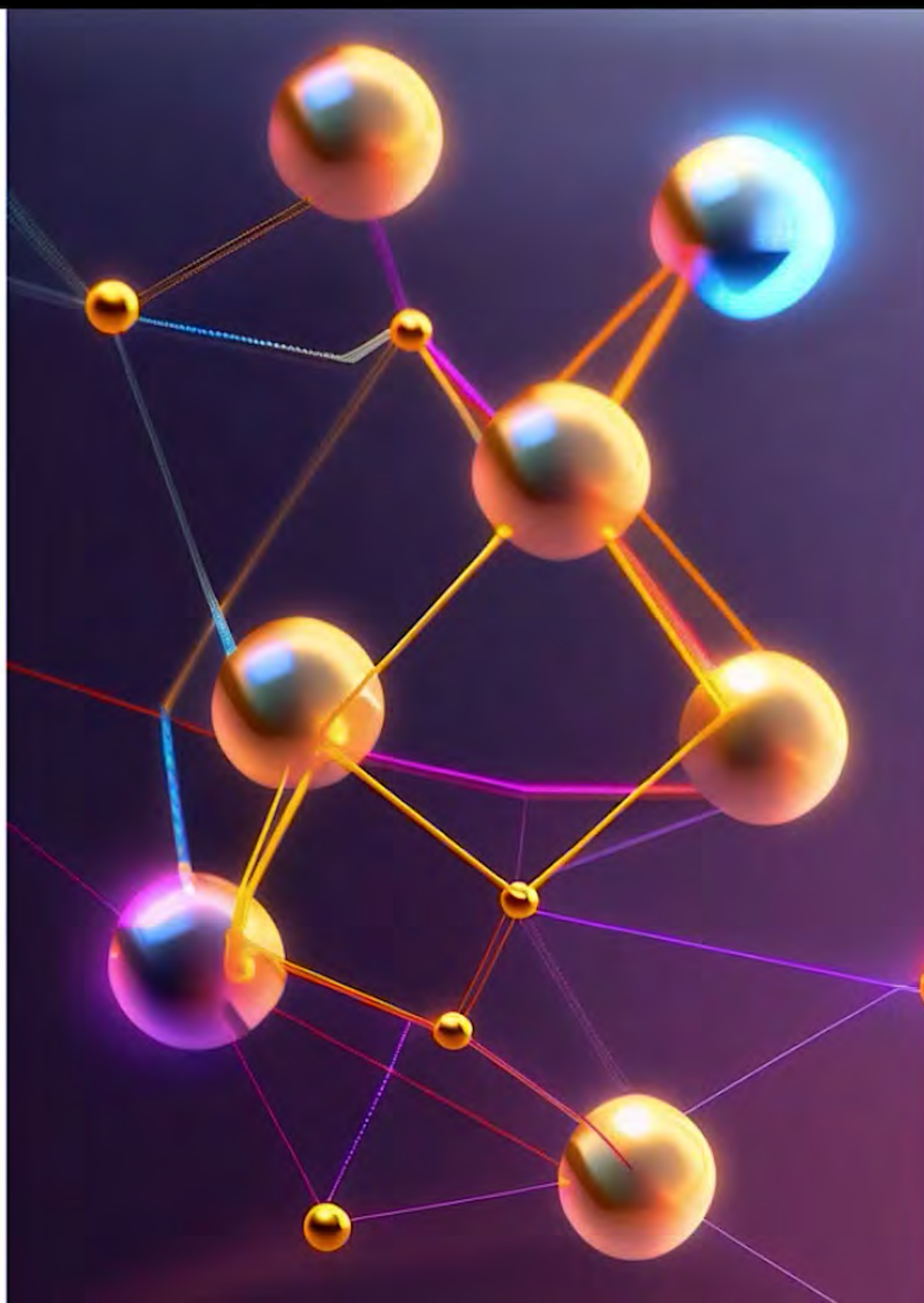
<https://doi.org/10.1038/s42256-024-00643-5>

