



香港中文大學

The Chinese University of Hong Kong



Auto-QChem: an automated workflow for the generation and storage of DFT calculations for organic molecules

Jingyi Liu

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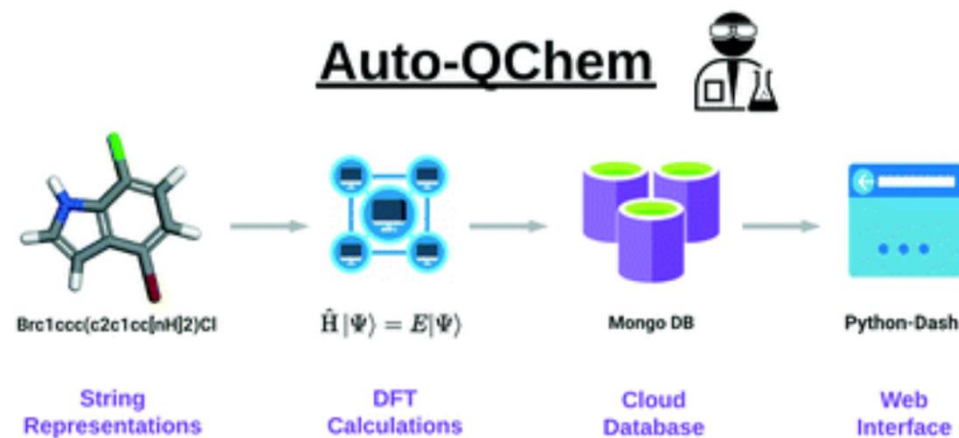
Outline

1. Background

2. Auto-Qchem workflow

3. Applications

4. Discussion



Research Background

Research Background

- The application of machine learning models in organic chemistry requires effective representations of chemical structures.
- ML models trained with chemical descriptors are more interpretable than molecular fingerprints and other representations.
- High throughput DFT presents a significant barrier to experimental chemists.
- Previous DFT automation packages are mainly designed for material science rather than small organic molecules.

Auto-Qchem

- An automatic, high-throughput and end-to-end DFT calculation workflow
- Computes chemical descriptors for organic molecules

[1] <https://github.com/PrincetonUniversity/auto-qchem>;

[2] <https://princetonuniversity.github.io/auto-qchem>

[3] <https://autoqchem.org>

Auto-Qchem workflow

➤ Features

1. Generate user-specified input.
2. HPC interface.
3. Automatic information extraction from DFT results.
4. Convenient storage and data accessing.

