

# Towards Autonomous Computational Chemistry Research with AI and ML

Pavlo O. Dral

Xiamen University, China

Nicolaus Copernicus University in Toruń, Poland

Aitomistic, Shenzhen, China

9 September 2026

dr-dral.com

Behind the Paper

## Artificial intelligence makes accurate quantum chemical simulations more affordable

We have developed artificial intelligence-enhanced quantum mechanical method 1 (AIQM1), which can be used out of the box for very fast quantum chemical calculations with the accuracy of the gold-standard coupled-cluster method.

Published in Chemistry

Dec 02, 2021

 **Pavlo O. Dral**  
Professor, Xiamen University; Nicolaus Copernicus University in Toruń; Aitomistic

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*E<sub>SQM</sub>*

*E<sub>disp</sub>*

*E<sub>ex</sub>*



ChatGPT:  
Nov 30, 2022

<https://communities.springernature.com/posts/artificial-intelligence-makes-accurate-quantum-chemical-simulations-more-affordable>



learning (ML) – the very tool of AI we actually used in AIQM1), in school, I was fascinated by nature and was learning about it as much as I could. I quickly realized that it is impossible to learn everything I am interested in. The more I thought about it, the more obvious was to me that the human mind is just incapable of coping with all the information, which humanity generated, and this is likely to impose a natural barrier on how we can actually make a qualitative jump in our understanding of the world. Computers, in the long run, may overcome this barrier as they are not limited by biological constraints and can have access to all digitalized information. However, can computers by themselves make a qualitative jump in our understanding of nature? This is the first question.

Dec 2, 2021

<https://communities.springernature.com/posts/artificial-intelligence-makes-accurate-quantum-chemical-simulations-more-affordable>

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Later, already in the university doing my master degree, I have had to run many quantum chemical (QC) simulations. It was exciting at first to witness how a computer can predict correctly, e.g., the geometry of a water molecule, but after I run many such simulations, I realized that many of them are quite repetitive and the second question is: can computers by themselves perform all these simulations, analyze results and write papers?

Dec 2, 2021

<https://communities.springernature.com/posts/artificial-intelligence-makes-accurate-quantum-chemical-simulations-more-affordable>



**Protomia**

for Diels-Alder reaction between ethylene and 1,3-butadiene, what's the reaction energy?

reaction

**Ask about reaction energy**

RECENT ACTIVITY Hide

- Interpreting instructions
- Starting structure preparation
- Structure prepared
- Starting structure preparation
- Structure prepared

5 item(s)

**Protomia**

I've successfully completed the computational workflow for the Diels-Alder reaction between ethylene and 1,3-butadiene to form cyclohexene. All steps — from structure preparation through geometry optimization, frequency analysis, a **Autonomous simulation** aggregation — ran v

Here are the key results:

- Reaction: ethylene + 1,3-butadiene → cyclohexene
- Reaction Energy: -38.14 kcal/mol (exothermic)
- Reaction Enthalpy (298K): -33.82 kcal/mol
- Reaction Gibbs Free Energy (298K): -19.73 kcal/mol (spontaneous at room temperature)

| type                        | system/reaction                         | units    | coefficient | E        | H        | G        | method      | output_molecule       |
|-----------------------------|-----------------------------------------|----------|-------------|----------|----------|----------|-------------|-----------------------|
| System                      |                                         |          |             |          |          |          |             |                       |
| reactant                    | ethylene                                | hartree  | 1           | -78.572  | -78.5165 | -78.542  | B3LYP/6-31G | /aitomistic/hub/home/ |
| reactant                    | 1,3-butadiene                           | hartree  | 1           | -155.96  | -155.868 | -155.9   | B3LYP/6-31G | /aitomistic/hub/home/ |
| product                     | cyclohexene                             | hartree  | 1           | -234.592 | -234.438 | -234.473 | B3LYP/6-31G | /aitomistic/hub/home/ |
| Reactions/Relative energies |                                         |          |             |          |          |          |             |                       |
| reaction                    | ethylene + 1,3-butadiene -> cyclohexene | hartree  |             | -0.06078 | -0.05389 | -0.03144 |             |                       |
| reaction                    | ethylene + 1,3-butadiene -> cyclohexene | kcal/mol |             | -38.1393 | -33.8159 | -19.7303 |             |                       |
| reaction                    | ethylene + 1,3-butadiene -> cyclohexene | kJ/mol   |             | -159.575 | -141.486 | -82.5515 |             |                       |
| constant                    | hartree_to_kcal_mol                     | kcal/mol |             | 627.509  |          |          |             |                       |
| constant                    | hartree_to_kj_mol                       | kJ/mol   |             | 2625.5   |          |          |             |                       |

**Spreadsheet of reaction**

protomia\_demo

DA\_ethene\_butadiene

1-3-butadiene\_opt\_freq\_AIQM2

cyclohexene\_opt\_freq\_AIQM2

scr

README.md

cyclohexene\_opt\_freq\_AIQM2.inp

cyclohexene\_opt\_freq\_AIQM2.log

cyclohexene\_opt\_freq\_AIQM2.xyz

slurm.sh

slurm\_6655.err

slurm\_6655.out

ethene\_opt\_freq\_AIQM2

1-3-butadiene\_init.json

1-3-butadiene\_init.xyz

DA\_ethene\_butadiene.xlsx

README.md

cyclohexene\_init.json

cyclohexene\_init.xyz

ethene\_init.json

ethene\_init.xyz

mini\_paper.aux

mini\_paper.log

mini\_paper.out

mini\_paper.pdf

mini\_paper.tex

test

SHARED WITH ME

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?

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cyclohexene\_opt\_freq\_AIQM2.xyz

cyclohexene\_opt\_freq\_AIQM2.xyz

cyclohexene\_opt\_freq\_AIQM2.inp

mini\_paper.tex

mini\_paper.pdf

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AIQM2 Study of the Diels–Alder Reaction Between Ethene and 1,3-Butadiene

Computational Chemistry Report — Generated by Protomia

May 27, 2026

Abstract

The Diels–Alder reaction between ethene and 1,3-butadiene to form cyclohexene was investigated using the AIQM2 machine-learning quantum mechanical method. Geometry optimizations and harmonic frequency calculations were performed for all reactants and the product. The reaction is found to be strongly exergonic with  $\Delta G = -26.16 \text{ kcal mol}^{-1}$  and exothermic with  $\Delta H = -40.31 \text{ kcal mol}^{-1}$  at standard conditions (298.15 K, 1 atm), consistent with the formation of a stable six-membered ring via a concerted [4 + 2] cycloaddition mechanism.

1 Introduction

The Diels–Alder reaction is one of the most important pericyclic reactions in organic chemistry, providing a powerful and atom-economical route to six-membered carbocycles. Discovered by Otto Diels and Kurt Alder in 1928 (for which they were awarded the Nobel Prize in Chemistry in 1950), this [4 + 2] cycloaddition involves a conjugated diene and a dienophile reacting in a concerted, suprafacial manner to form a cyclohexene derivative [1].

The prototypical example—the reaction between 1,3-butadiene (the diene) and ethene (the dienophile)—serves as a textbook illustration of the reaction:

$$\text{C}_4\text{H}_6 + \text{C}_2\text{H}_4 \longrightarrow \text{C}_6\text{H}_{10} \quad (1)$$

Despite its simplicity, this reaction is of fundamental interest for benchmarking computational methods, as it involves significant changes in bonding, conjugation, and strain upon cyclization.

2 Computational Methods

All calculations were performed with the AIQM2 method, a machine-learning-enhanced quantum mechanical approach that combines the efficiency of semi-empirical methods with the accuracy of

Protomia

(exergonic) under standard conditions. The reaction is strongly enthalpically driven ( $\Delta H = -40.31 \text{ kcal/mol}$ ), with the entropy penalty from combining two molecules into one partially offsetting the favorable enthalpy.

All optimized geometries, frequencies, and thermochemical data have been saved to the project directory (protomia\_demo/DA\_ethene\_butadiene, along with a summary spreadsheet (DA\_ethene\_butadiene.xlsx) and a README.md. These results can be reused for follow-up work such as transition-state searches or solvent-effect studies without re-running the optimizations.

now write a mini-paper in latex and make a pdf

I'll write a mini-paper in LaTeX about this Diels–Alder reaction and compile it to PDF. Let me first check the project directory for any existing files, then create the LaTeX document.

> Ran 1 action

Starting step 1: listing the project directory contents to see existing results.

> Ran 1 action

Updated while you were away

Step 1 done. Found the project directory with all species calculation folders and the summary spreadsheet. Now moving to step 2: writing the LaTeX mini-paper.

> Ran 1 action

Step 2 done — the LaTeX mini-paper has been written. Now moving to step 3: compiling the LaTeX document to PDF using pdflatex.

> Ran 3 actions

Step 3 done — the LaTeX compilation succeeded and the PDF is ready.

Here's a summary of what was produced:

- LaTeX source: mini\_paper.tex
- Compiled PDF: mini\_paper.pdf (3 pages, ~197 KB)

The mini-paper includes:

- An abstract summarizing the AIQM2 thermodynamic findings
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- A Results and Discussion section with a formatted table of  $\Delta E$ ,  $\Delta H$ , and  $\Delta G$ , plus analysis of the enthalpy vs. entropy balance
- Conclusions and a short bibliography

All cross-references (table label, citations) were resolved cleanly in the two-pass compilation. The paper is ready for reading or further editing if you'd like to add figures, additional discussion, or refine any sections.

copy

+ mini\_paper.pdf

Press Enter to send message

dr b

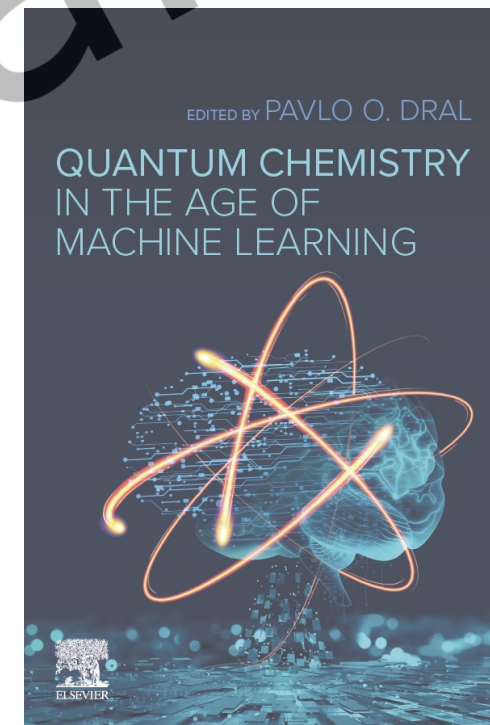
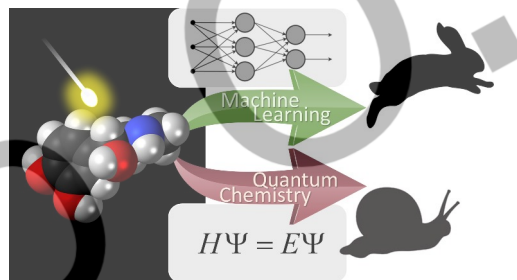




Professor

Email: [dral@xmu.edu.cn](mailto:dral@xmu.edu.cn)

**Research Areas:** AI, AI agents, autonomous research, quantum chemistry, dynamics, excited states, reaction mechanisms, theoretical spectroscopy



- 2025-Present : Aitomistic, co-founder
- 2024-Present : Nicolaus Copernicus University in Toruń, Professor
- 2021-Present : Xiamen University, Full Professor
- 2019-2021 : Xiamen University, Associate Professor
- 2013-2019 : Max-Planck-Institut für Kohlenforschung, Postdoc
- 2008-2013 : University of Erlangen-Nürnberg, M.Sc. & Ph.D.
- 2004-2010 : National Technical University of Ukraine “KPI”, B.Sc. & M.Sc.

### ✓ Selected papers:

- AI-DFT:** *Adv. Sci.* **2024**, 2408239
- AI platform:** *JCTC* **2024**, 20, 1193 | **2026**, 22, 6557
- AI-quantum dynamics:** *Nat. Commun.* **2022**, 13, 1930
- AI-quantum mechanics:** *Nat. Commun.* **2021**, 12, 7022
- AI-excited states:** *Nat. Commun.* **2026**, 17, 4949

Group website: [dr-dral.com](http://dr-dral.com)





# CHEMISTRY\WORLD

NEWS

## AI agents set to democratise computational chemistry

BY  JULIA ROBINSON | 2 JANUARY 2026

Aitomia

Status: [Publicly available](#) since 11 May 2025.

What it does: Aitomia assists researchers in all stages of their work, from setting up the calculations, providing possible options, automatically performing the simulations, and then helping to analyse the results in an understandable and reusable format. It supports a wide-array of AI-driven atomistic and quantum chemistry methods enabling essential computations such as energy calculations, geometry optimisations, molecular dynamics, thermochemistry, reaction and spectra simulations.

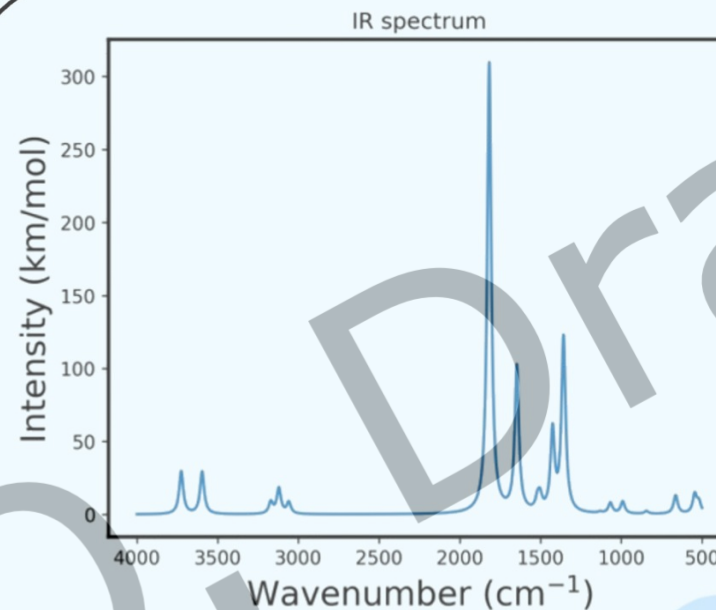
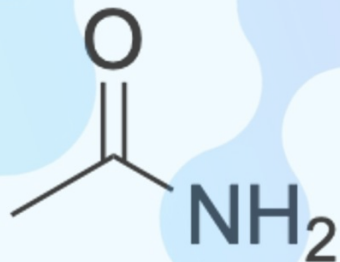
While we'll have to wait to find that out, it's clear that significant change is already afoot. 'I'm pretty sure that in the future, people will wonder how people previously did quantum chemical simulations,' says Dral. 'Like we are now astonished at how people built the pyramids in Egypt.'

I used Aitomistic for teaching Applied Quantum Chemistry ... It was amazing and students liked it so much.

By one of the independent professors



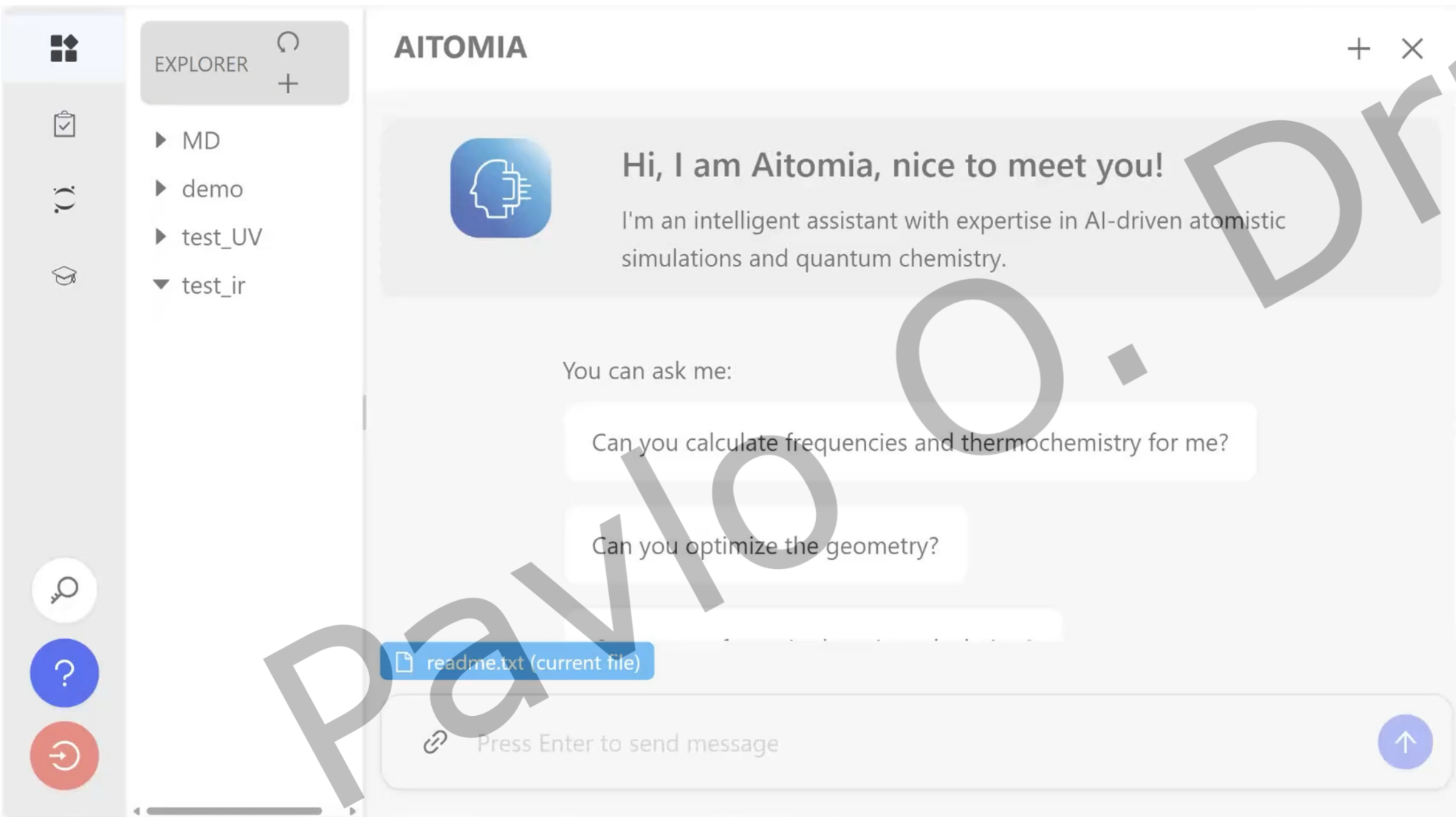
Simulate IR spectrum for  
this molecule



Sure!