Table 4 | Experimentally validated small-molecule generative design case studies

Model	Input	Output	Design task	Target	Hit rate	Outcome	Publication year
Distribution learning							
LSTM-RNN ¹	SMILES	SMILES	De novo	RXR	4/5 (80%)	nM agonist	2018
LSTM-RNN ²	SMILES	SMILES	De novo	RXR	2/4 (50%)	µM agonist	2018
g	JAK1				7/7 (100%)	nM inhibitor	2021
LXR					17/25 (68%)	µM agonist	2021
ROy					3/3 (100%)	µM agonist	2021
FLT-3					1/1 (100%)	µM inhibitor	2022
g	CDK8				9/43 (21%)	nM inhibitor	2022
Bacteria					0/1 (0%)	µM inhibitor	2022
DDR1					2/2 (100%)	nM inhibitor	2021
MERTK					15/17 (88%)	µM inhibitor	2022
PI3Ky					3/18 (17%)	nM inhibitor	2023
g	TBK1				1/1 (100%)	nM inhibitor	2023
CDK2					17/23 (74%)	nM inhibitor (MC) ³	2023
Nurr1y					2/6 (33%)	nM inhibitor	2023
PPARy					2/2 (100%)	µM agonist	2023
Kinases							
JAK3					1/1 (100%)	µM inhibitor	2018
DDR1					4/6 (67%)	nM inhibitor ³	2019
DDR1					4/6 (67%)	nM inhibitor	2021
EGFR					4/15 (27%)	nM inhibitor	2022
RPK1					4/8 (50%)	nM inhibitor ³	2022
re	CDK20				6/13 (46%)	nM inhibitor	2023
re	CDK8				1/1 (100%)	nM inhibitor ³	2023
re	SIK2				6/6 (100%)	nM inhibitor	2023
re	KOR				2/5 (40%)	µM antagonist	2023
re	PHD enzymes				1/1 (100%)	nM inhibitor ³	2024
re	NLRP3				0 ⁴	nM inhibitor ³	2024
re	Tuberculosis ClpP				1/6 (17%)	µM inhibitor	2024
re	KRAS				1/12 (8%)	µM inhibitor	2024
re	MGLL				1/3 (33%)	µM inhibitor	2024
g	PolR				4/6 (67%)	µM inhibitor ³	2024
re	TRK				Unknown ⁵	nM inhibitor ³	2024
re	Factor Xa				Unknown ⁶	µM inhibitor	2024
re	HAT1 and YTHDC1				0/2 and 0/3 (0%)	Both µM inhibitor ³	2024

Nature Machine Intelligence

<https://github.com/GuoJeff/generative-drug-design-with-experimental-validation>
45 studies with experimentally validated drug candidates



AdsMT: Multi-modal Transformer for Predicting Global Minimum Adsorption Energy

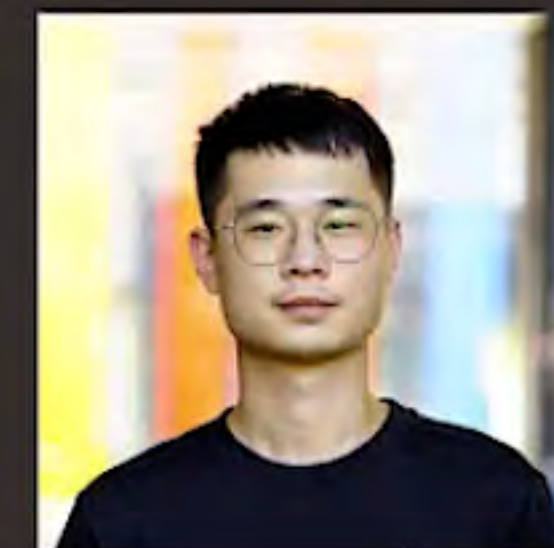
AdsMT: A multi-modal transformer for predicting global minimum adsorption energy