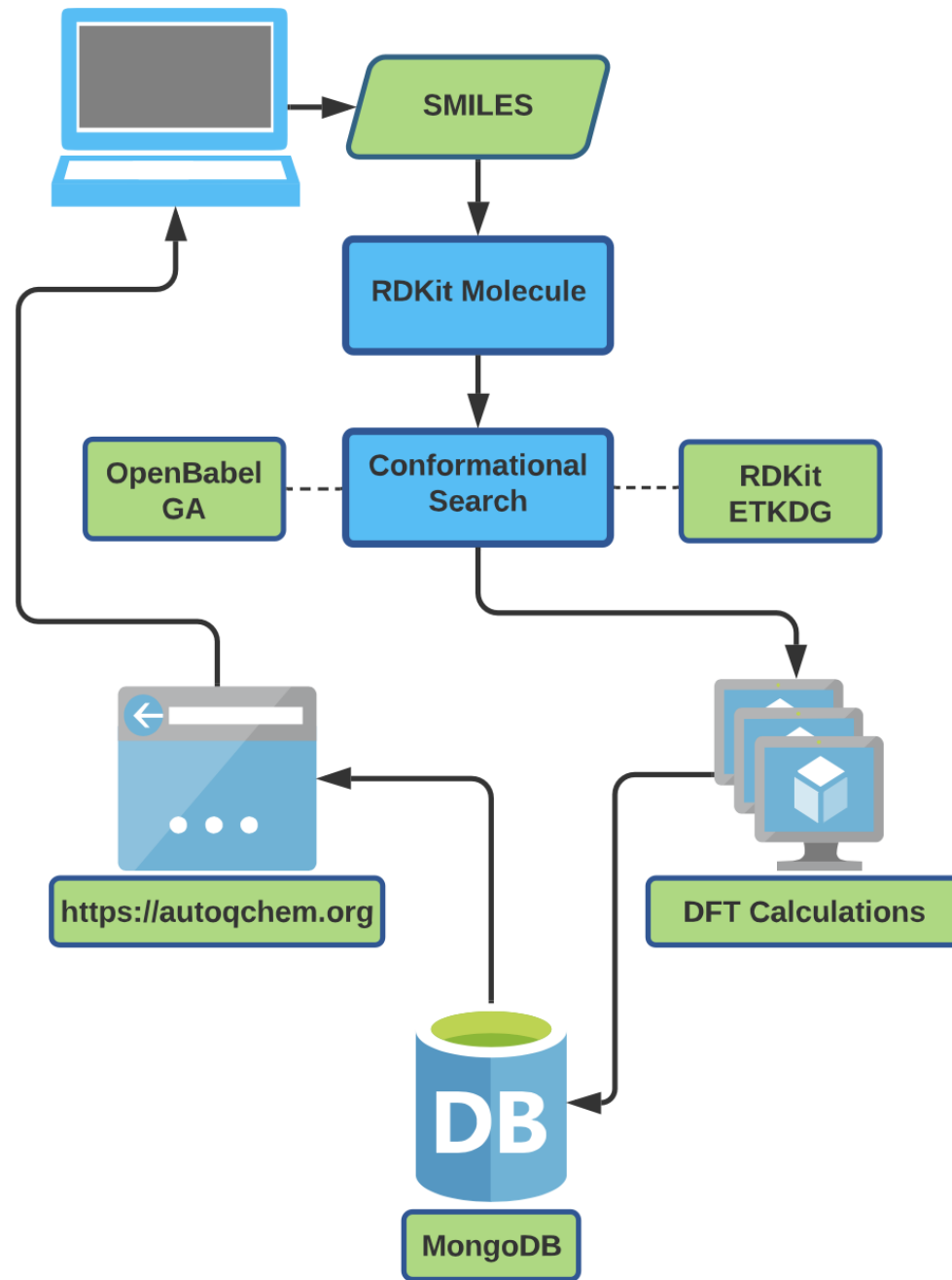


Fig. 1 Computational workflow of Auto-QChem.

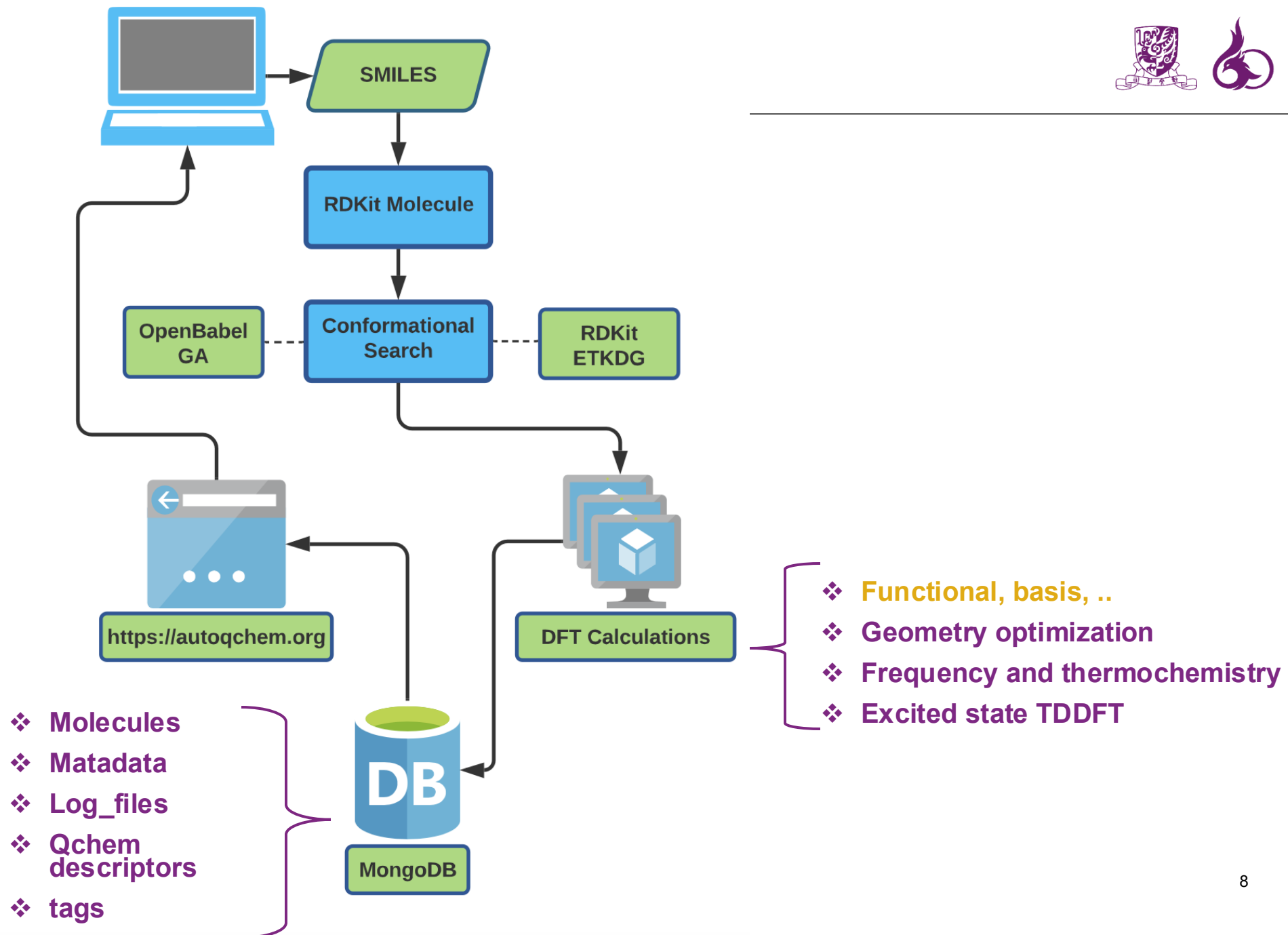


Fig. 1 Computational workflow of Auto-QChem.