Hello! I am *Aitomia*



\* Aitomia is the research framework introduced in 2025; **Protomia** is the modern architecture

<https://atom.xmu.edu.cn> (available for free when registered with an academic email)

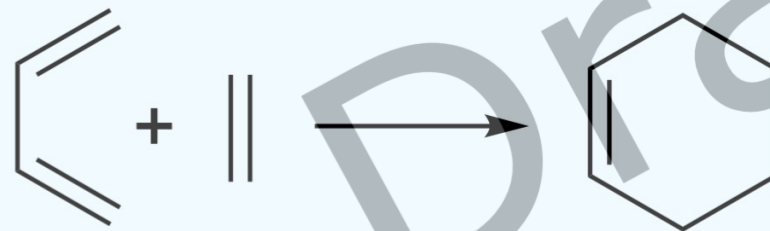
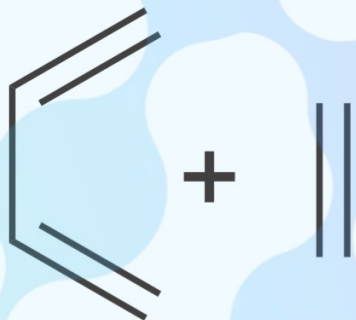
<https://aitomistic.xyz> (registration and basic use free for everyone)



# Aitomia: Intelligent Assistant for Aitomistic (AI+atomistic) and QM Simulations



Simulate this reaction

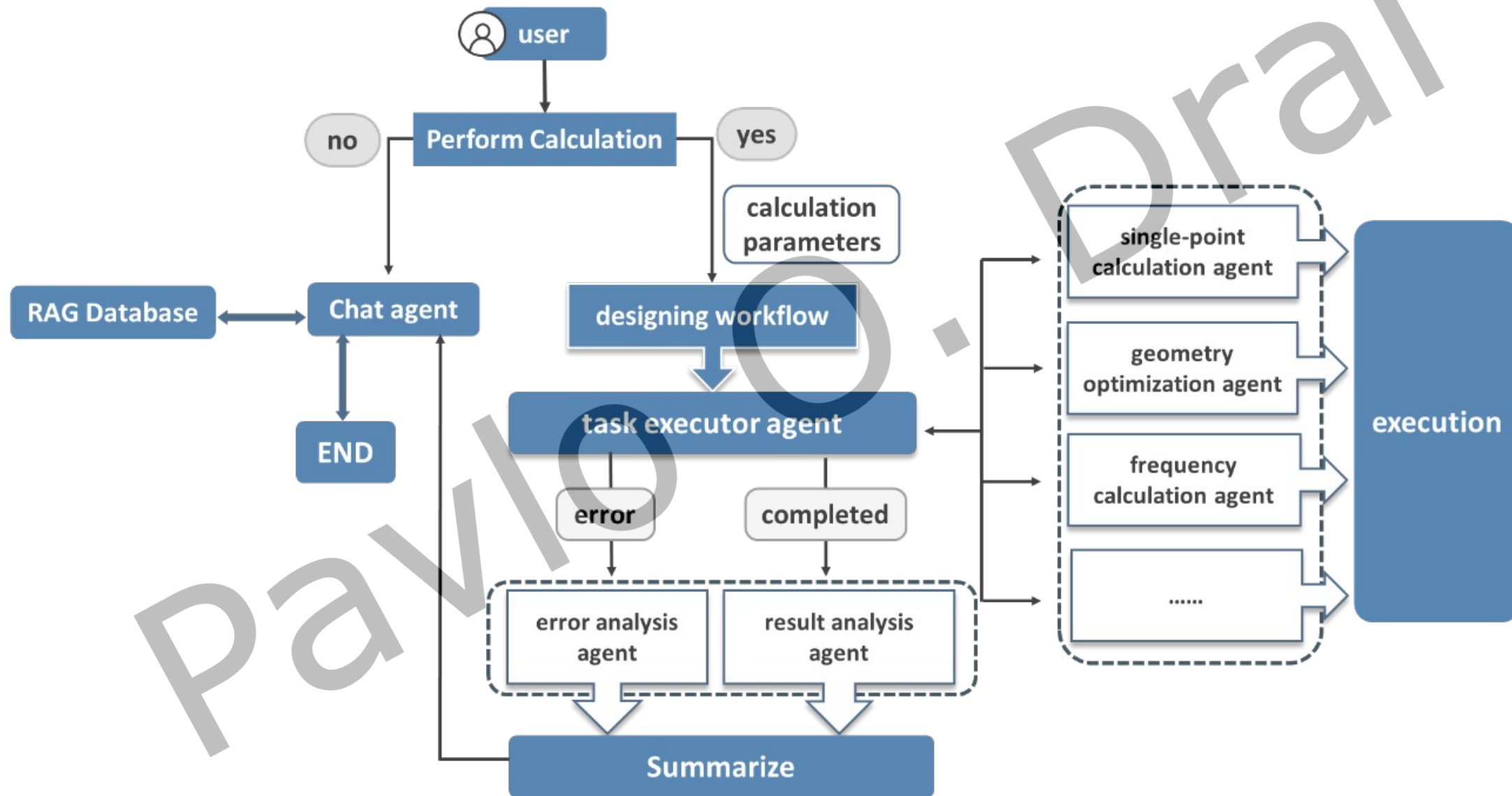


$\Delta G = -25.7$  kcal/mol

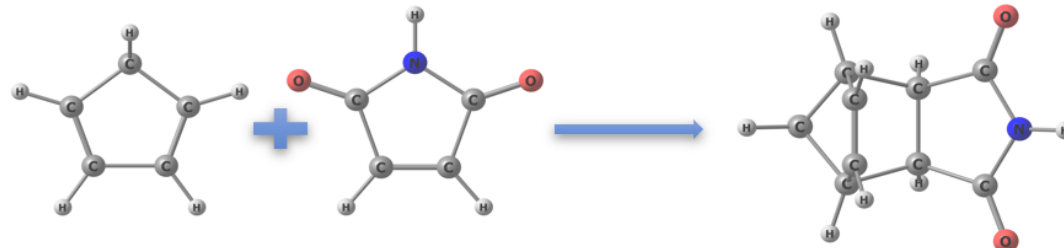
Sure!



Hello! I am *Aitomia*







J. Hu, H. Nawaz, Y.-F. Hou, Y. Rui, L. Chi, Y. Chen\*, A. Ullah\*, P. O. Dral\*. *JCTC* **2026**, 22, 6557–6571



Calculate the reaction energy and thermodynamic properties for the Diels-Alder reaction between cyclopentadiene and maleimide using AIQM2. I will provide the adduct. Please generate the other required structures for me.



The user only provides the adduct structure. For this task, the structures of maleimide and cyclopentadiene are also required. I will call the initial structure generation agent to generate the missing structures needed to solve the user's problem.



For **\*\*adduct\*\***, I will proceed with following calculations:

.....

For **\*\*maleimide\*\***, I'll follow this plan:

.....



We are calculating the reaction energy ( $\Delta E$ ), enthalpy change ( $\Delta H$ ), and Gibbs free energy change ( $\Delta G$ ) for Diels-Alder reaction.



1. Reaction energy ( $\Delta E$ ) = -33.73 kcal/mol
2. Enthalpy Change ( $\Delta H$ ) = -30.41 kcal/mol
3. Gibbs Free Energy Change ( $\Delta G$ ) = -15.

EXPLORER

bugs

demo

md

md-ml

mltpa20250728\_140831

test

tpa

tpa2

Welcome to Aitomistic Lab

Please create a new project or open an existing file

You can try to

Create Demo Folder

Or

chat with chatbot

Save File: 

Command

 + 

S

Copy File: 

Command

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C

Paste File: 

Command

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Submit Current File: 

Command

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R

Select Multiple File: 

Shift

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Create New File

TERMINAL

OUTPUT

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AITOMIA



Hi, I am Aitomia, nice to meet you!  
I'm an intelligent assistant with expertise in AI-driven atomistic simulations and quantum chemistry.  
You can select a specific molecule file by open it in the explorer,  
or select a folder by the button  in the input box.

You can ask me:

Can you calculate frequencies and thermochemistry for me?

Can you optimize the geometry?

Can you perform single point calculation?

Can you perform transition state search?

Can you calculate the infrared spectra?

Can you calculate the ultraviolet spectra?

Can you calculate the raman spectra?

Press Enter to send message

A

Deep Thinking ☐





J. Hu, H. Nawaz, Y.-F. Hou, Y. Rui, L. Chi, Y. Chen\*, A. Ullah\*, P. O. Dral\*. *JCTC* **2026**, 22, 6557–6571

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Protomia

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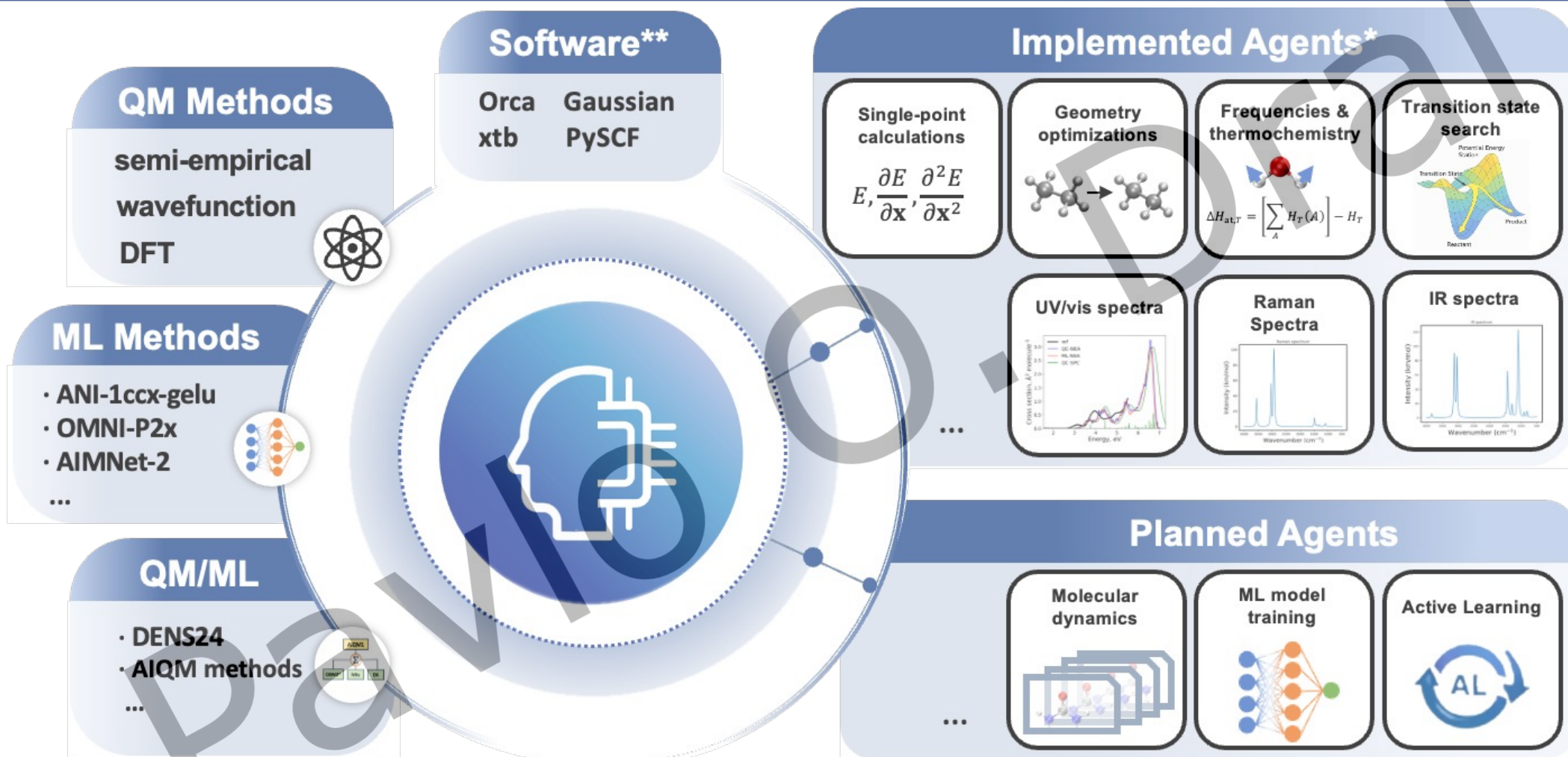
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+ mini\_paper.pdf

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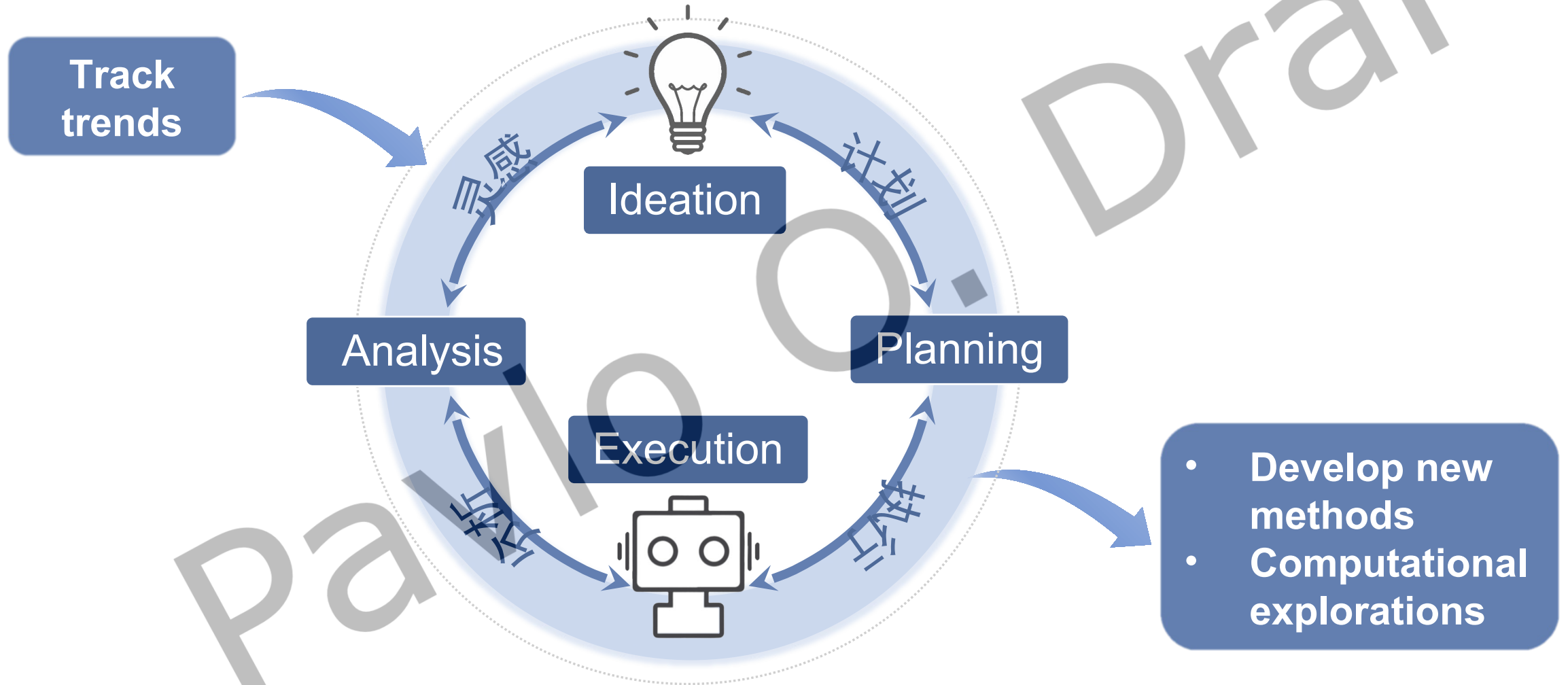
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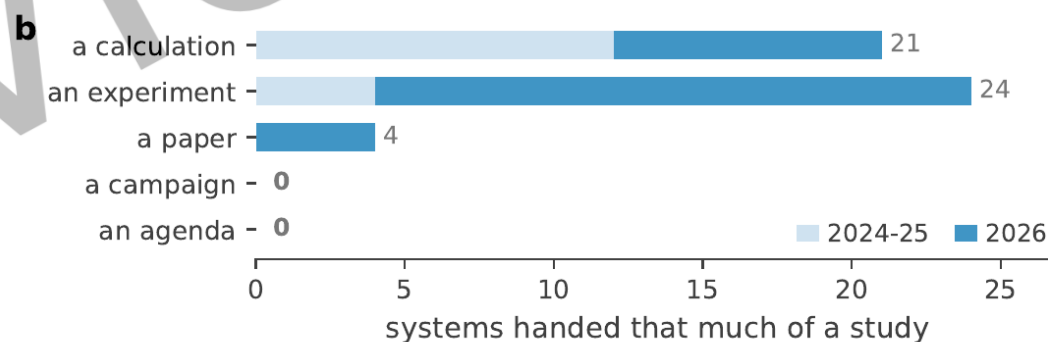
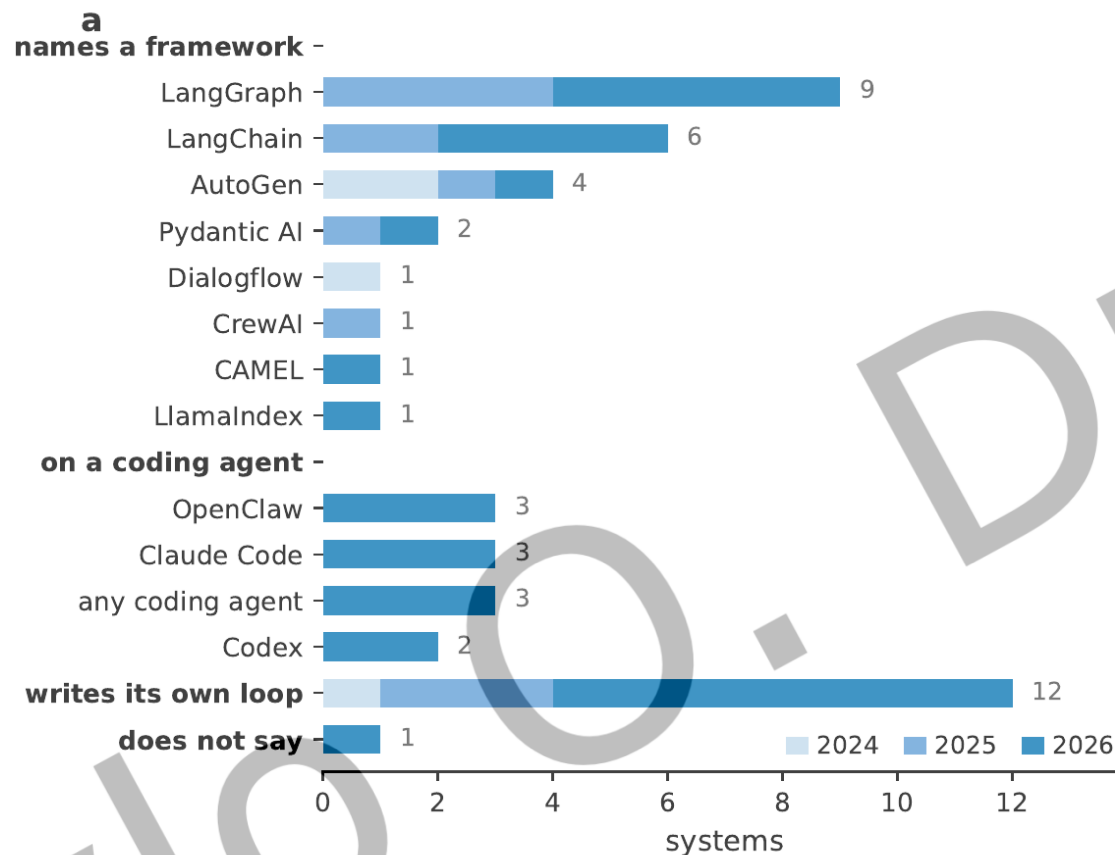