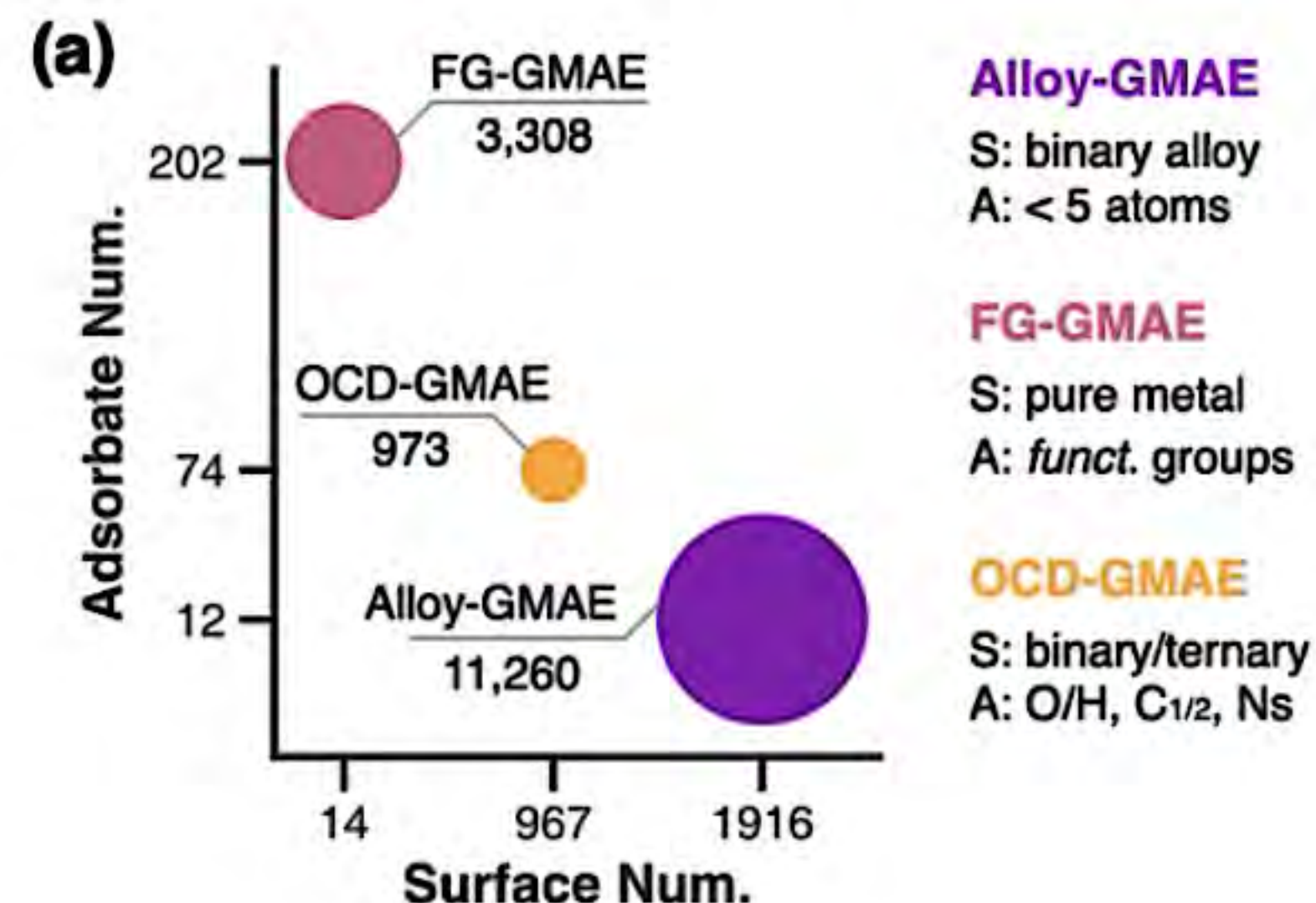