➤ Features

- Avoid duplicate calculations

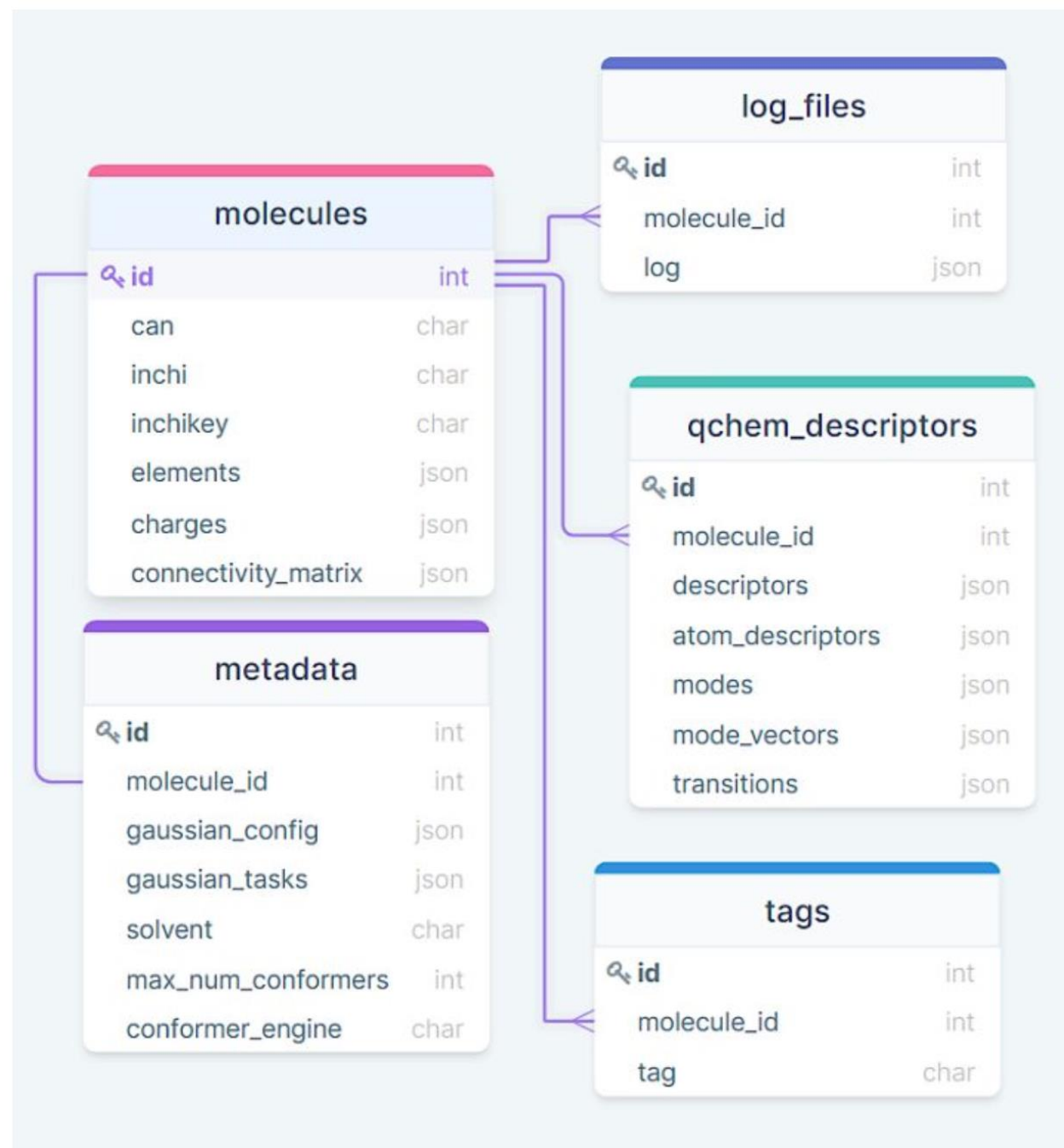


Fig. 2 Collection schema of Auto-QChem database.

➤ **Features:** Queries and data retrieval

➤ 1692 molecules in total






QUERY DOWNLOAD DESCRIPTORS EXPORT MOLECULE LIST

DATASET TAG: deoxyF_AM2 (32 molecules) SOLENT: TETRAHYDROFURAN FUNCTIONAL: M062X

BASE SET: ALL SUBSTRUCTURE: C[0;H1] SMILES: SMILES string

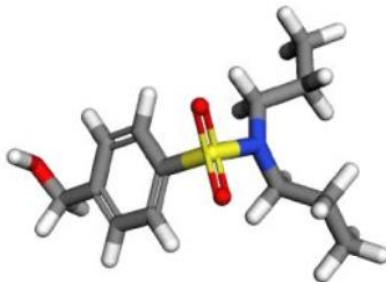
QUERY

Found 32 molecules

image	can	solvent	theory	light_basis_set	heavy_basis_set	num_conf/max_conf	detail
	CCCN(CCC)S(=O)(=O)c1ccc(CO)cc1	TetraHydroFuran	M062X	Def2TZVP	LANL2DZ	20/30	detail
	CCN(CCO)c1ccc(N=N/c2ccc([N+]=[O-])cc2Cl)cc1	TetraHydroFuran	M062X	Def2TZVP	LANL2DZ	29/30	detail
	COC(=O)[C@@H]1C[C@@H](O)CN1C(=O)OC(C)C	TetraHydroFuran	M062X	Def2TZVP	LANL2DZ	8/30	detail
	COC(=O)[C@@H]1C[C@@H](O)CN1C(=O)OC(C)C	TetraHydroFuran	M062X	Def2TZVP	LANL2DZ	10/30	detail
	COc1ccc2c(c1)c(CCO)c(C)n2C(=O)c1ccc(Cl)cc1	TetraHydroFuran	M062X	Def2TZVP	LANL2DZ	22/30	detail