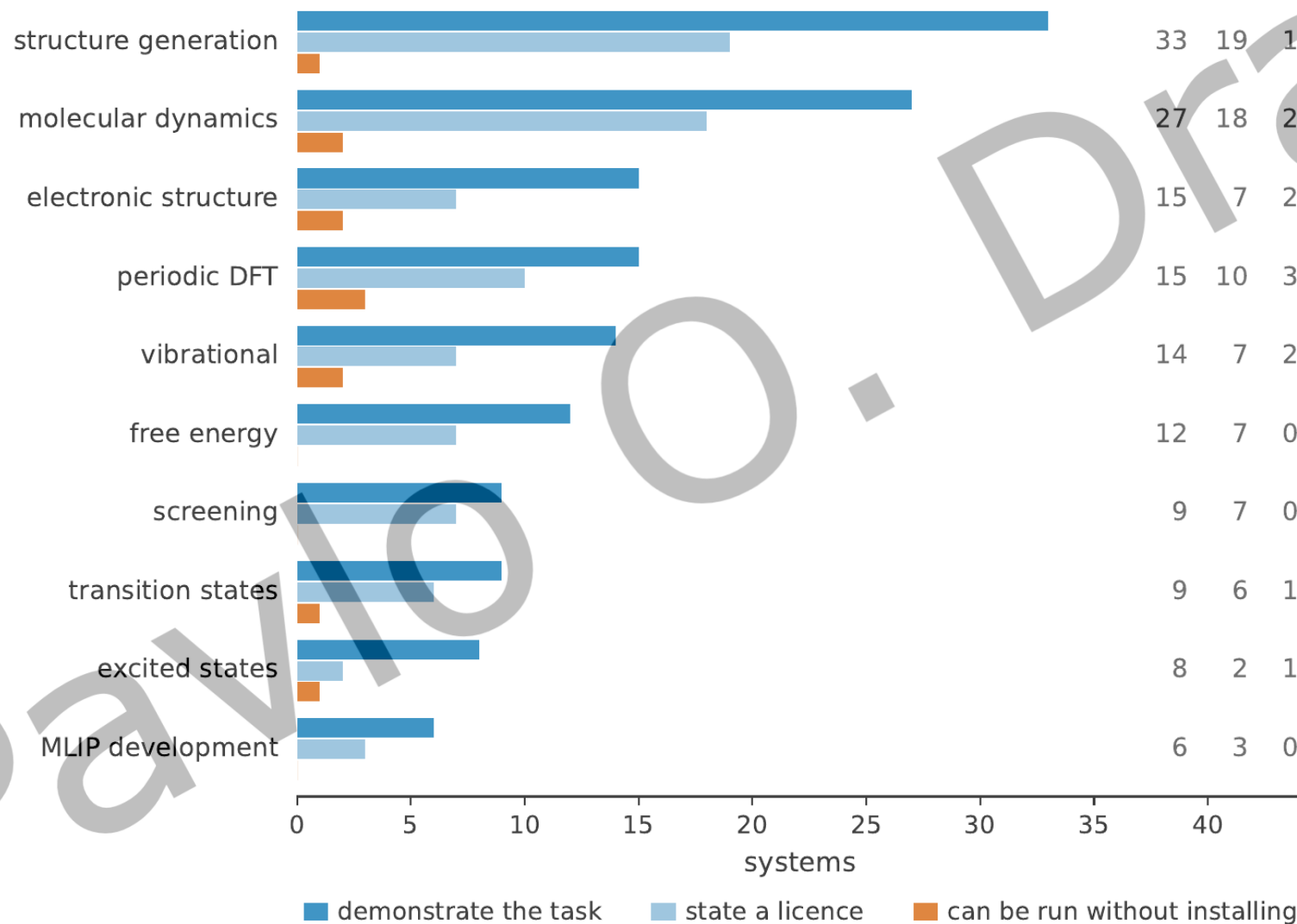
\*Implemented agents can be used in more complex workflows, such as reaction enthalpy calculations.

\*\*Only software with permissive licenses is available online for registered users, while Aitomia supports more programs.

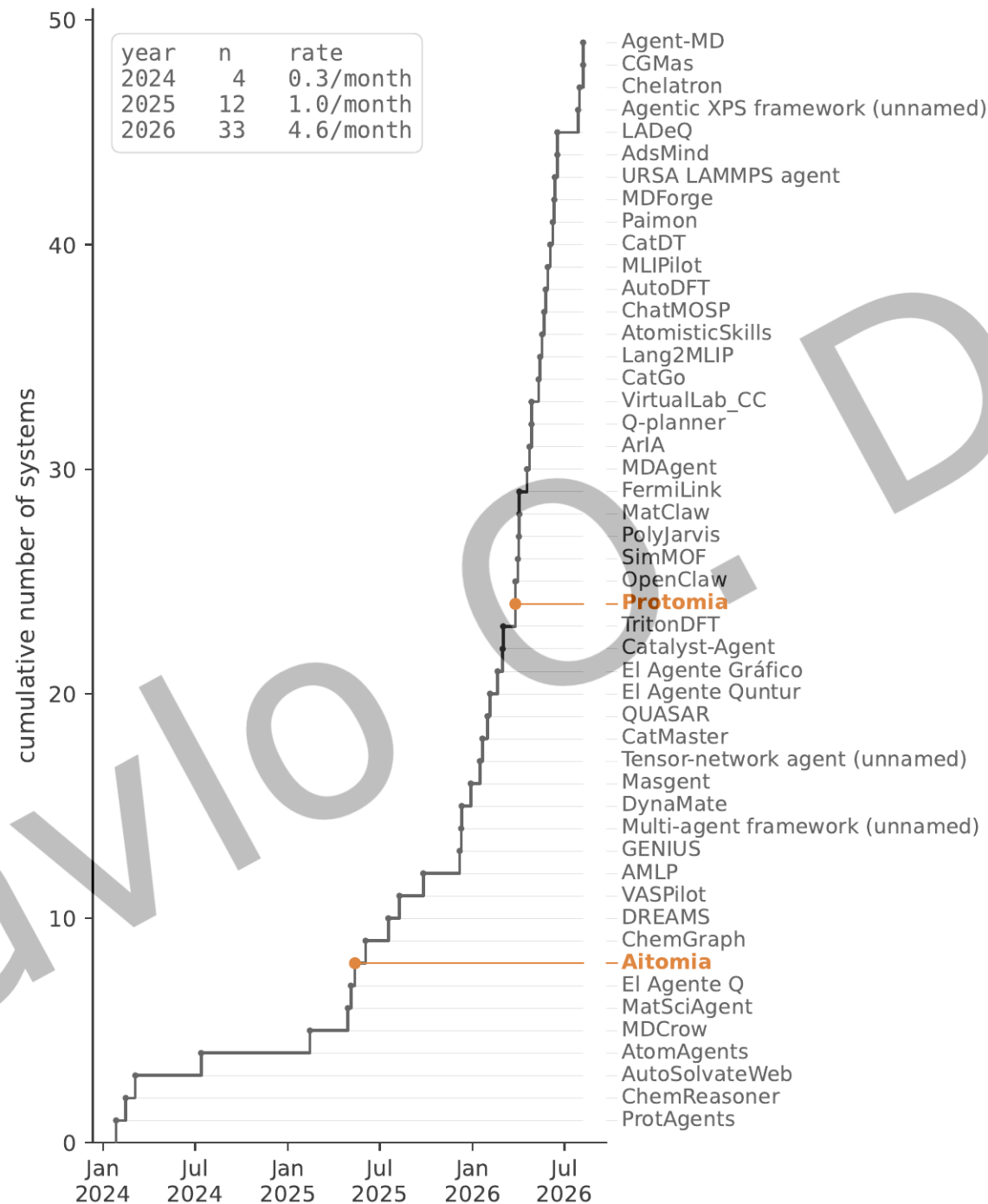


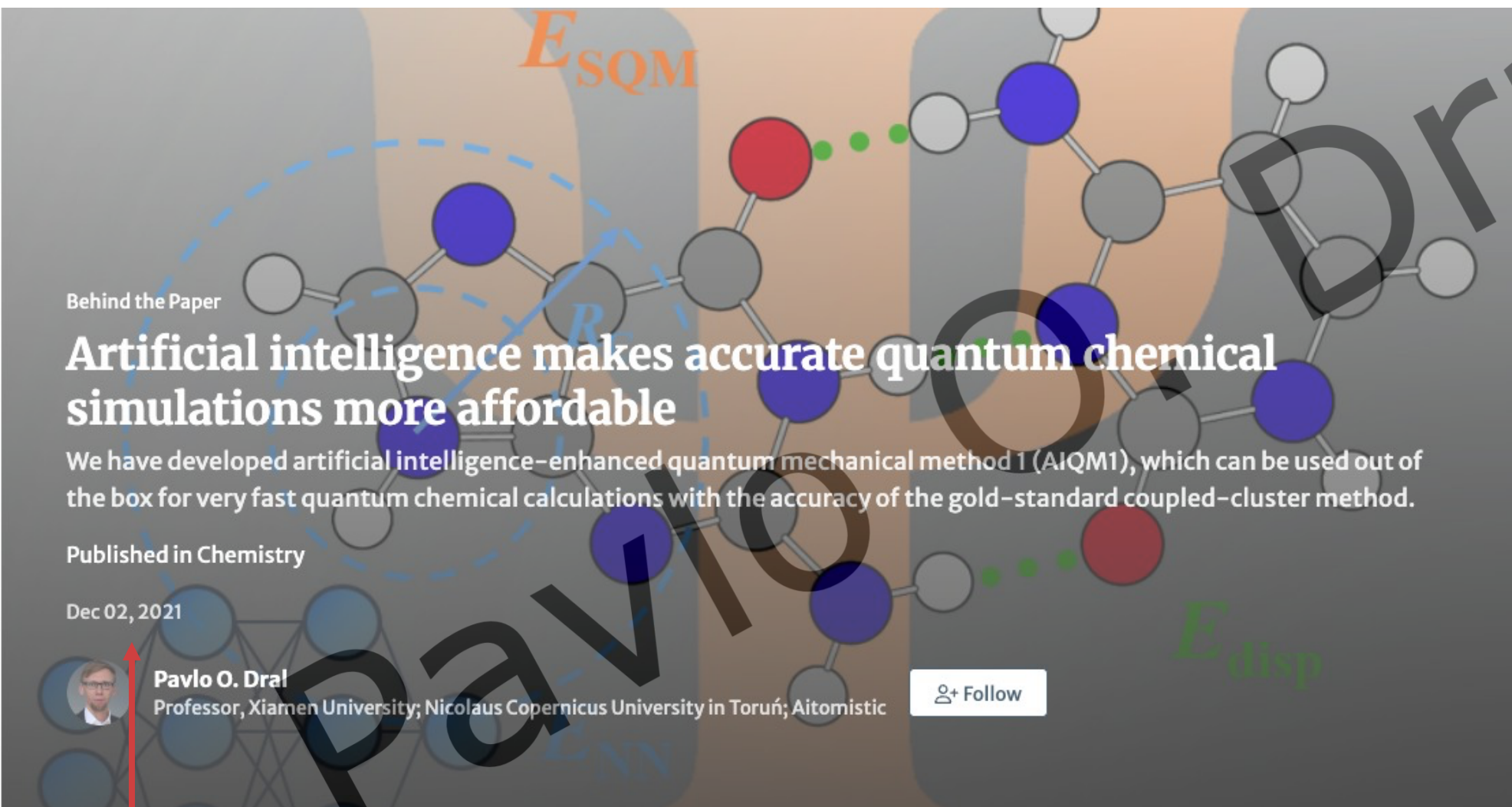






P. O. Dral, H. Nawaz, A. Ullah. Science Done on a Machine by a Machine: AI Agents in Computational Chemistry. **2026**. arXiv:2608.18508





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Published in Chemistry

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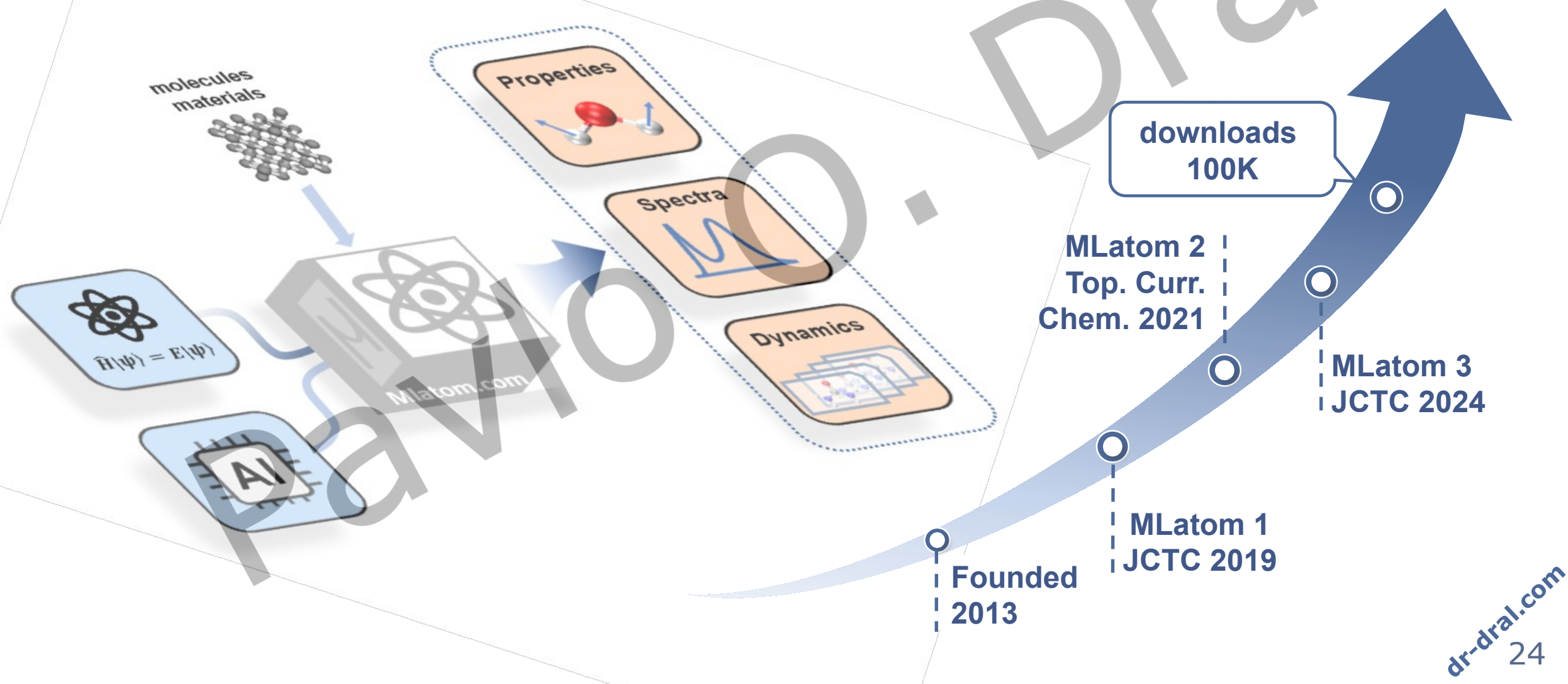
ChatGPT:  
Nov 30, 2022

<https://communities.springernature.com/posts/artificial-intelligence-makes-accurate-quantum-chemical-simulations-more-affordable>

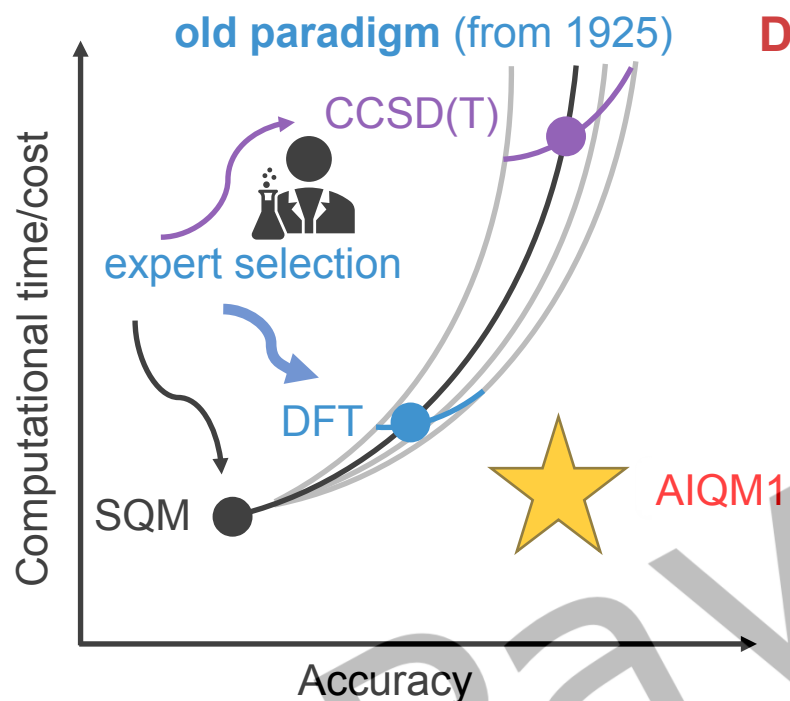


We develop MAtom – an open-source software ecosystem integrating AI and quantum chemistry