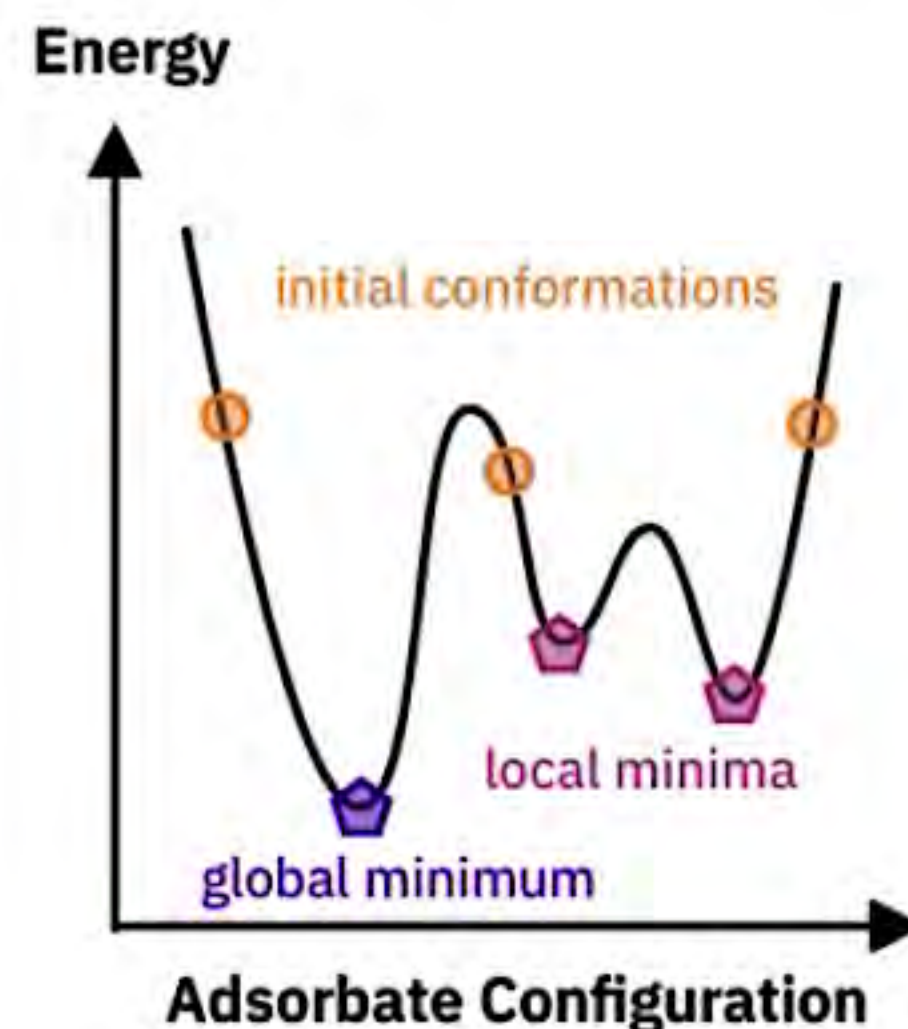