CCCN(CCC)S(=O)(=O)c1ccc(CO)cc1

2: G = -744410.4 + 0.10 kcal/mol



1 3 5 7 9 11 13 15 17 19

number_of_atoms	charge	multiplicity	dipole	molar_mass	molar_volume	electronic_spatial_extent	E_scf	zero_point_corrector
39	0	1	6.46	271.37	2275.5	6329.46	-1186.58	0.33

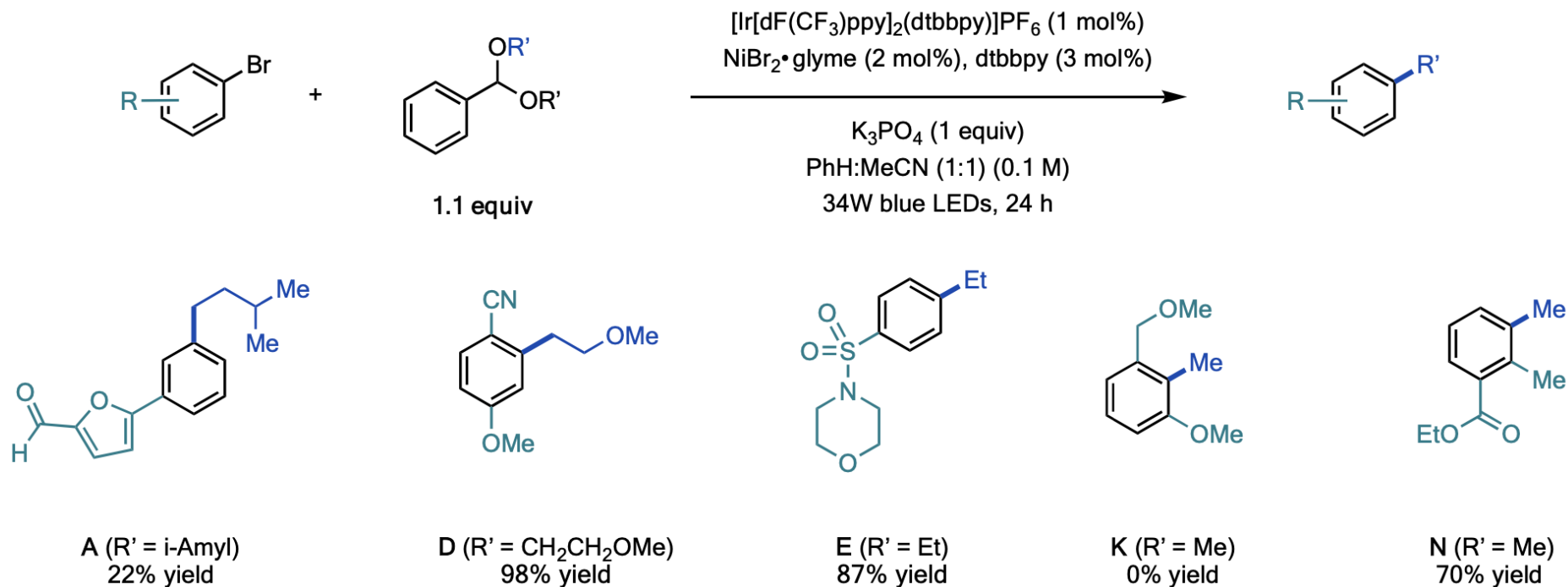
Atom-Level Descriptors

atom_idx	label	X	Y	Z	VBur	Mulliken_charge	APT_charge	NPA_charge	NPA_core	NPA_valence	NPA_Rydberg
0	C	-0.33	0.95	-0.77	0.42	-0.44	0.05	-0.6	2	4.6	0.01
1	C	-0.33	0.72	-0.76	0.56	-0.22	0.08	-0.41	2	4.4	0.01
2	C	-0.43	0.37	-0.51	0.64	-0.33	0.39	-0.21	2	4.2	0.02

Fig. 3 Query view (left) and the molecule view (right) of the web interface. The molecule view is a snapshot while viewing the second lowest energy conformation in 3D.

1. Substrate scope design in Ni/ photoredox methodology development

- Ni/ photoredox catalyzed alkylation reaction of aryl halides using acetals as alcohol-derived aliphatic radical sources.
- Data set: 2683 aryl bromides, with 168 electronic and steric features,
- 95 of 168 features were employed for clustering.