<https://github.com/dralgroup/mlatom>



**We developed AI models working out of the box, resolving their stability problem**



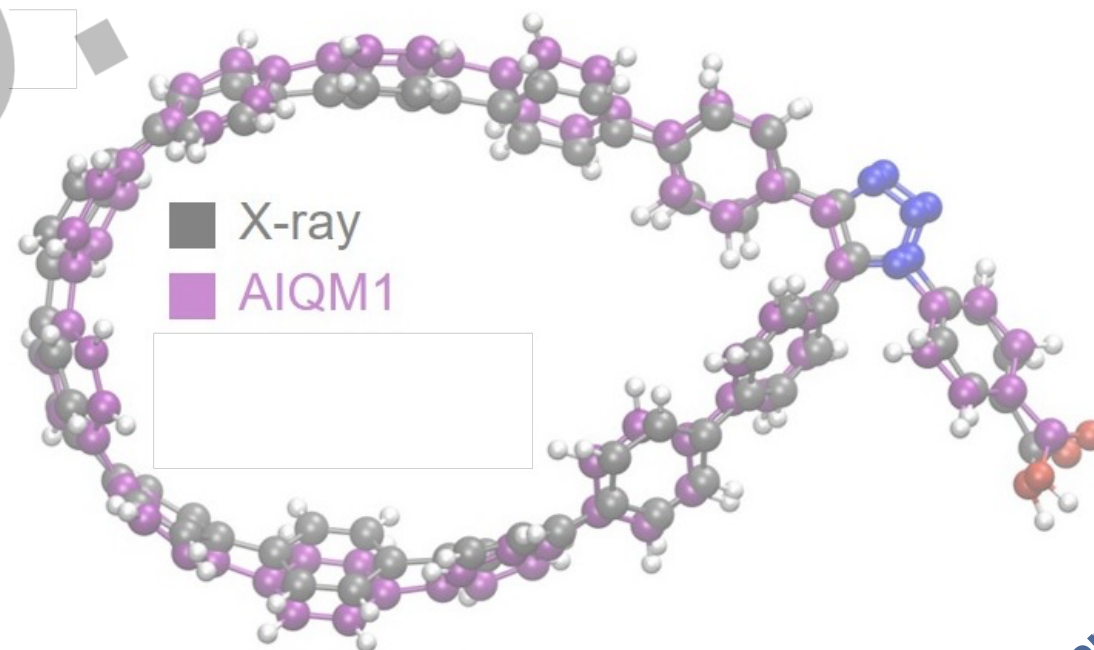
**Dral\*, et al. Nat. Commun. 2021**



**AIQM1 vs DFT**

**2 vs 1200 CPU-min**

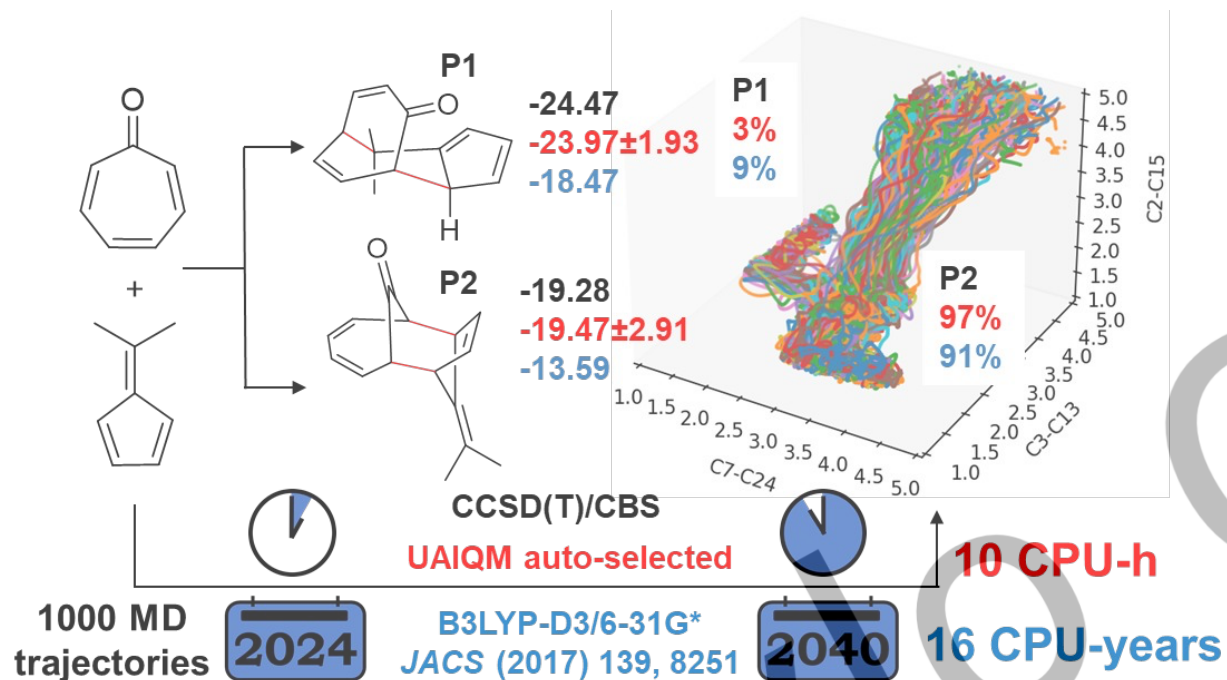
**0.625 vs 0.663 Å (RMSD)**



**AIQM1 vs typical DFT (B3LYP, PBE, ...)**

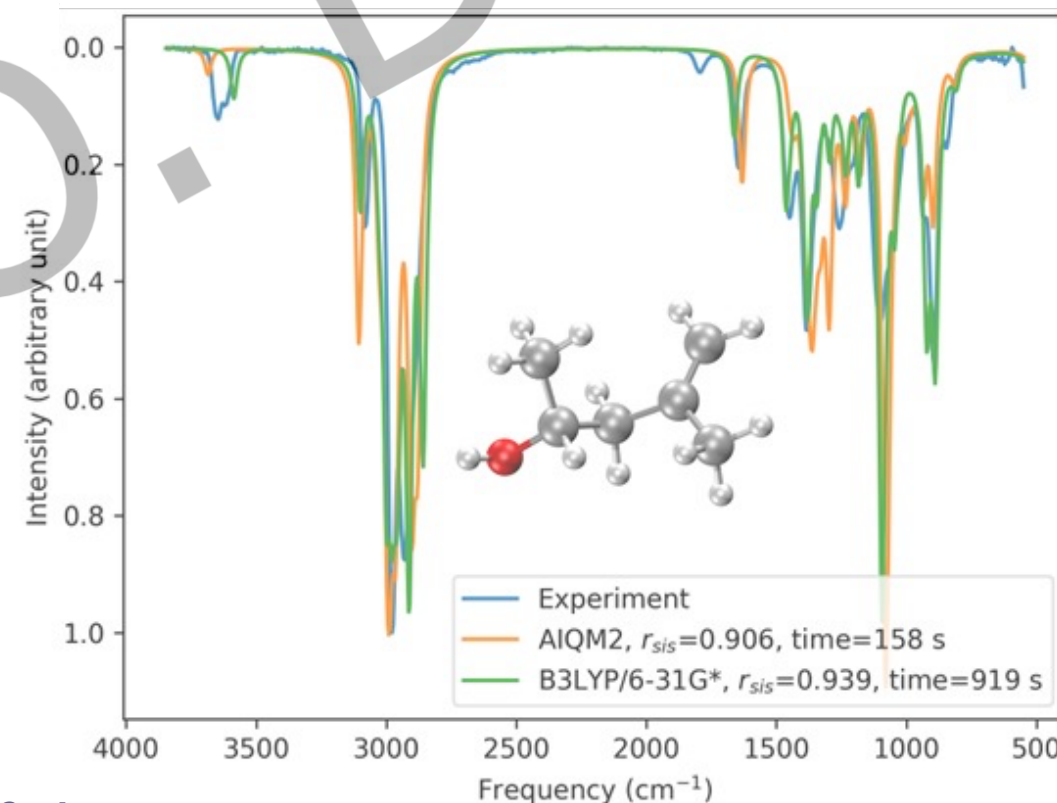
- **Orders of magnitude faster**
- **Accuracy is usually higher** for closed-shell, neutral, CHNO molecules

**Dral, et al. J. Chem. Eur. J. 2023, 29, e202300668**



Fast, accurate, and trustworthy AI models for  
快速、准确且可靠的人工智能模型用于

- reaction mechanisms, thermochemistry, IR and Raman spectra, MD, many chemical elements...

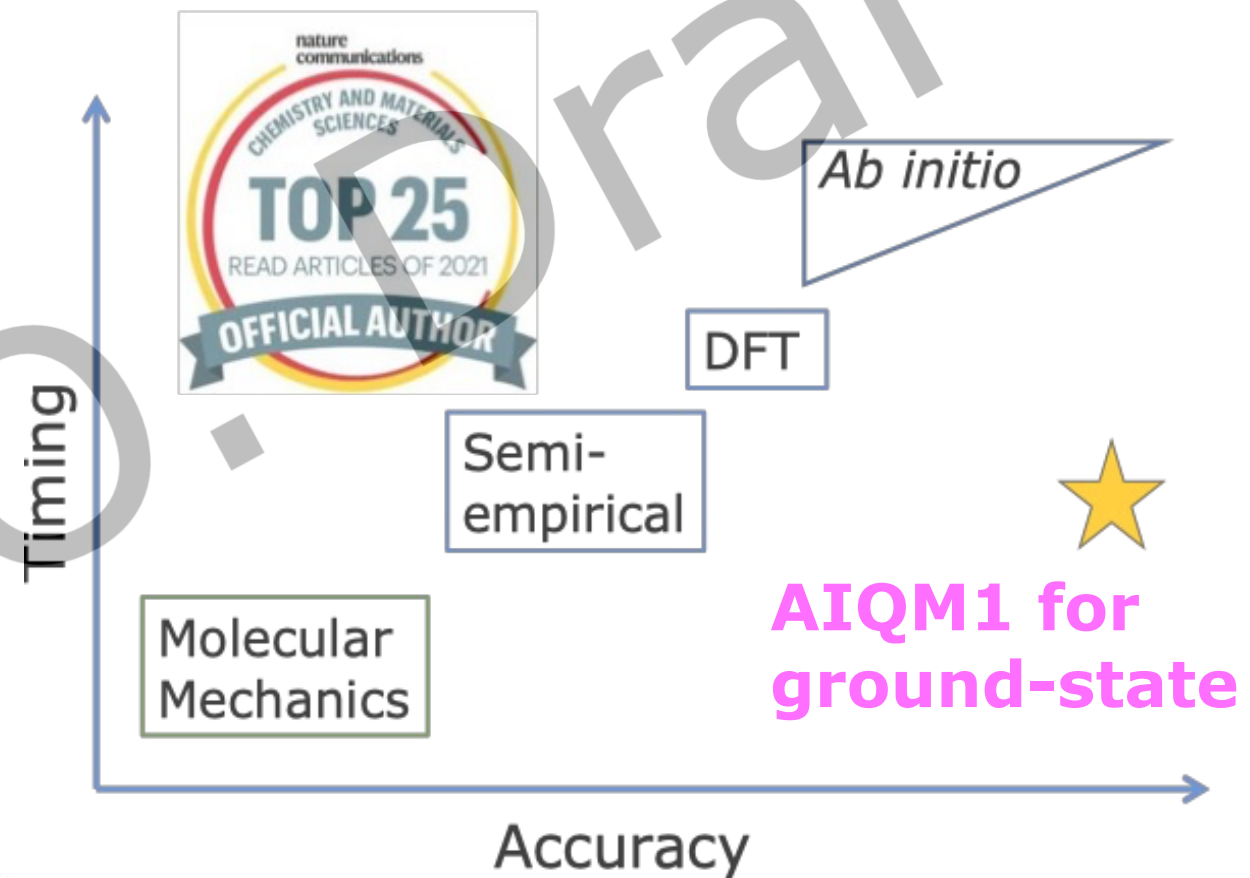
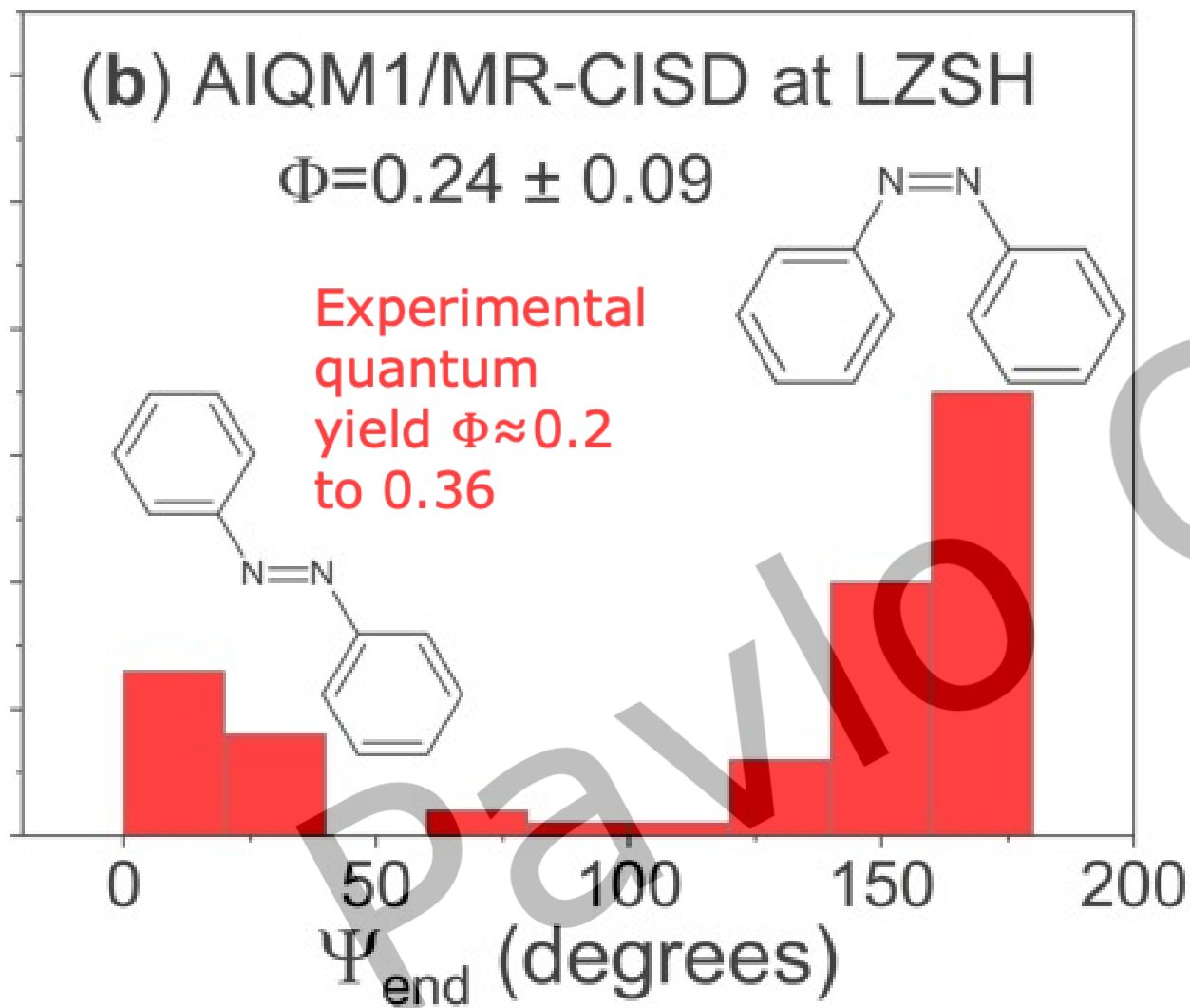


Selected publications on the models:

- Drál\*, et al. *Nat. Commun.* 2021, 12, 7022
- Drál\*, et al. *J. Phys. Chem. Lett.* 2022, 13, 3479; 2025, 16, 4
- Drál\*, et al. *Chem. Sci.* 2025, 16, 15901







AIQM1: P. Zheng, R. Zubatyuk, W. Wu, O. Isayev, P. O. Dral, *Nat. Commun.* **2021**, 12, 7022

# Handing off everything to Protomia

## Protomia Solution 2.8.3

The molecule in your folder is **methane (CH<sub>4</sub>)** (from `sp.xyz`), and I've submitted the AIQM2 orbital calculation to the cluster — it's running now (SLURM job 183853).

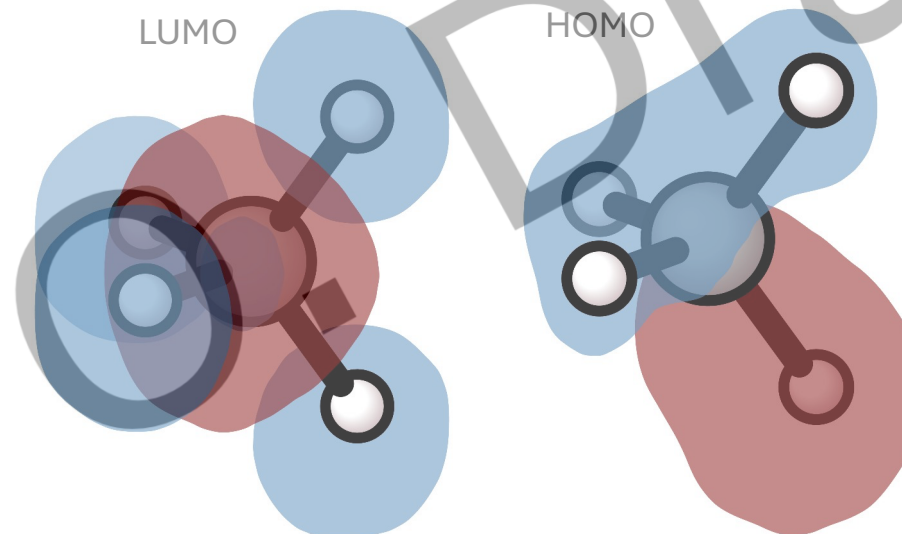
The job runs the single-point AIQM2 energy, then automatically renders the **HOMO** and **LUMO** molecular-orbital surface pictures, so you don't need to do anything else. It should finish quickly — I'll report the orbital energies (HOMO/LUMO in eV), the single-point energy, and the rendered orbital images to you as soon as it's done. Feel free to keep chatting in the meantime!