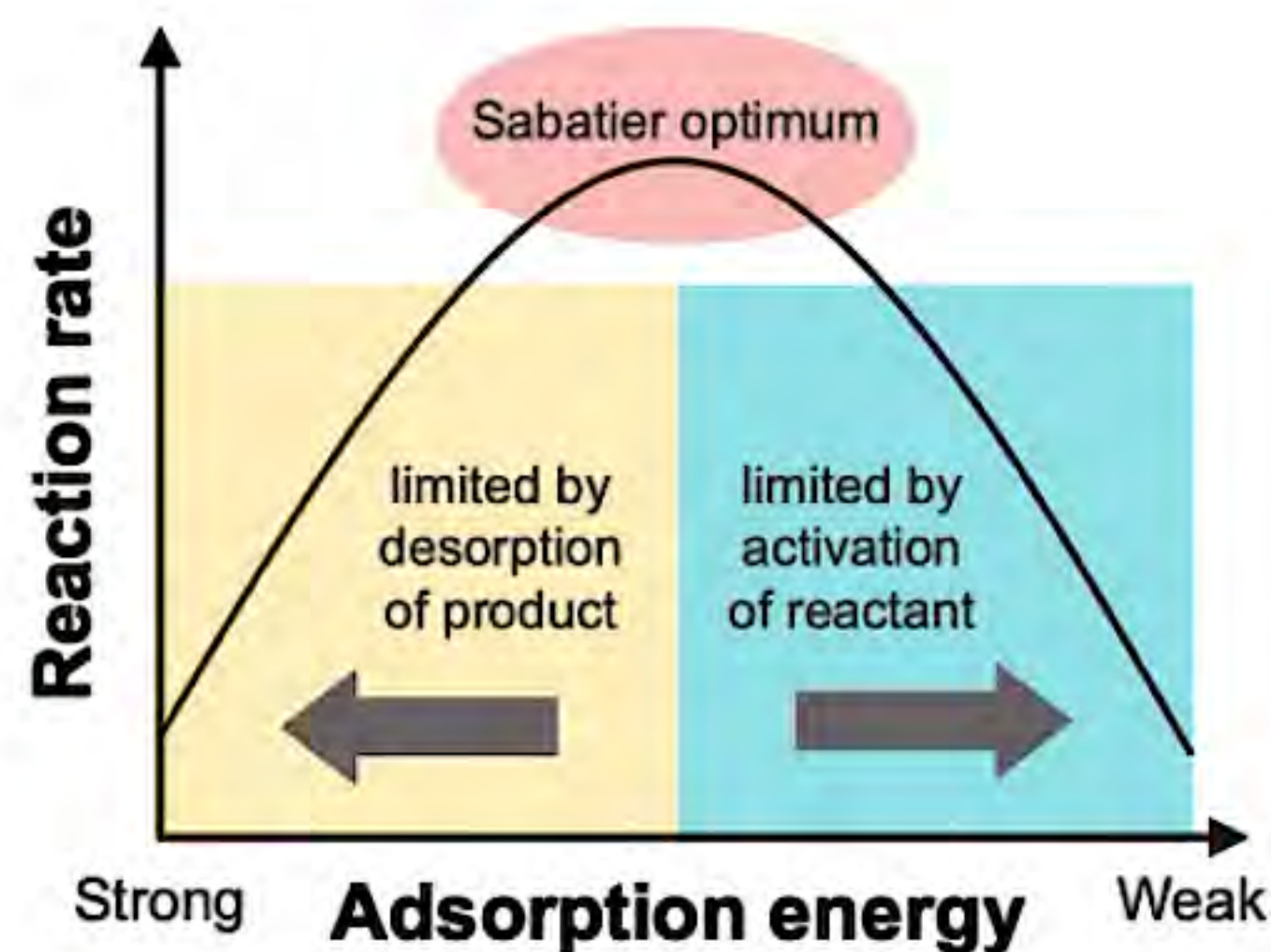
07 August 2024, Version 2