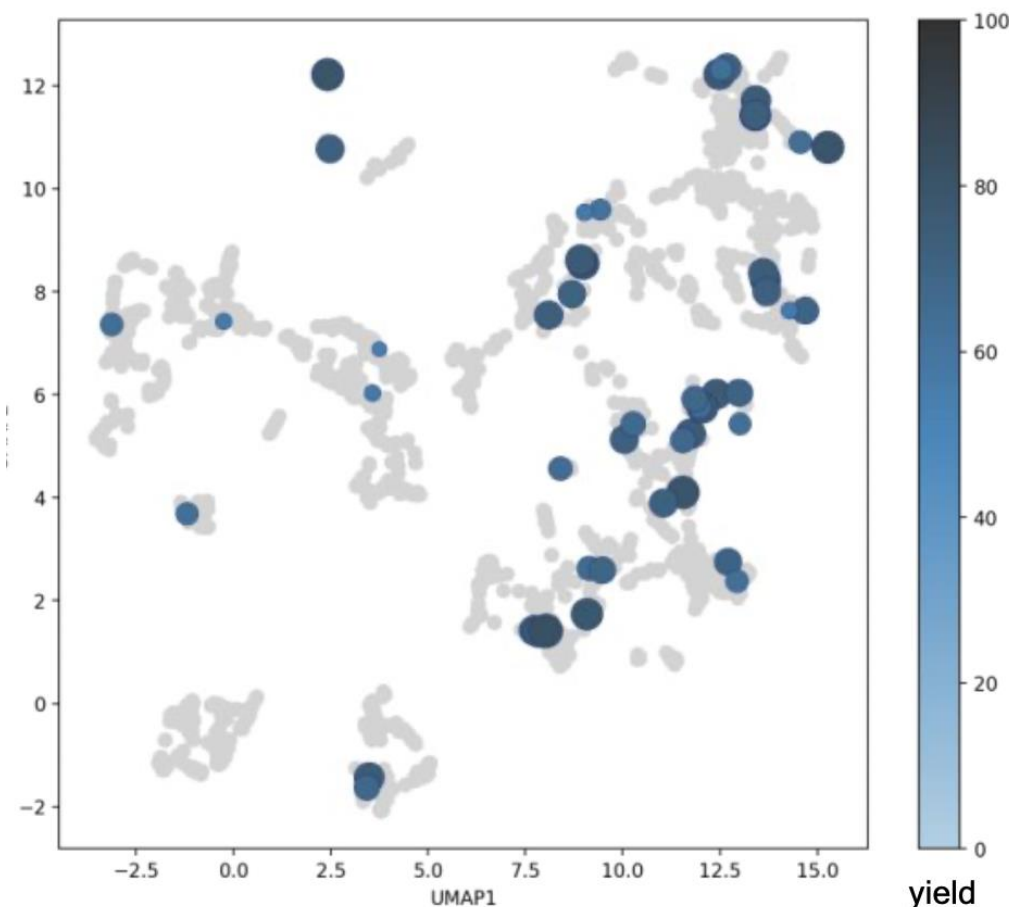
❖ Applications



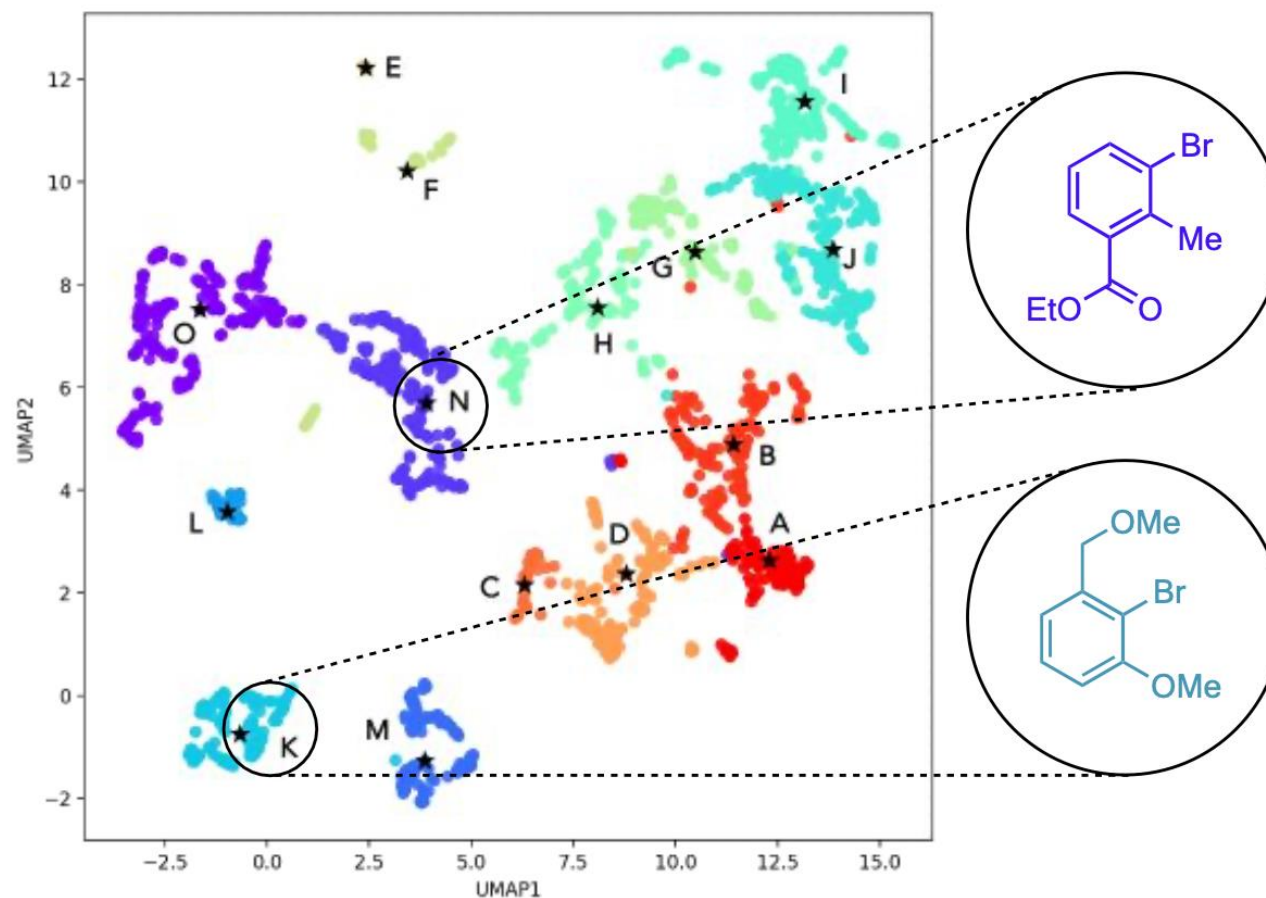
1. Substrate scope design in Ni/ photoredox methodology development