The methane (CH<sub>4</sub>) AIQM2 orbital calculation has finished — here are the results:

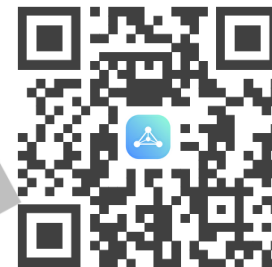
**Single-point energy (AIQM2): -40.464860 Hartree** (dipole moment: 0.0 D, as expected for this T<sub>4h</sub>-symmetric geometry)

### Frontier molecular orbitals:

- **HOMO: -12.6931 eV**
- **LUMO: 4.4160 eV**
- **HOMO-LUMO gap: 17.1092 eV**



Visualization of HOMO and LUMO orbitals of methane calculated with AIQM2 by Protomia



Aitomistic Lab@XMU  
[atom.xmu.edu.cn](http://atom.xmu.edu.cn)



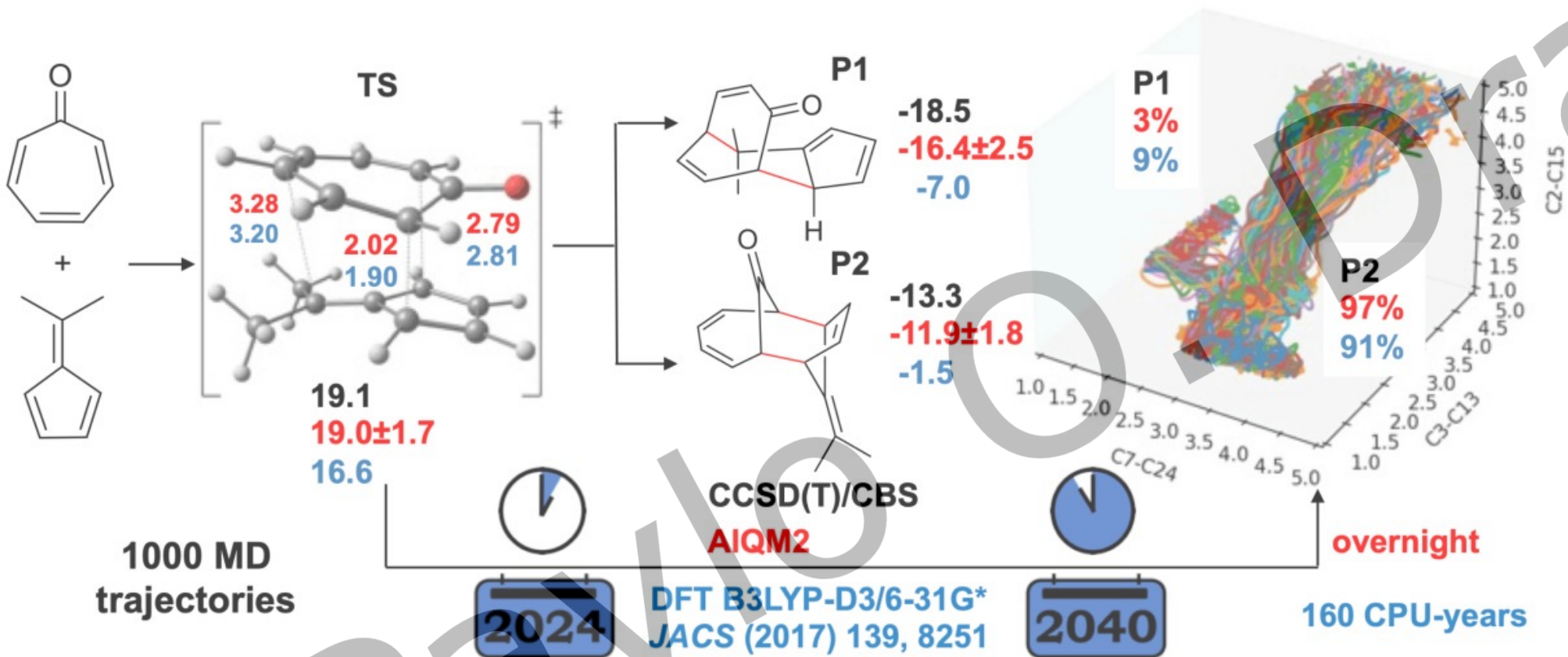
Docs of MLatom3  
[mlatom.com/docs](http://mlatom.com/docs)

rsc.li/chemical-science

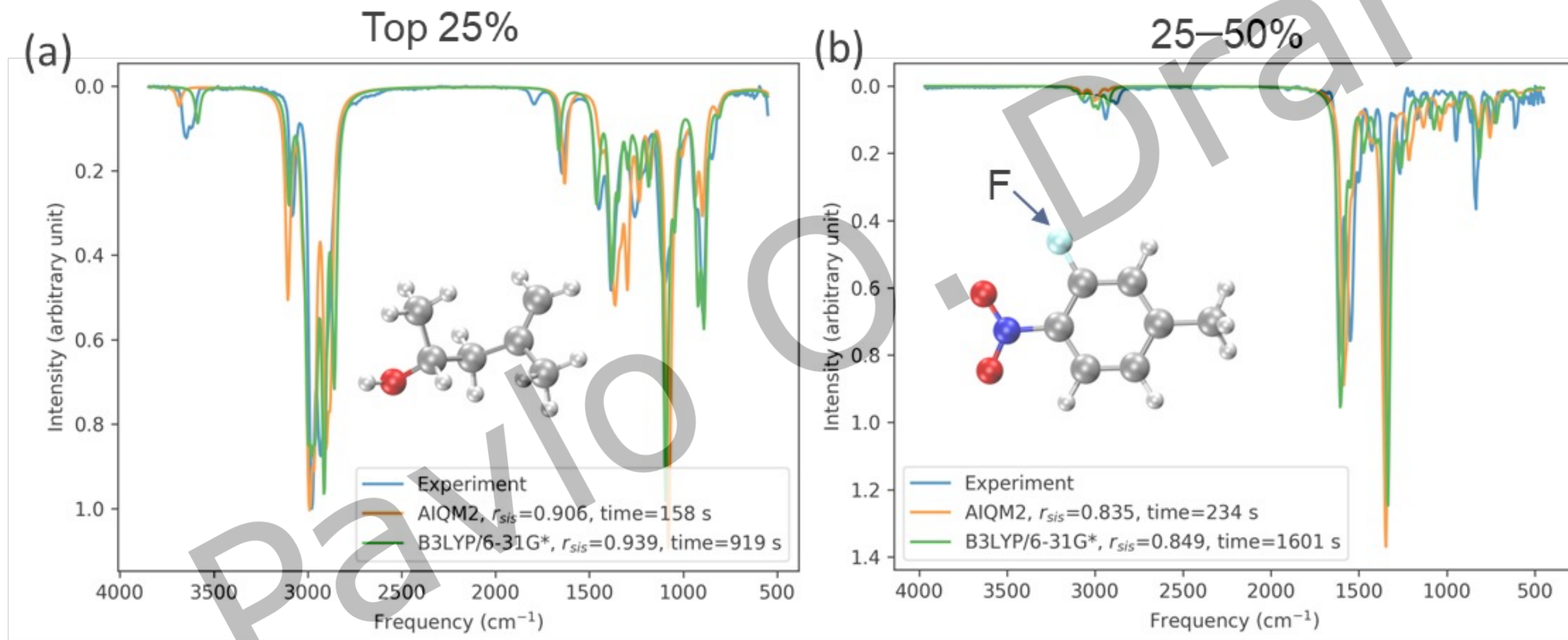


ISSN 2041-6539





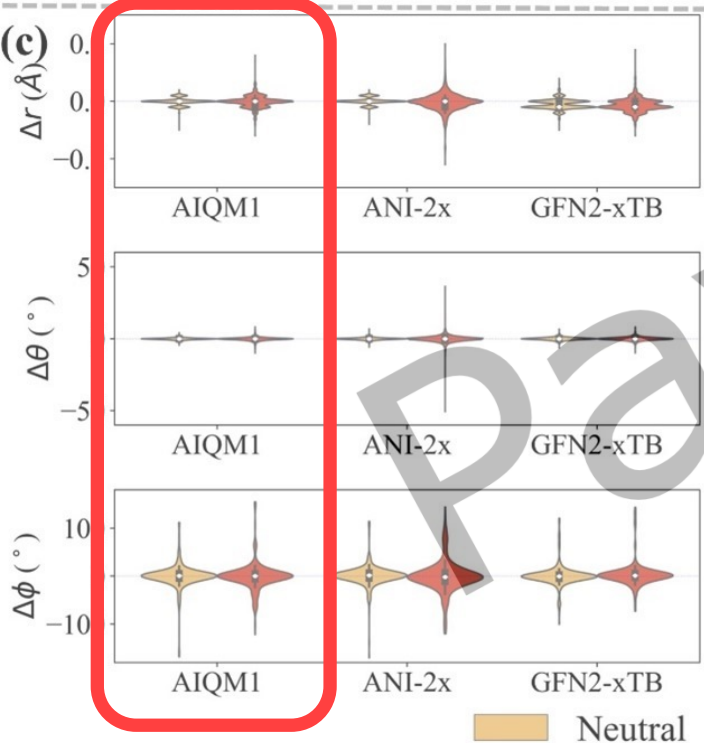
- Fast
- Accurate
- Trustworthy



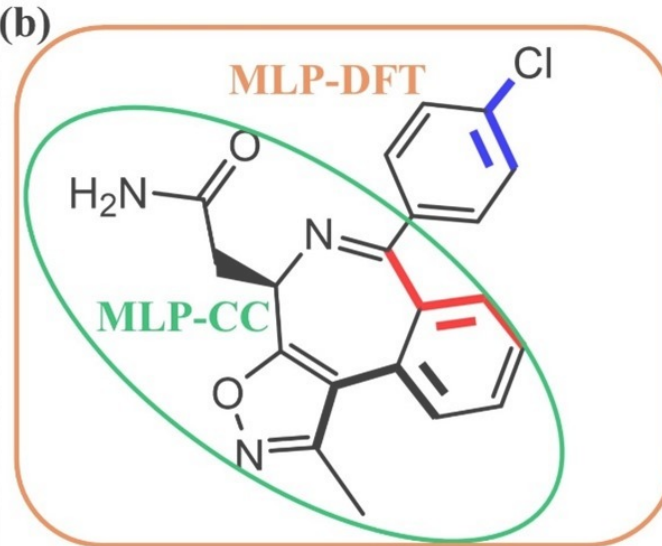
(a)

|      | Methods                                    |
|------|--------------------------------------------|
| M1   | ONIOM2(DFT:MM)                             |
| M2   | ONIOM2-EE(DFT:MM)                          |
| M3   | ONIOM2(ANI-2x:MM)                          |
| M4   | ONIOM2(ANI-1ccx:MM)                        |
| M4a  | ONIOM3(ANI-1ccx:ANI-2x:MM) <sup>b</sup>    |
| M5   | ONIOM2(AIQM1:MM) <sup>c</sup>              |
| M5a  | ONIOM3(AIQM1:ANI-2x:MM) <sup>b</sup>       |
| M6   | ONIOM2(SE:MM)                              |
| M7   | ONIOM3(DFT:SE:MM)                          |
| M8   | ONIOM3(ANI-2x:SE:MM)                       |
| M9   | ONIOM3(ANI-1ccx:SE:MM)                     |
| M9a  | ONIOM4(ANI-1ccx:ANI-2x:SE:MM) <sup>b</sup> |
| M10  | ONIOM3(AIQM1:SE:MM) <sup>c</sup>           |
| M10a | ONIOM4(AIQM1:ANI-2x:SE:MM) <sup>b</sup>    |
| M6R  | ONIOM2(SE:MM) <sup>d</sup>                 |

(c)

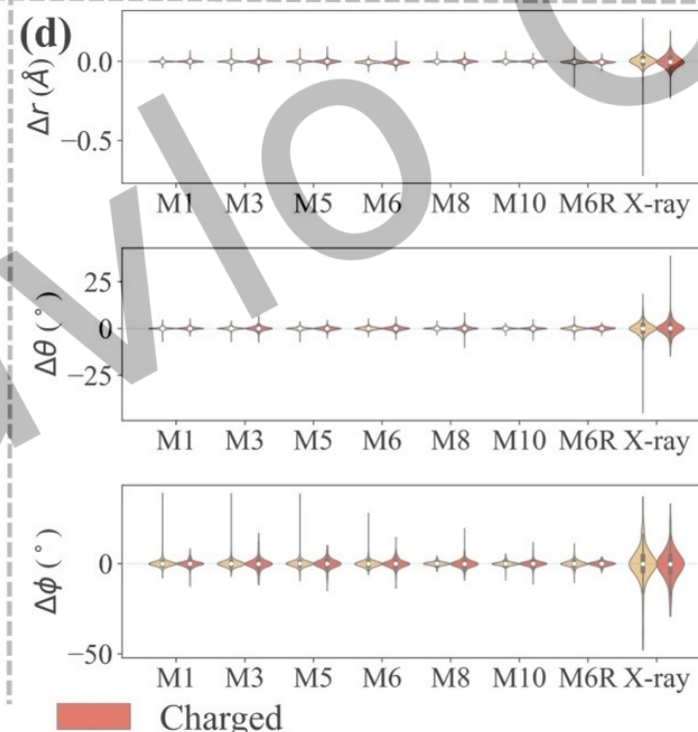


(b)



ONIOM2(MLP-CC:MLP-DFT)  
ONIOM2(MLP:xTB)

(d)



Article | [Open access](#) | Published: 16 May 2024

# Accelerating reliable multiscale quantum refinement of protein–drug systems enabled by machine learning

[Zeyin Yan](#), [Dacong Wei](#), [Xin Li](#) & [Lung Wa Chung](#)

[Nature Communications](#) **15**, Article number: 4181 (2024) | [Cite this article](#)



# Integrating Machine Learning Interatomic Potentials with MMPBSA for Accurate Protein–Ligand Binding Free Energy Calculations

Xue-Xin Wei, Yuxinxin Chen, Yuedong Yang, Mingyuan Xu\*, Pavlo O. Dral\*, and Hongming Chen\*

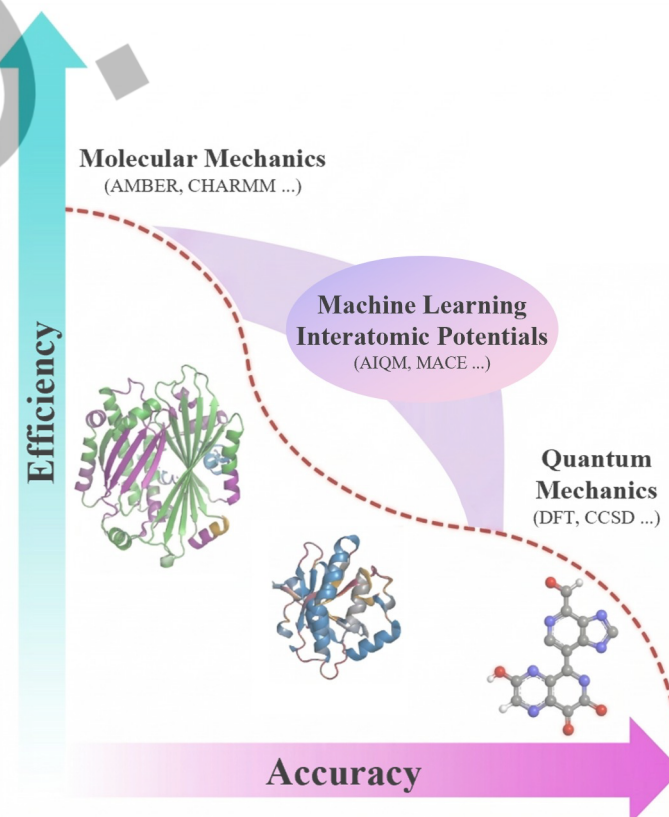
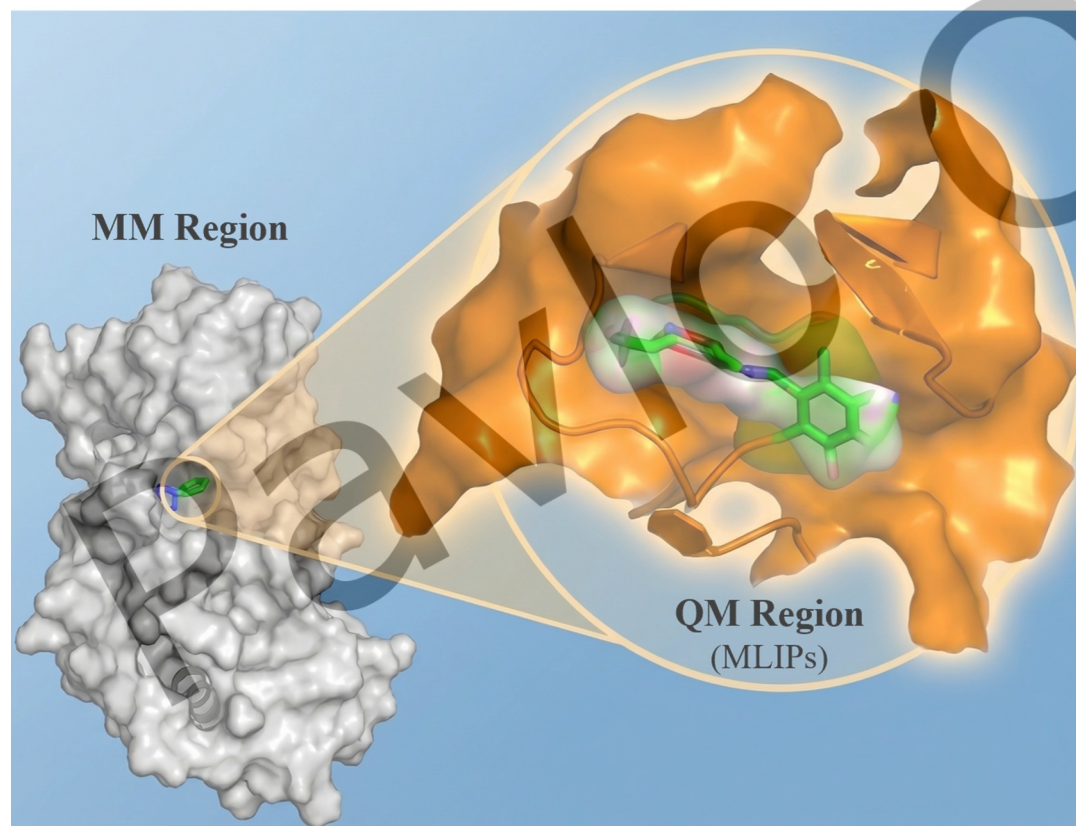
Open PDF

Supporting Information (1)

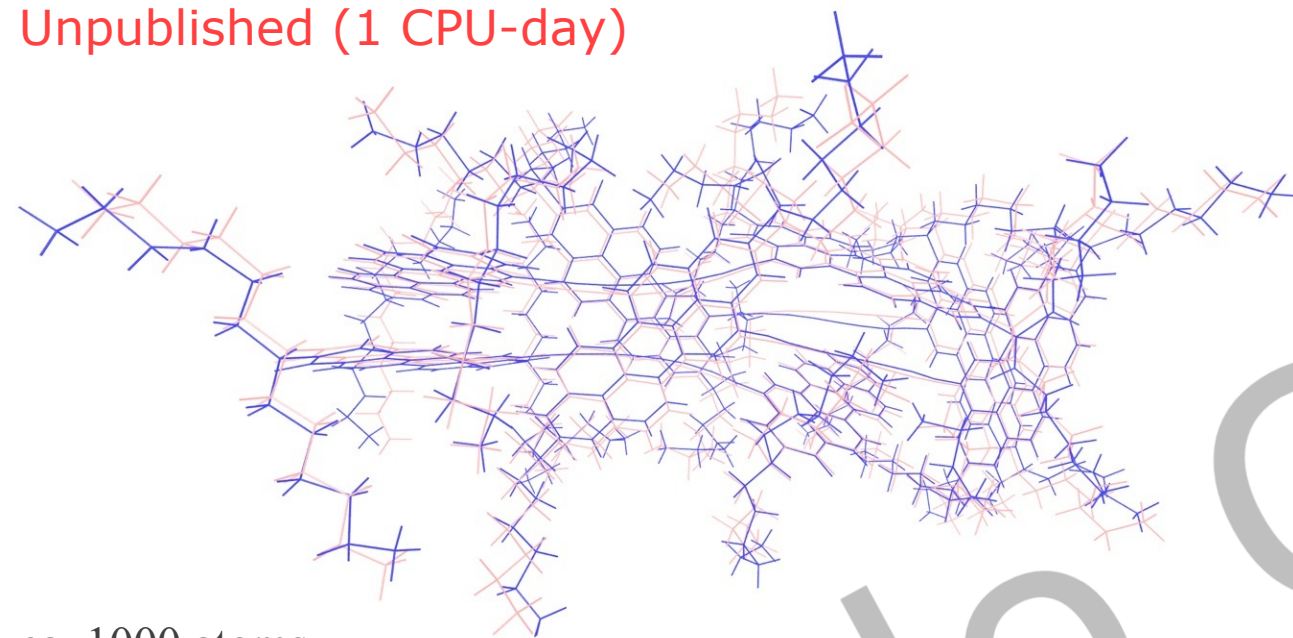
XMU

ONIOM(MLIP:MM)

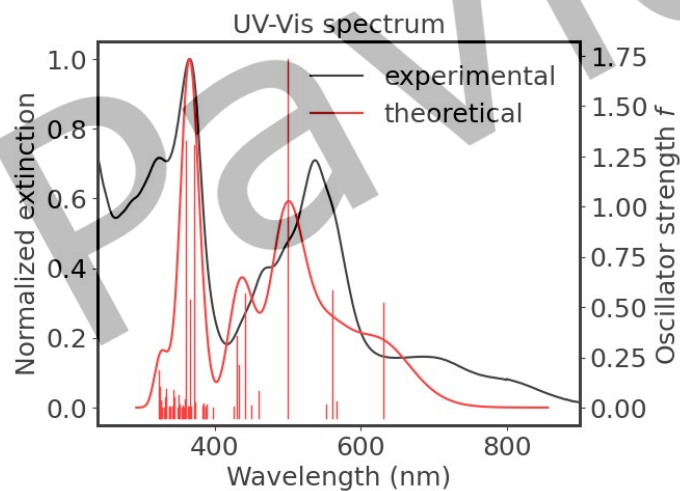
Accuracy Meets Efficiency



Unpublished (1 CPU-day)