Working Paper

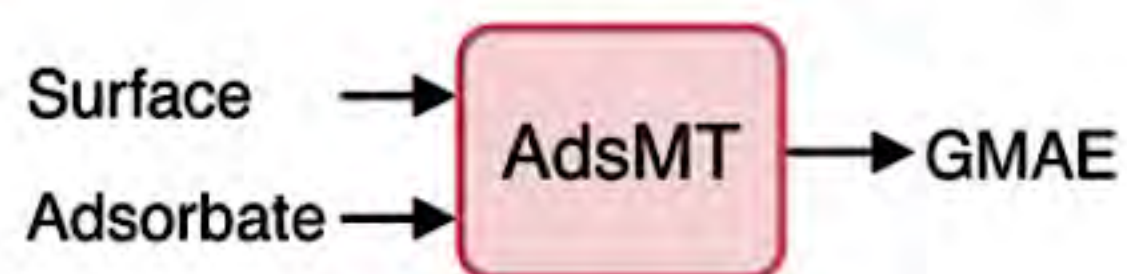
[Junwu Chen](#), [Xu Huang](#), [Cheng Hua](#), [Yulian He](#), [Philippe Schwaller](#)

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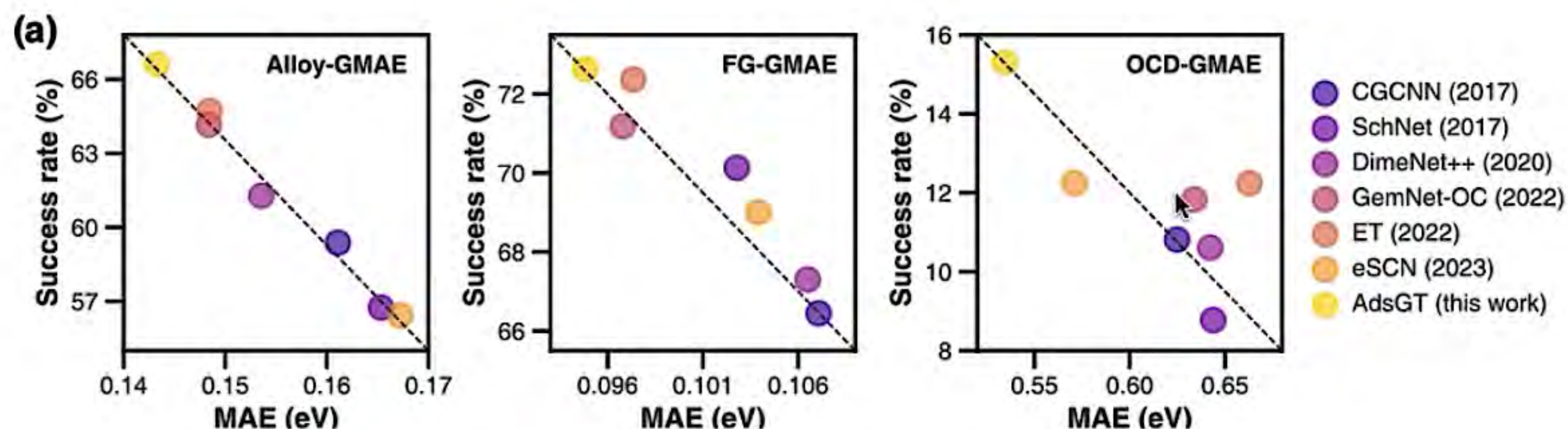
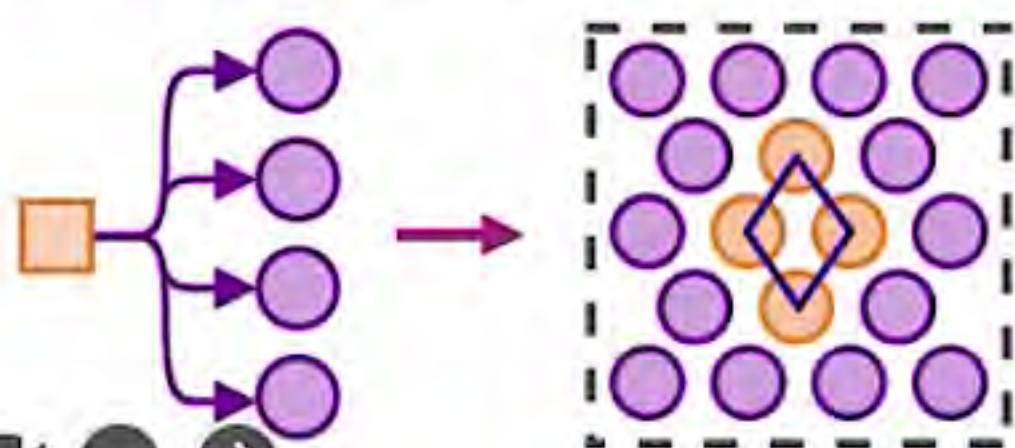