- Electronegativity of the aryl bromides was highly correlated with yield.
- Generalized additive model (GAM) trained on electronegativity outperformed ML models trained on literatures

(a). Substrates from literature and yields



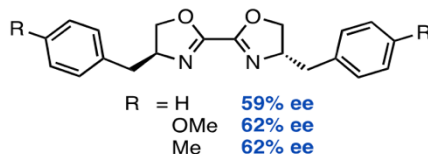
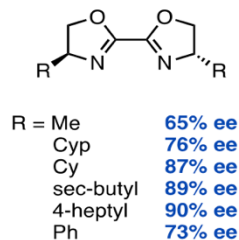
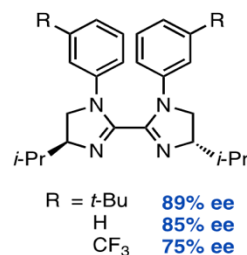
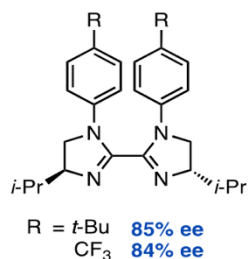
(b). Clustering results and selected aryl bromides



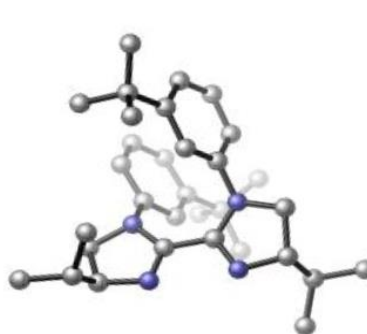
2. Ligand parametrization and enantioselectivity prediction in nickel catalysis

- Previous used ligand: Bioxazoline (BiOx)
 - Good enantioselectivity, low yield
- Optimal ligand: chiral biimidazoline (Bilm) : time-consuming to discover

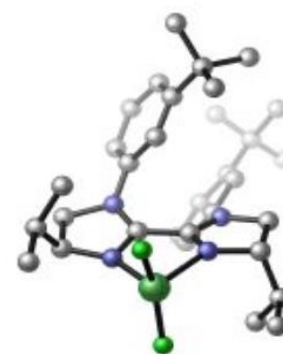
(a). Model reaction system and representative examples of ligands tested.



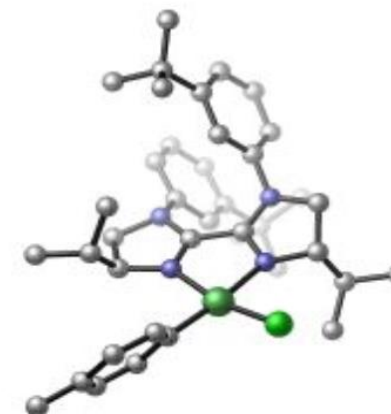
(b). Ligand environment models.