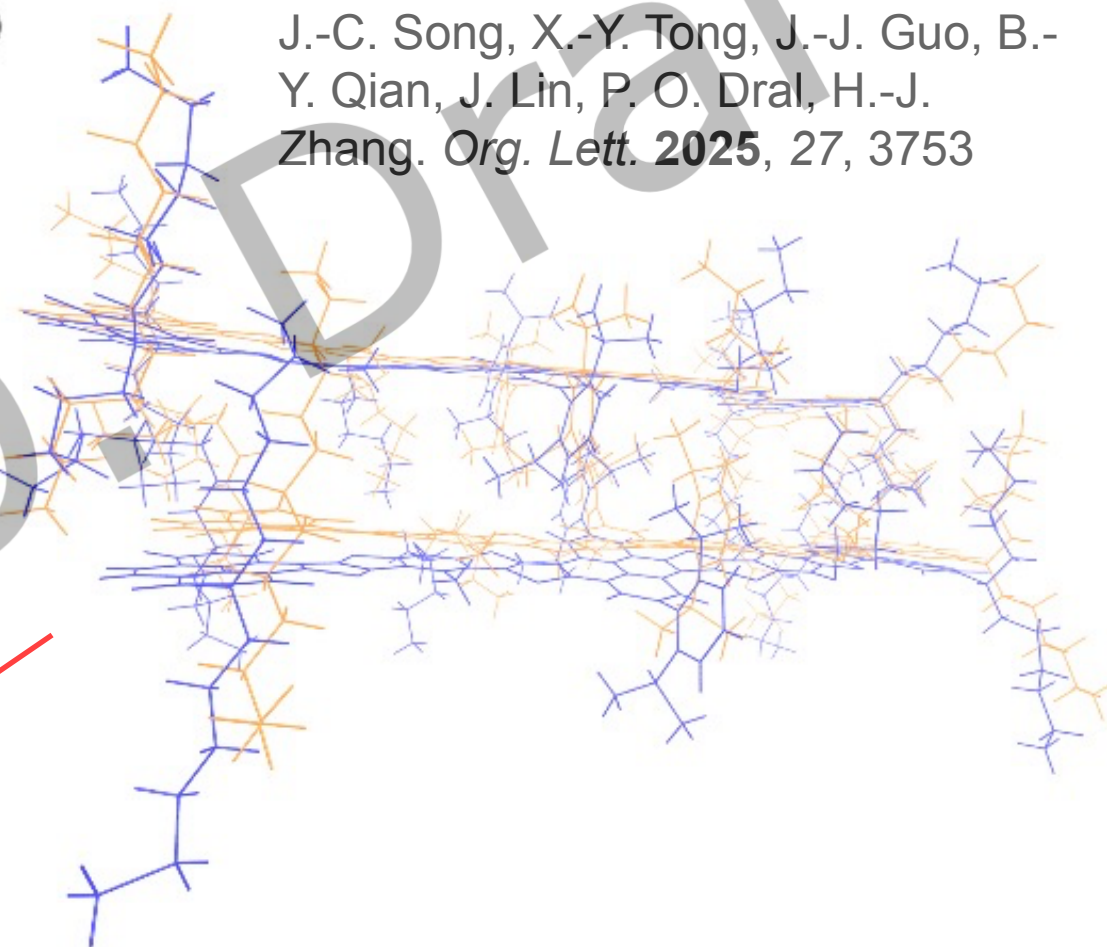


ca. 1000 atoms,  
agreement with crystallographic structure: 0.63 Å

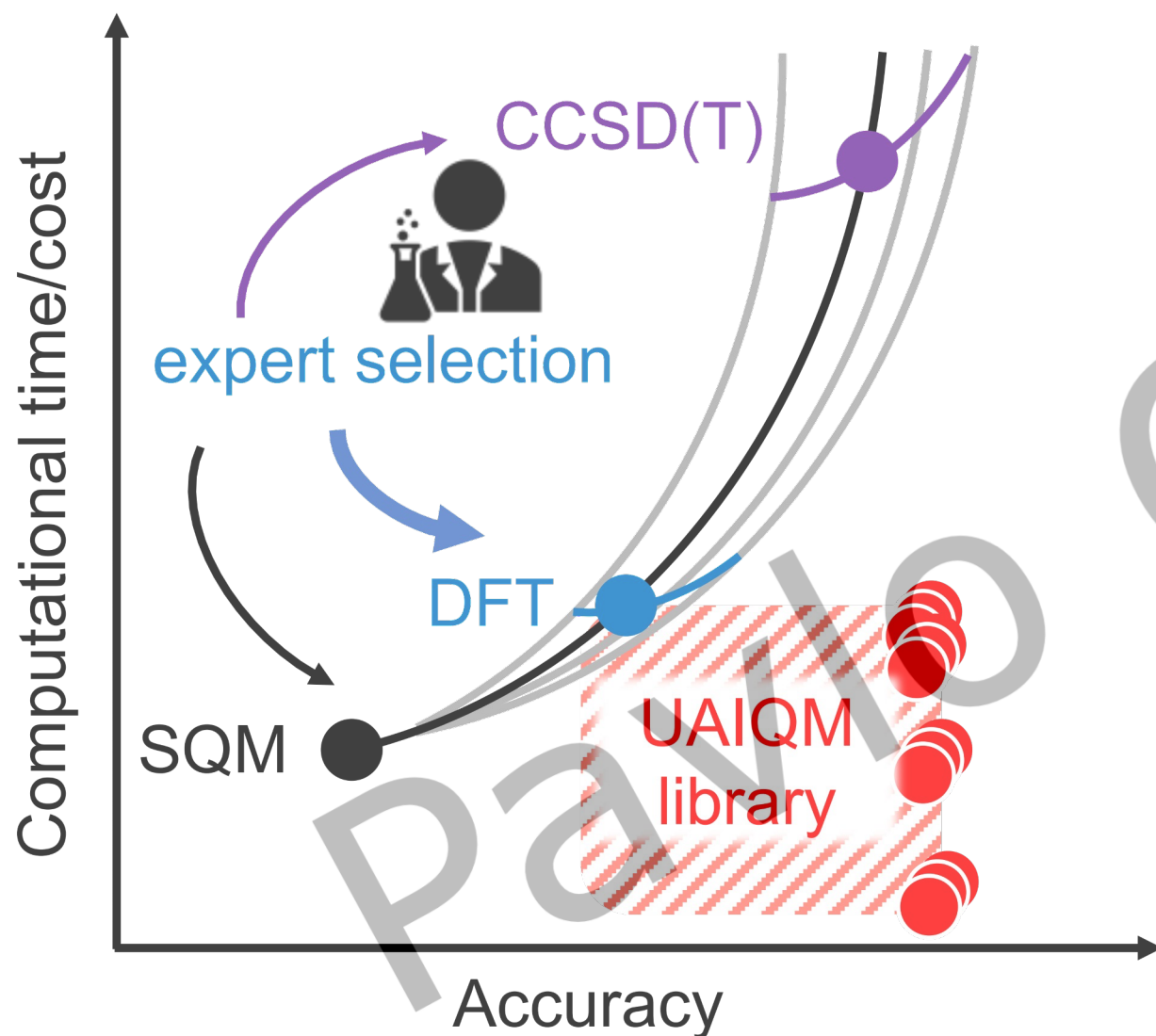


(b)

J.-C. Song, X.-Y. Tong, J.-J. Guo, B.-Y. Qian, J. Lin, P. O. Dral, H.-J. Zhang. *Org. Lett.* **2025**, 27, 3753



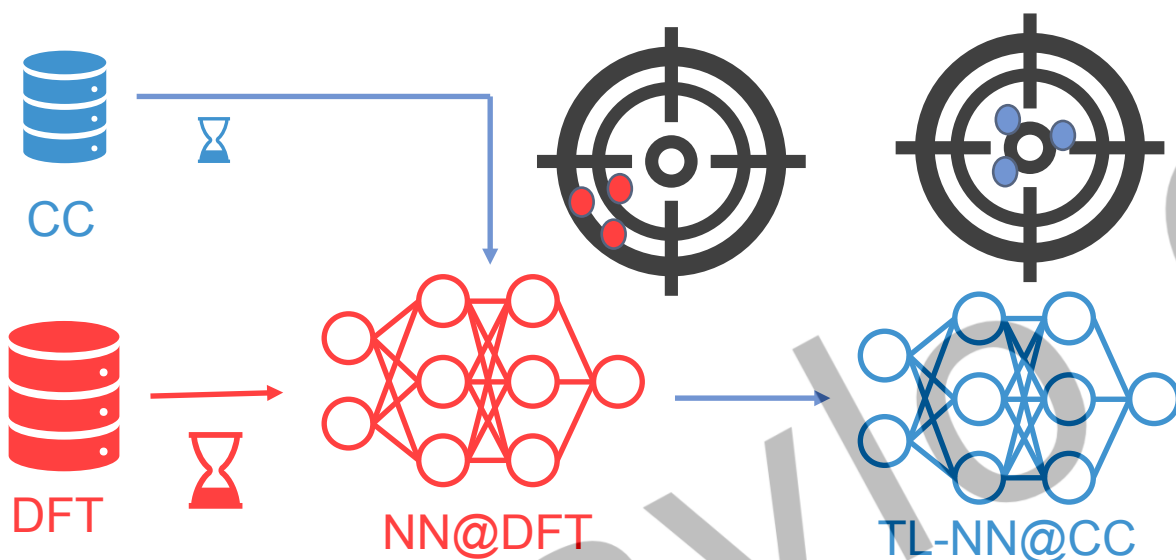
714 atoms,  
agreement with crystallographic structure: 2.948 Å





| name          | keywords in MLatom      | version  | target level | baseline                 | dispersion          | recommended elements             |
|---------------|-------------------------|----------|--------------|--------------------------|---------------------|----------------------------------|
| -             | uaiqm_nobaseline@dft    | 20240606 | DFT          | -                        | D4 ( $\omega$ B97X) | H C N O                          |
| ANI-1ccx      | uaiqm_nobaseline@cc     | 20190701 | CC           | -                        | -                   | H C N O                          |
| -             | uaiqm_nobaseline@cc     | 20240606 | CC           | -                        | D4 ( $\omega$ B97X) | H C N O                          |
| AIQM1@DFT     | uaiqm_odm2star@dft      | 20211202 | DFT          | ODM2*                    | D4 ( $\omega$ B97X) | H C N O                          |
| AIQM1         | uaiqm_odm2star@cc       | 20211202 | CC           | ODM2*                    | D4 ( $\omega$ B97X) | H C N O                          |
| -             | uaiqm_odm2star@cc       | 20240308 | CC           | ODM2*                    | D4 ( $\omega$ B97X) | H C N O                          |
| AIQM2@DFT     | uaiqm_gfn2xtbstar@dft   | 20240106 | DFT          | GFN2-xTB*                | D4 ( $\omega$ B97X) | H C N O                          |
| AIQM3@DFT     | uaiqm_gfn2xtbstar@dft   | 20240619 | DFT          | GFN2-xTB*                | D3bj (b97-3c)       | H C N O F S Cl                   |
| AIQM4@DFT (?) | uaiqm_gfn2xtbstar@dft   | 20240730 | DFT          | GFN2-xTB*                | D3bj (b97-3c)       | H B C N O F Si P S Cl As Se Br I |
| AIQM2         | uaiqm_gfn2xtbstar@cc    | 20240106 | CC           | GFN2-xTB*                | D4 ( $\omega$ B97X) | H C N O                          |
| AIQM3         | uaiqm_gfn2xtbstar@cc    | 20250115 | CC           | GFN2-xTB*                | D3bj (b97-3c)       | H C N O F S Cl                   |
| -             | uaiqm_wb97x631gp@cc     | 20230924 | CC           | $\omega$ B97X/6-31G*     | D4 ( $\omega$ B97X) | H C N O                          |
| -             | uaiqm_wb97xdef2tzvpp@cc | 20230923 | CC           | $\omega$ B97X/def2-TZVPP | D4 ( $\omega$ B97X) | H C N O                          |
| -             | uaiqm_wb97xdef2tzvpp@cc | 20240227 | CC           | $\omega$ B97X/def2-TZVPP | D4 ( $\omega$ B97X) | H C N O                          |
| -             | uaiqm_wb97xdef2tzvpp@cc | 20240228 | CC           | $\omega$ B97X/def2-TZVPP | D4 ( $\omega$ B97X) | H C N O                          |





- × two-step training
- × separate models
- × cannot be trained on combined databases

Fine-tuning of AIQM, ANI, OMNI-P models: [https://aitomistic.com/mlatom/tutorial\\_finetuning.html](https://aitomistic.com/mlatom/tutorial_finetuning.html)

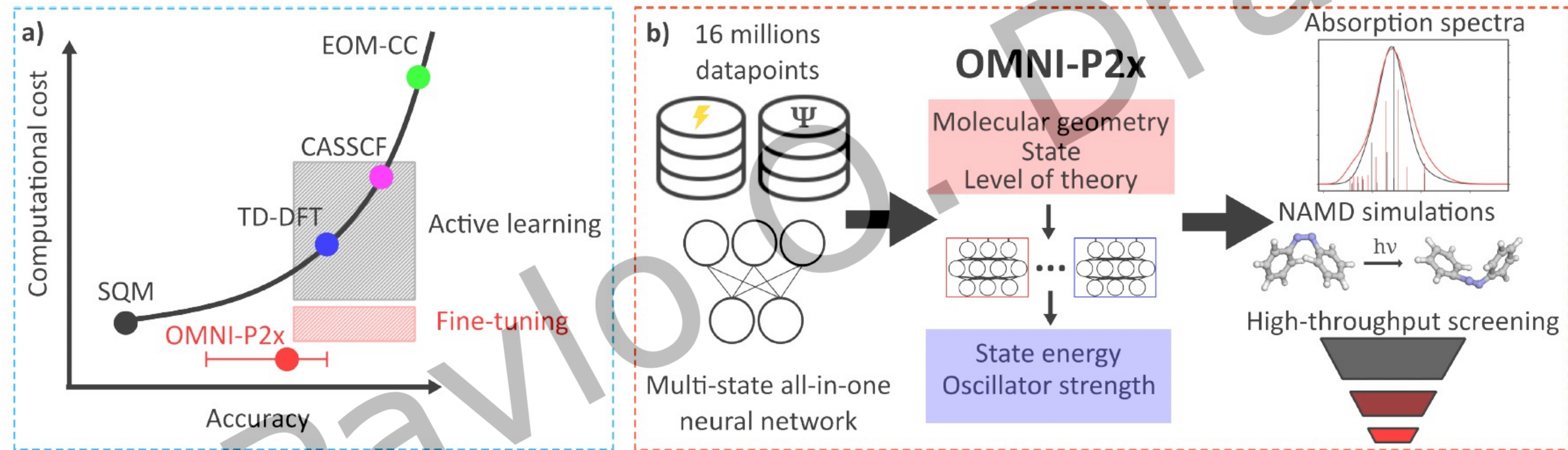
Seyedeh Fatemeh Alavi, Yuxinxin Chen, Yi-Fan Hou, Fuchun Ge, Peikun Zheng, Pavlo O. Dral. ANI-1ccx-gelu Universal Interatomic Potential and Its Fine-Tuning: Toward Accurate and Efficient Anharmonic Vibrational Frequencies. *J. Phys. Chem. Lett.* **2025**, 16, 483





## First-ever excited-state NNP

- Much more difficult than ground state

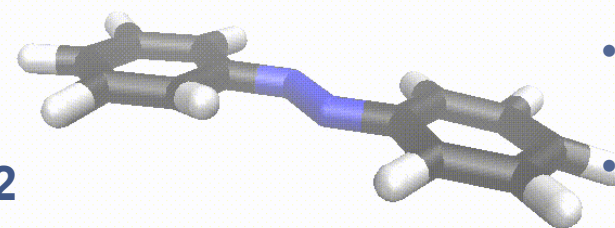


Selected publications:

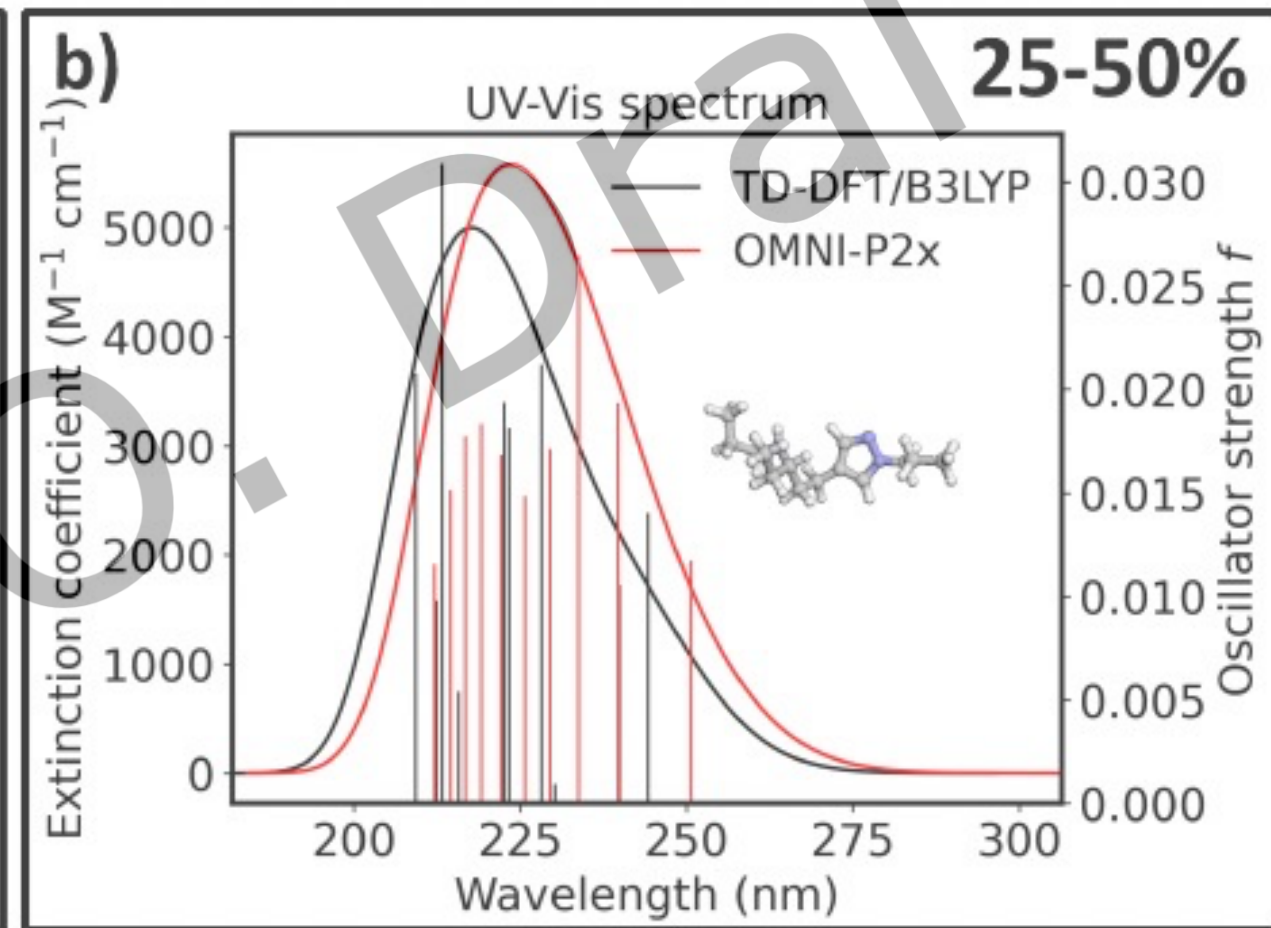
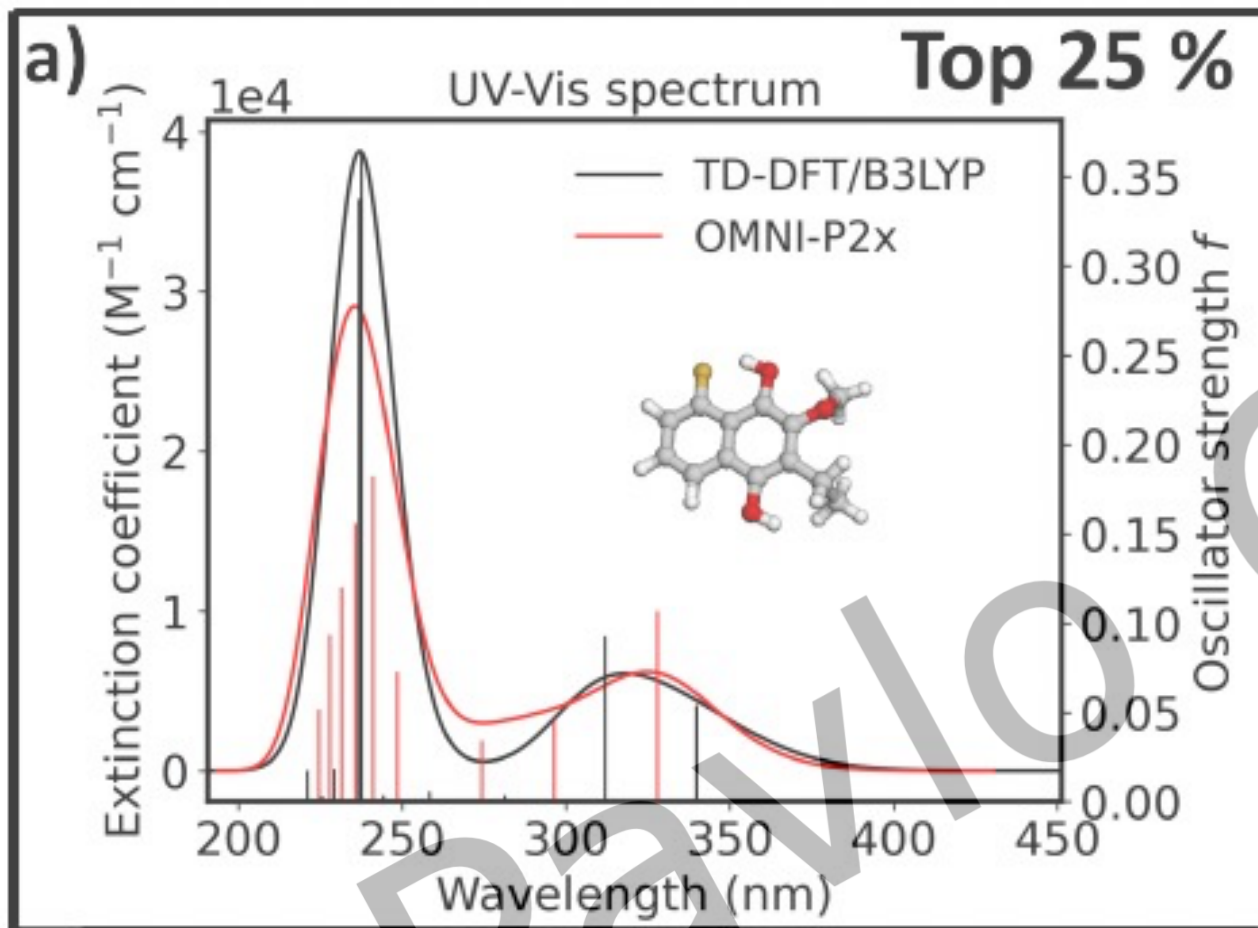
**OMNI-P2x:** Dral. *Nat. Commun.*, 2026, 17, 4949.

Novel architecture: Dral\*, et al. *npj Comput. Mater.* 2025, 11, 132

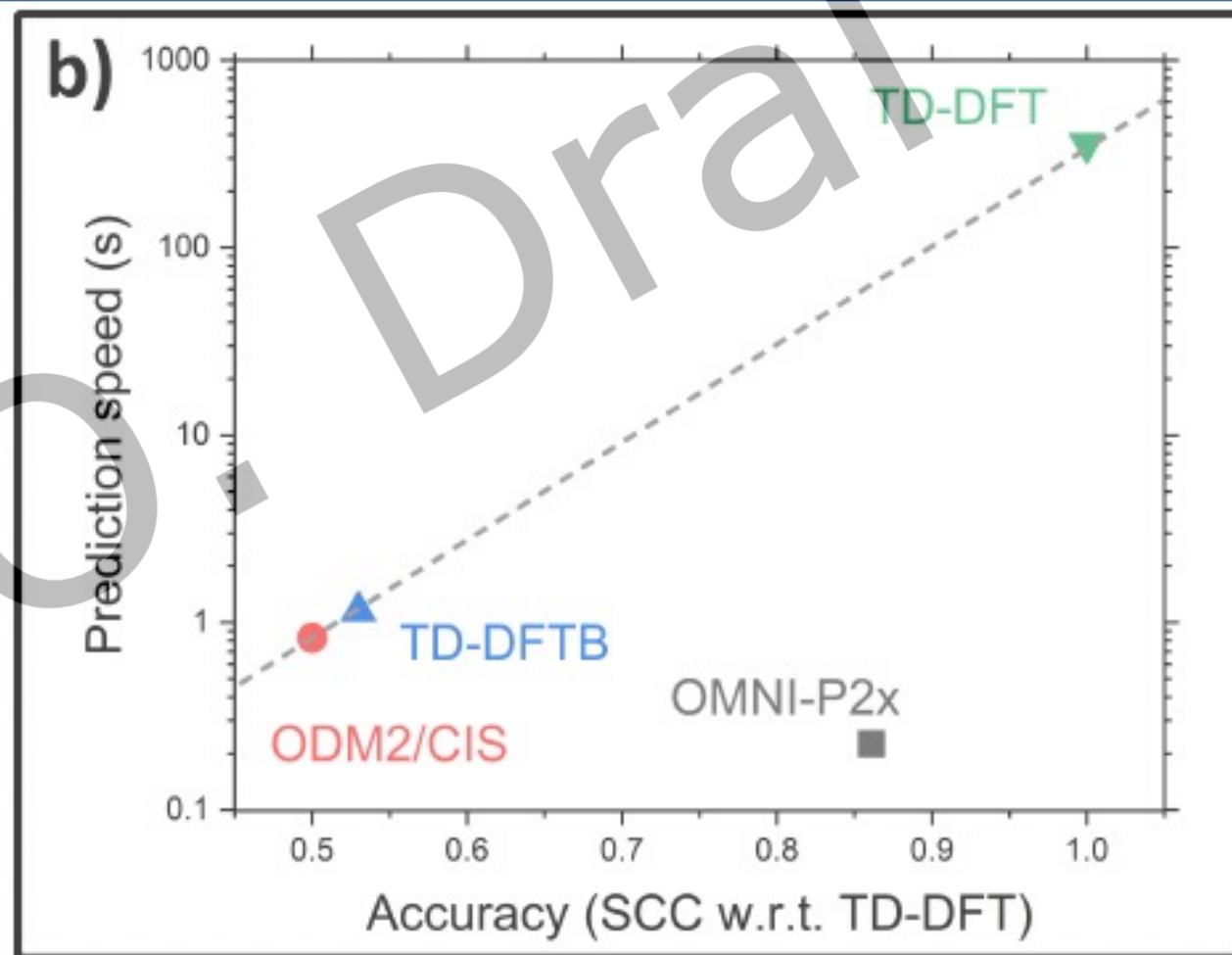
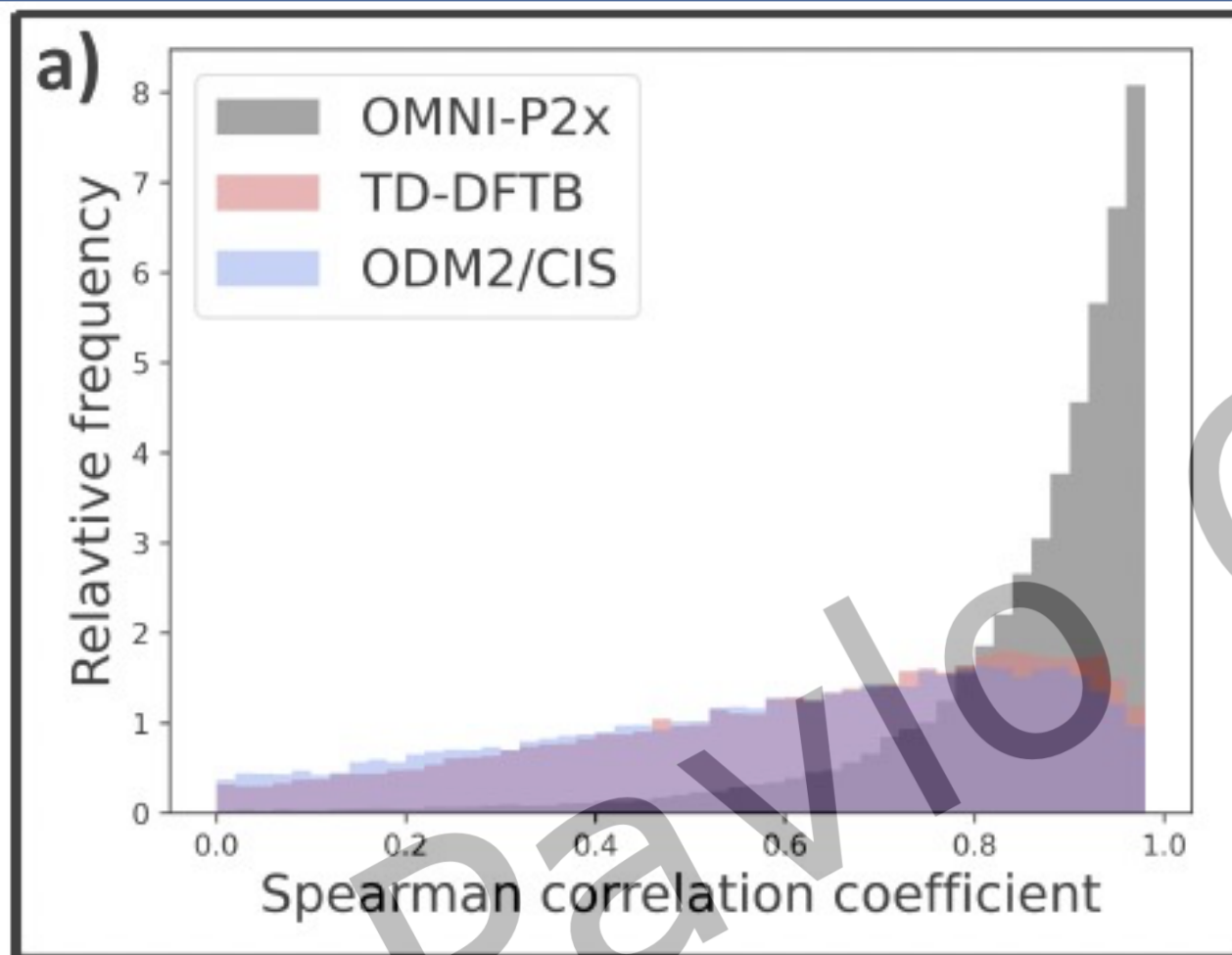
Dral\*, Barbatti\*. *Nat. Rev. Chem.* 2021, 5, 388

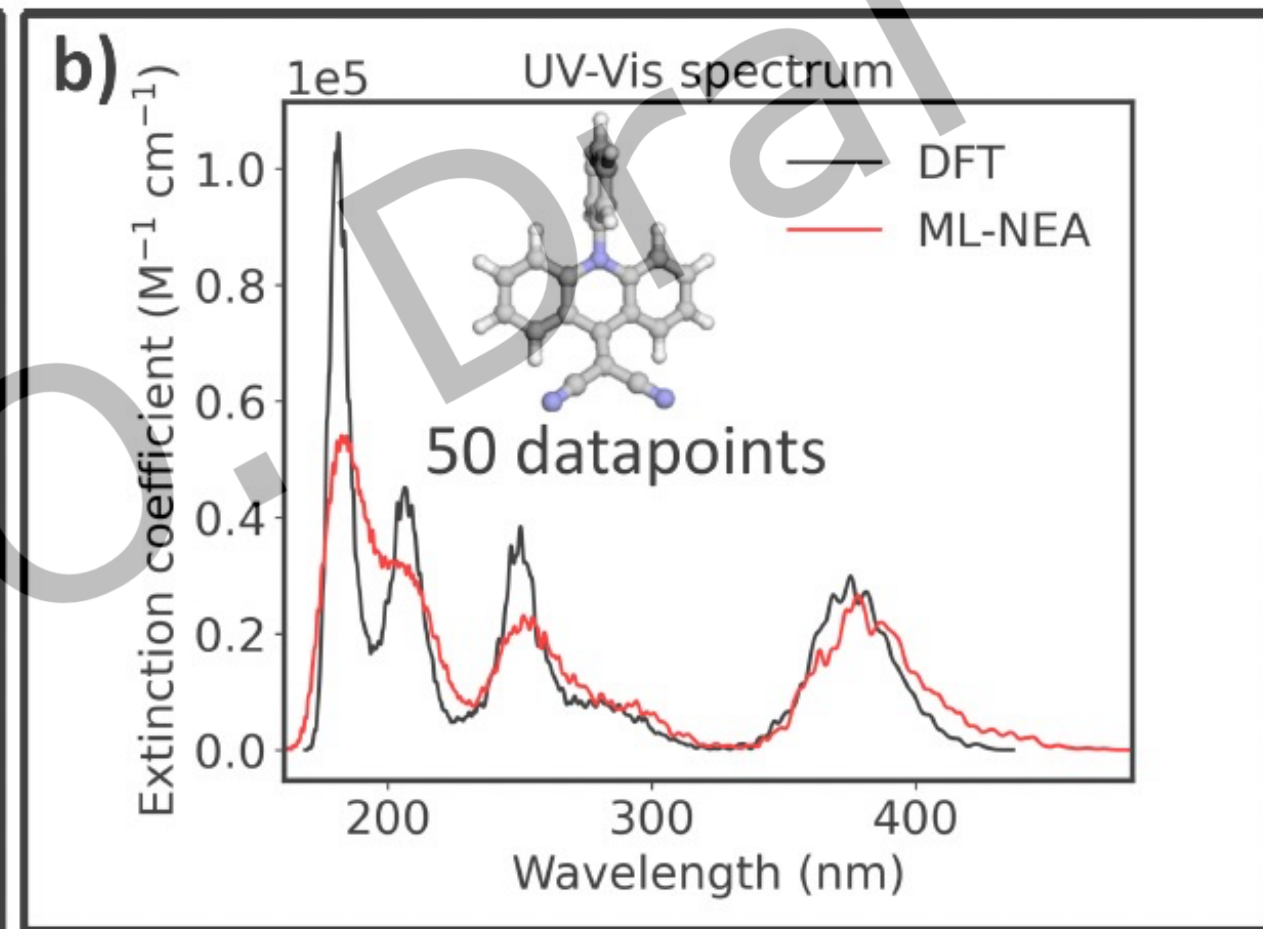
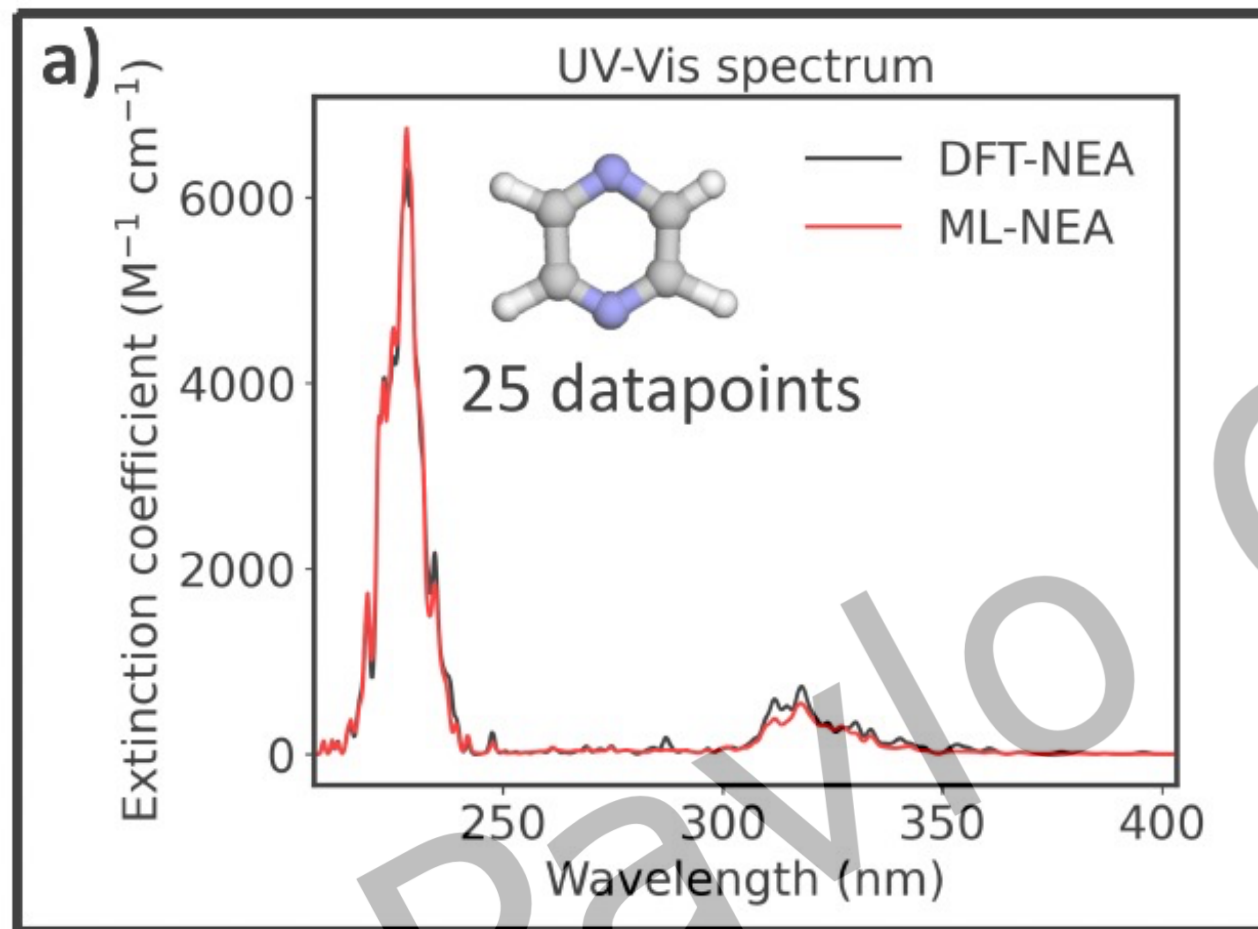


- photodynamics
- photoreaction quantum yields
- ...