Multi-modal model for GMAE

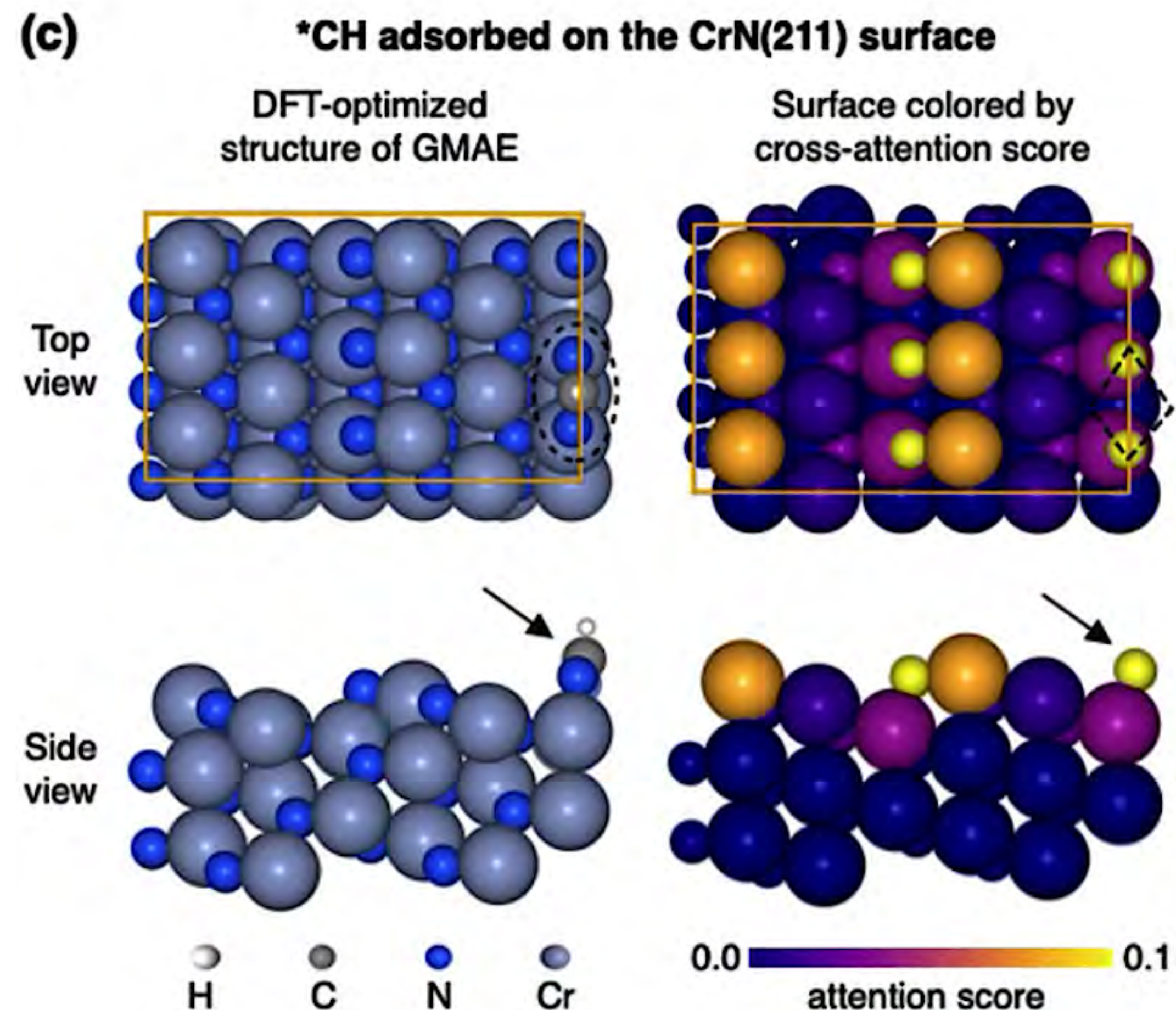
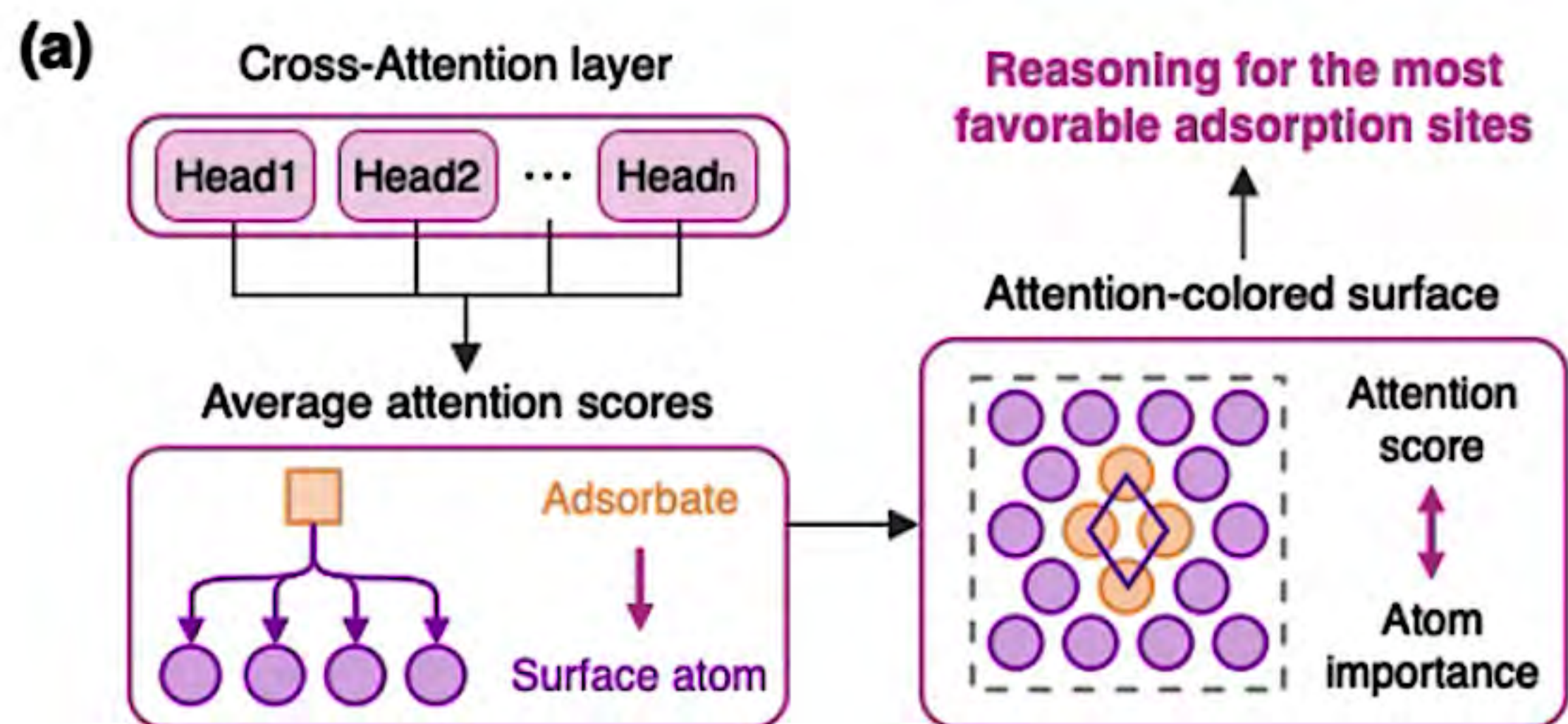


Attention for adsorption sites

Cross attention Atom importance



The **success rate** is the percentage of predicted GMAEs within **0.1 eV** of the DFT-computed ground truth GMAEs.



- **Transformers** capture the language of chemical reactions and more...
- LLM-Powered agents have the potential to **lower access barrier** to ML/computational tools.
- ChemCrow autonomously **solves generic tasks** in chemistry. Evaluation challenging.
- **Augmented Memory/Saturn** is a step towards sample efficient de novo design.
- **FSScore** for tunable and differentiable synthetic feasibility.



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Funding:



IBM Research



VANTAI



>10 nationalities — one team!
<https://schwallergroup.github.io>

Many thanks for your attention!



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Many thanks for your attention!



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