free ligand



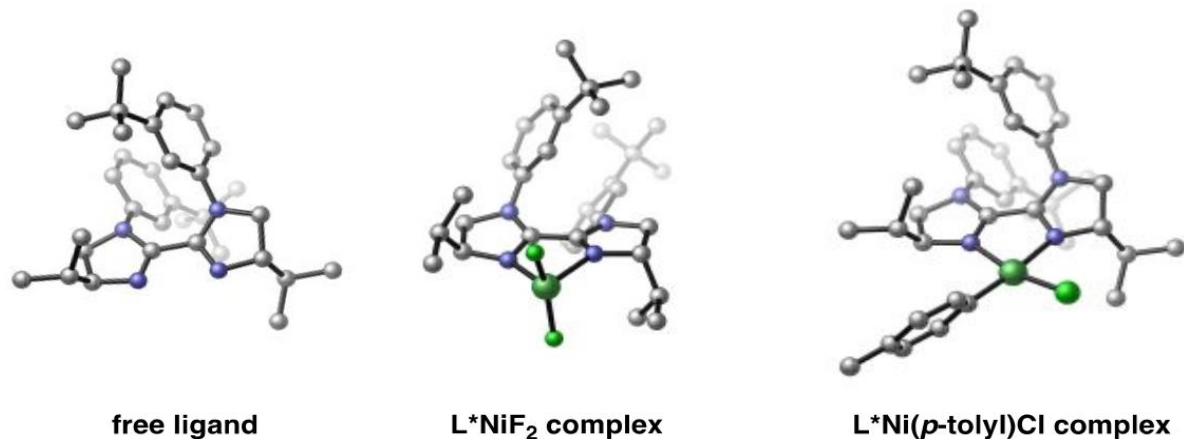
L*NiF₂ complex



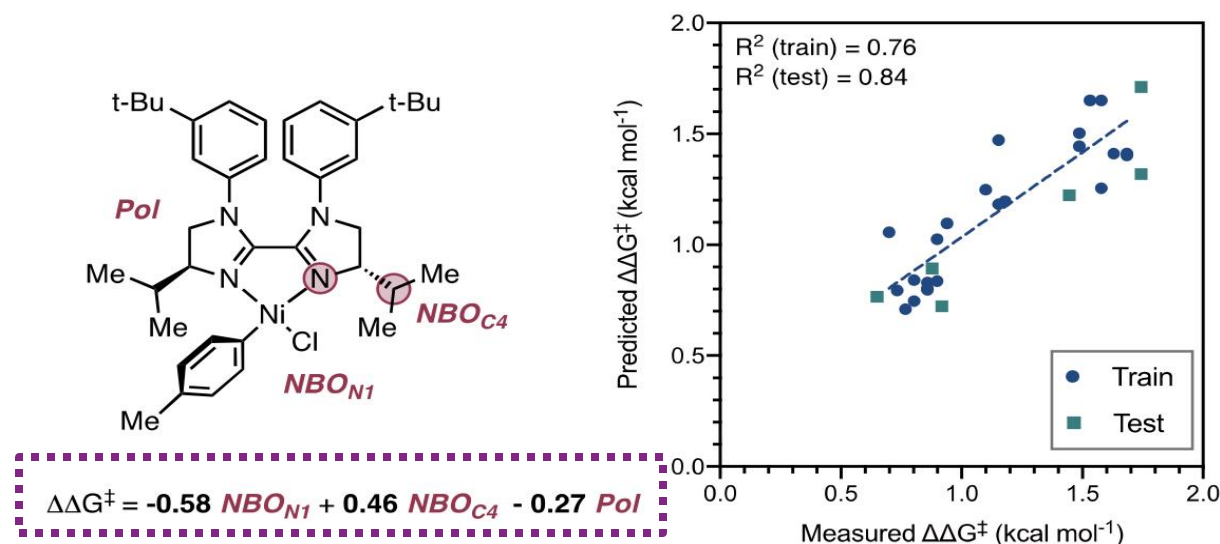
L*Ni(*p*-tolyl)Cl complex

2. Ligand parametrization and enantioselectivity prediction in nickel catalysis

(b). Ligand environment models.



(c). Regression modeling for L*Ni(*p*-tolyl)Cl with DFT-derived features.



- Hypothesis: computed features depend on ligand environment.
- Manually generated conformers ← Auto-Qchem cannot handle transition metal complexes
- Conclusion: electronic, rather than steric attributes of Bilm ligands govern the enantioselectivity of this reaction

3. Reaction condition optimization via Bayesian optimization

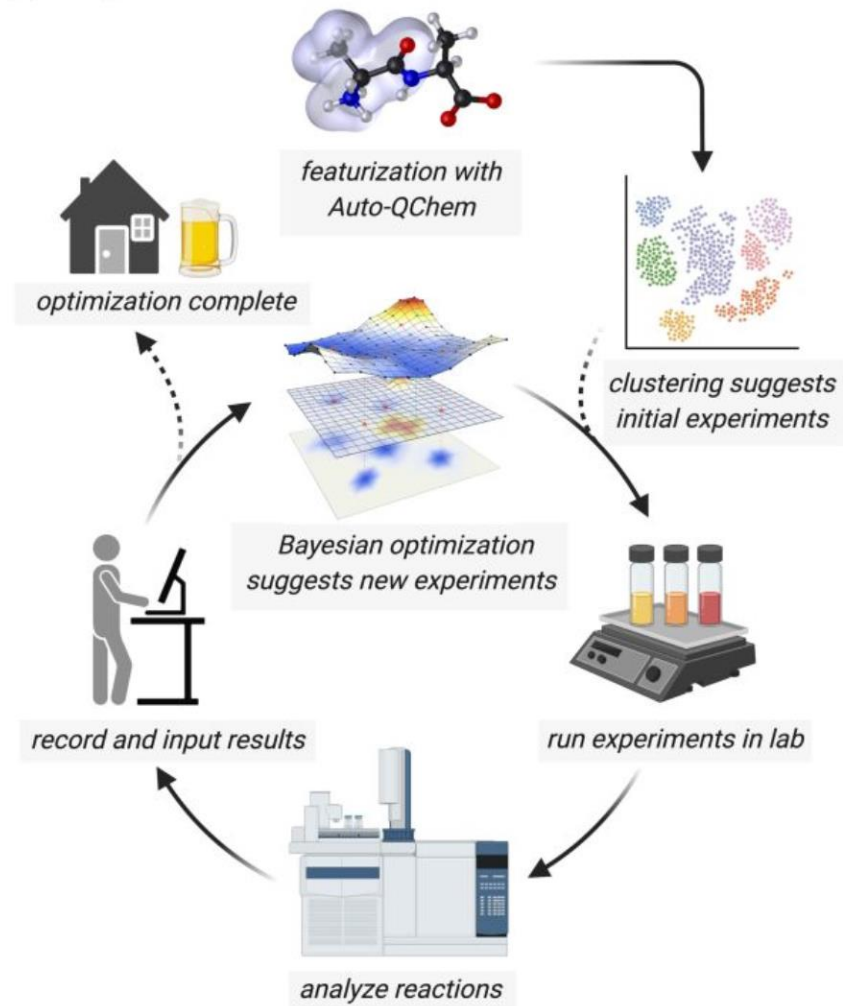
- Traditional reaction optimization is slow and resource-intensive, relying on trial-and-error or costly methods, with limited exploration of chemical space.
- Bayesian optimization, a sequential design algorithm for global optimization of black-box functions, in efficient reaction condition optimization.
- EDBO (Experimental Design via Bayesian Optimization)

❖ Applications

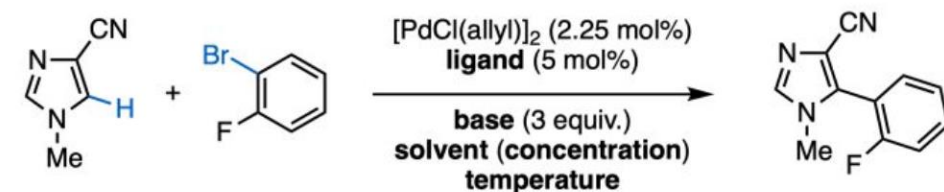


➤ 3. Reaction condition optimization via Bayesian optimization

(a). the general optimization workflow of EDBO.



(b). HTE study of Pd-catalyzed C-H arylation of imidazoles.



PCy ₃	CgMePPh	PPh ₃	GorlosPhos
XPhos	BrettPhos	<i>t</i> -BuPh-CPhos	PPh(<i>t</i> -Bu) ₂
PPh ₂ Me	PPhMe ₂	P(fur) ₃	JackiePhos

Ligand

KOAc	CsOAc	BuOAc	BuCN
KOPiv	CsOPiv	<i>p</i> -Xylene	DMAc

Base

Solvent

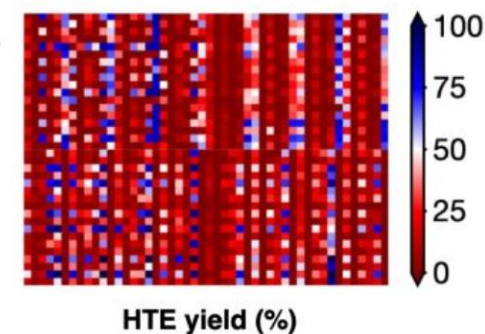
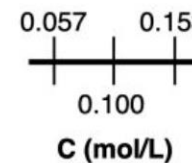
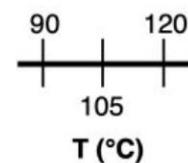
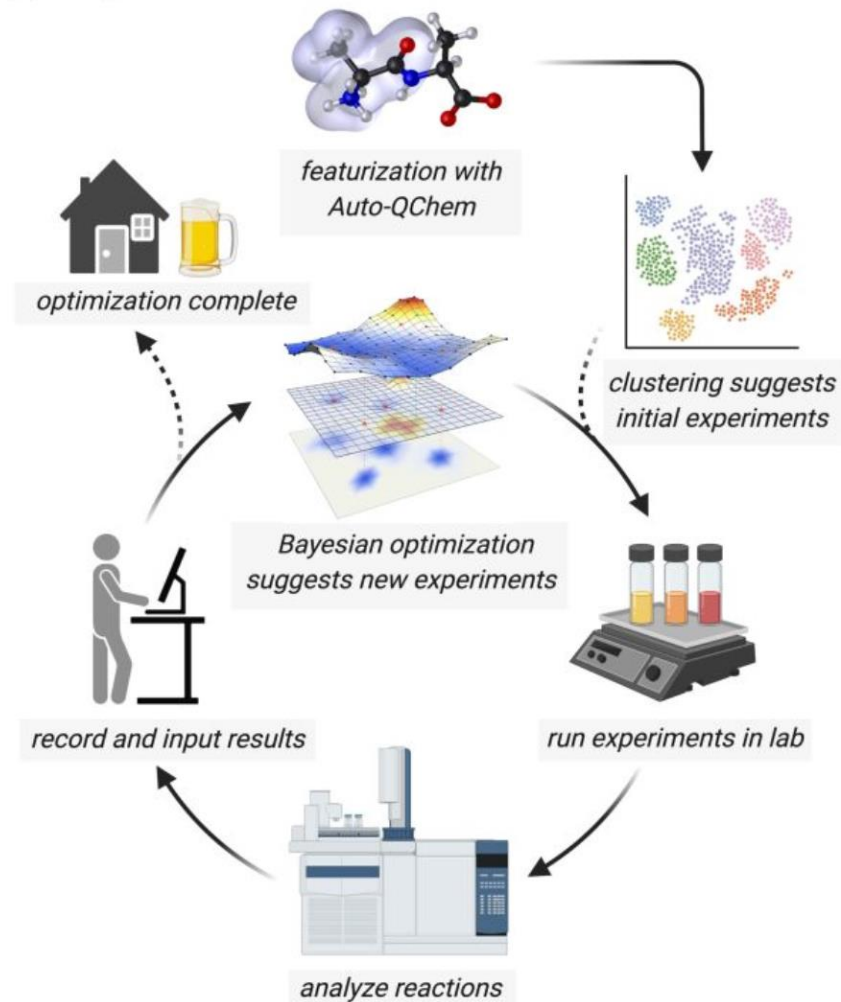


Fig. 6 Use case 3: reaction condition optimization via Bayesian optimization.

3. Reaction condition optimization via Bayesian optimization

(a). the general optimization workflow of EDBO.



EDBO (simulated 50 times) achieved a higher average performance within the first 15 experiments even with random initialization and found conditions with >99% yield 100% of the time!

Discussion

❖ Limits

- Cannot generate accurate conformers for transition metal complexes and molecules with non-canonical bonds.
- Lack of supports for other cluster schedulers

❖ Future perspective

- External packages support
- Data insufficiency

Questions? Comments?

Thank You