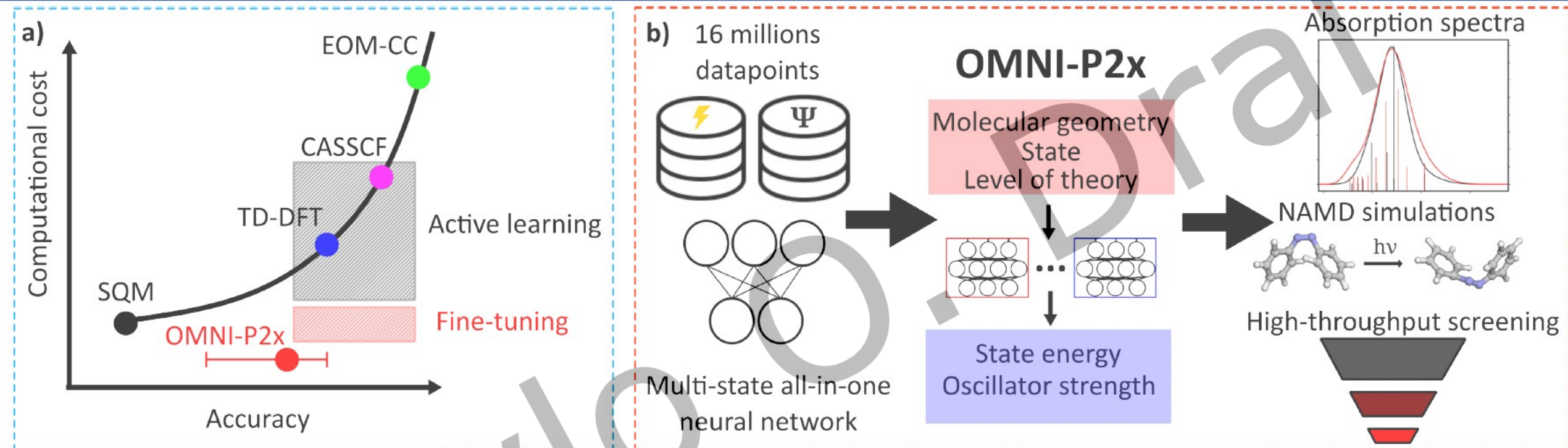




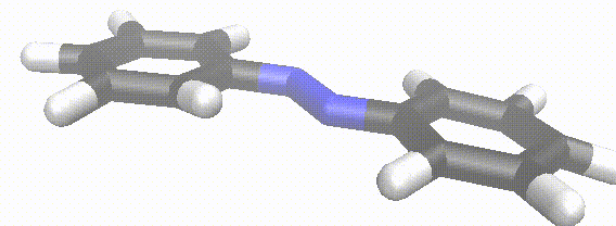


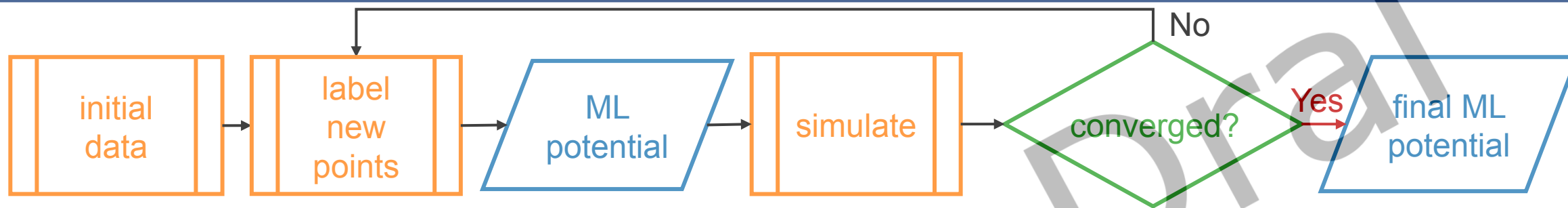
# NONADIABATIC DYNAMICS

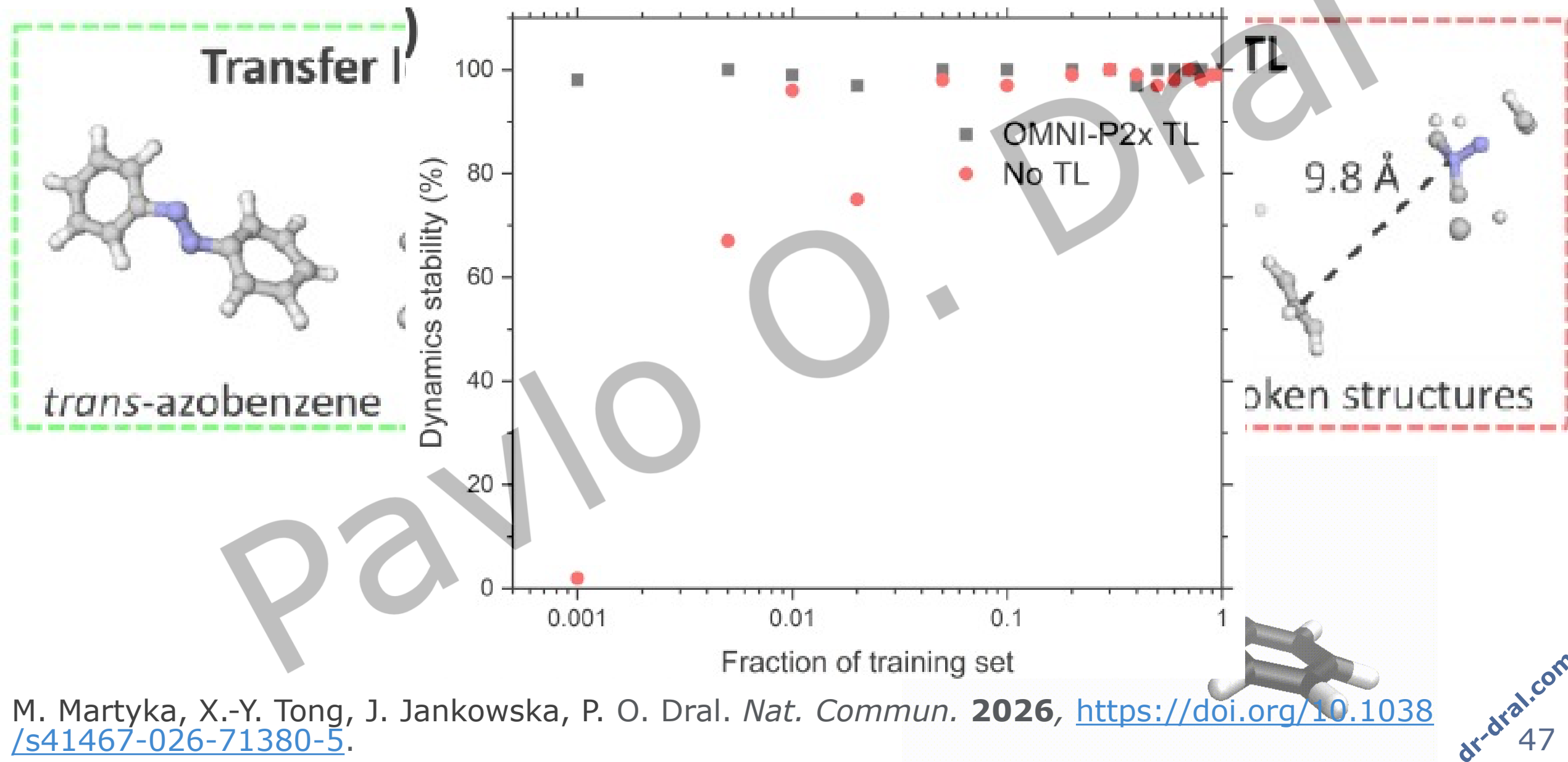




**Active transfer learning (ATL)** → robust nonadiabatic dynamics:









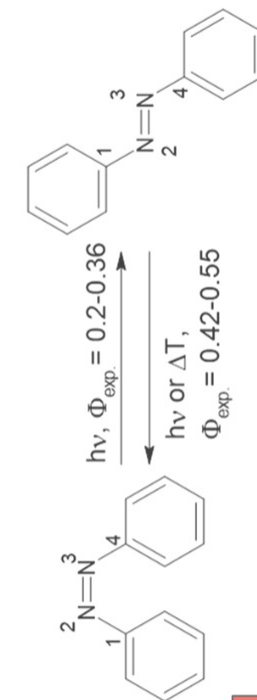
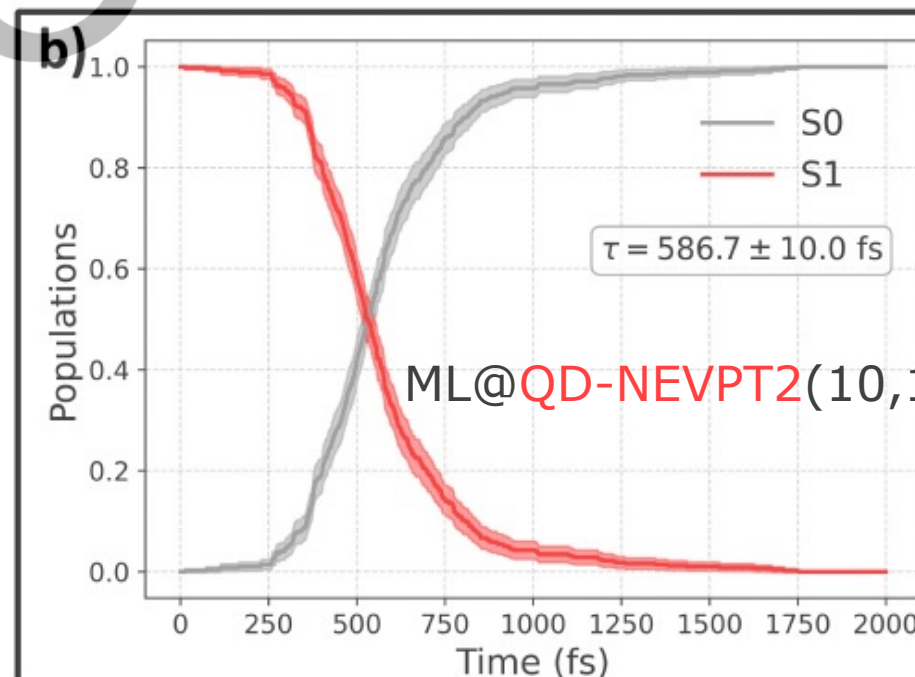
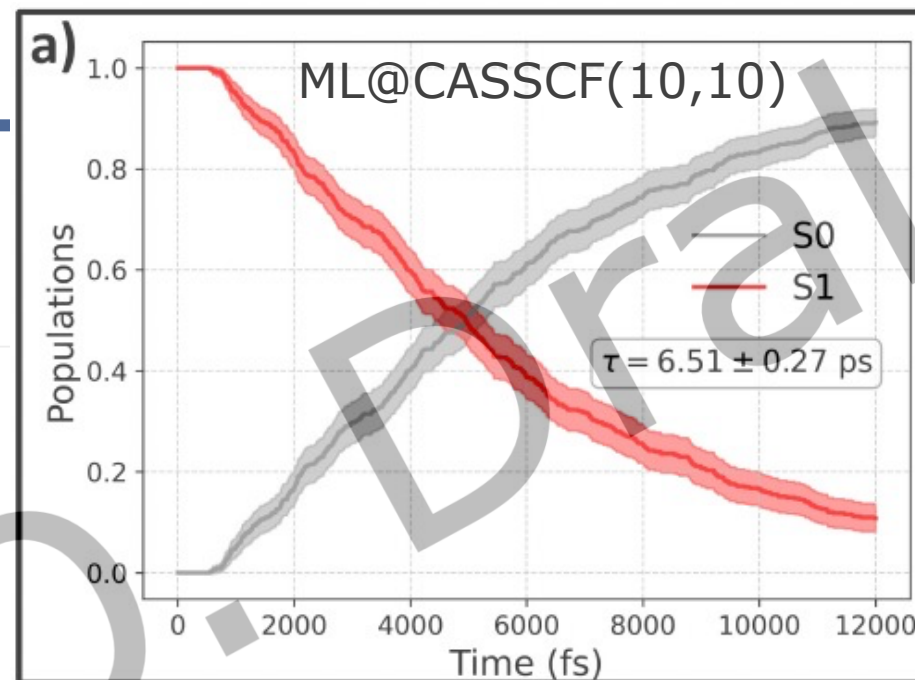
From the journal:  
**Chemical Science**

# Gradients not needed: ML-driven propagation of nonadiabatic molecular dynamics without reference gradients

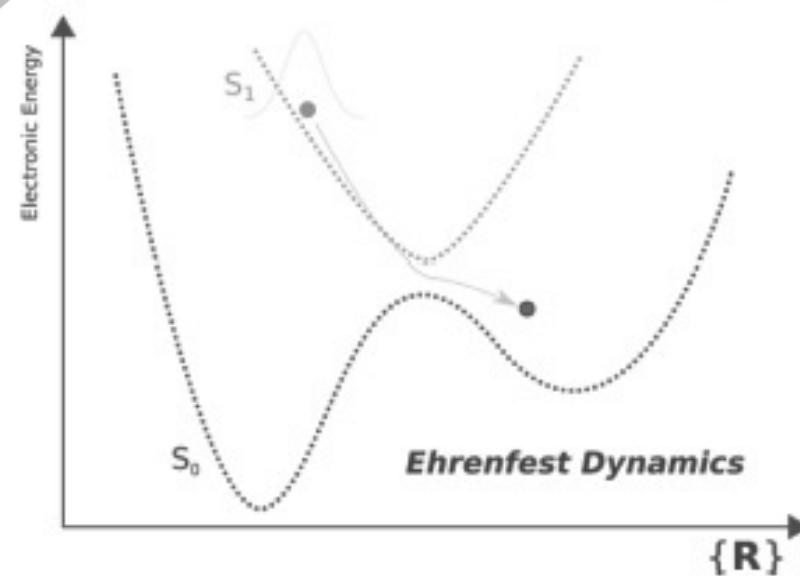
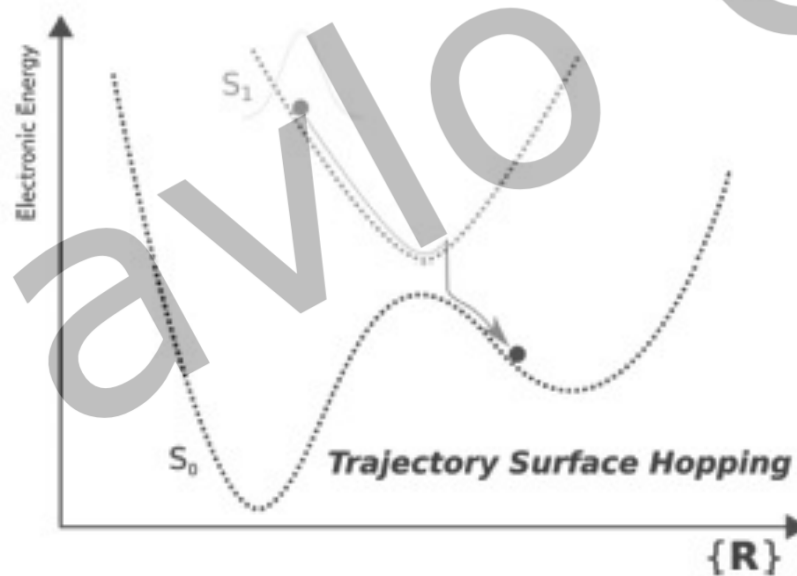
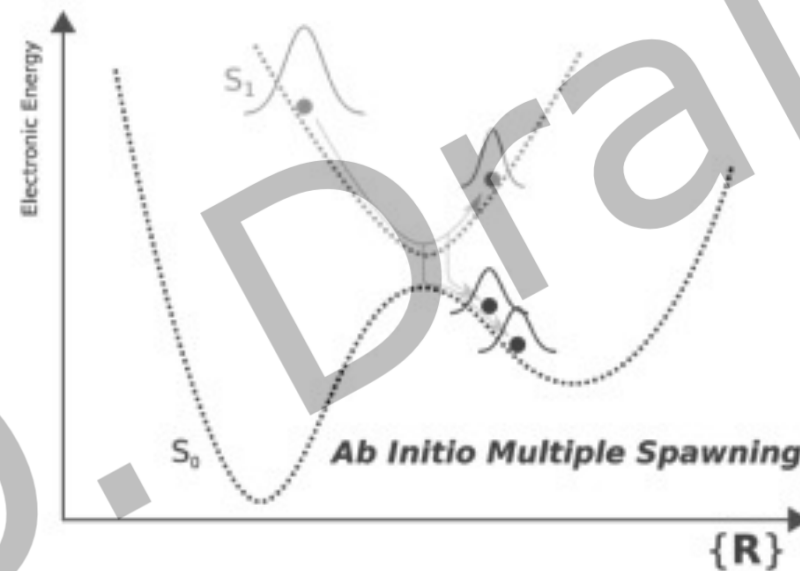
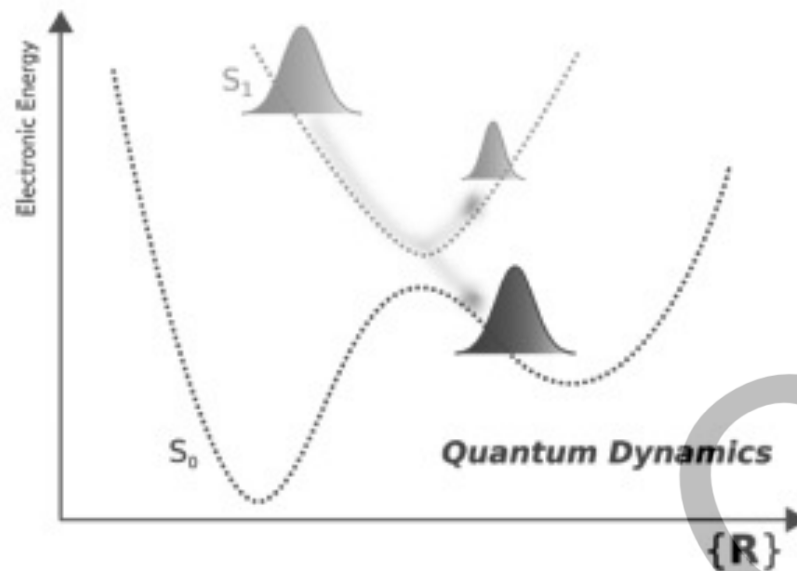
[Mikołaj Martyka](#), [Joanna Jankowska](#), [Hans Lischka](#) and [Pavlo O. Dral](#)

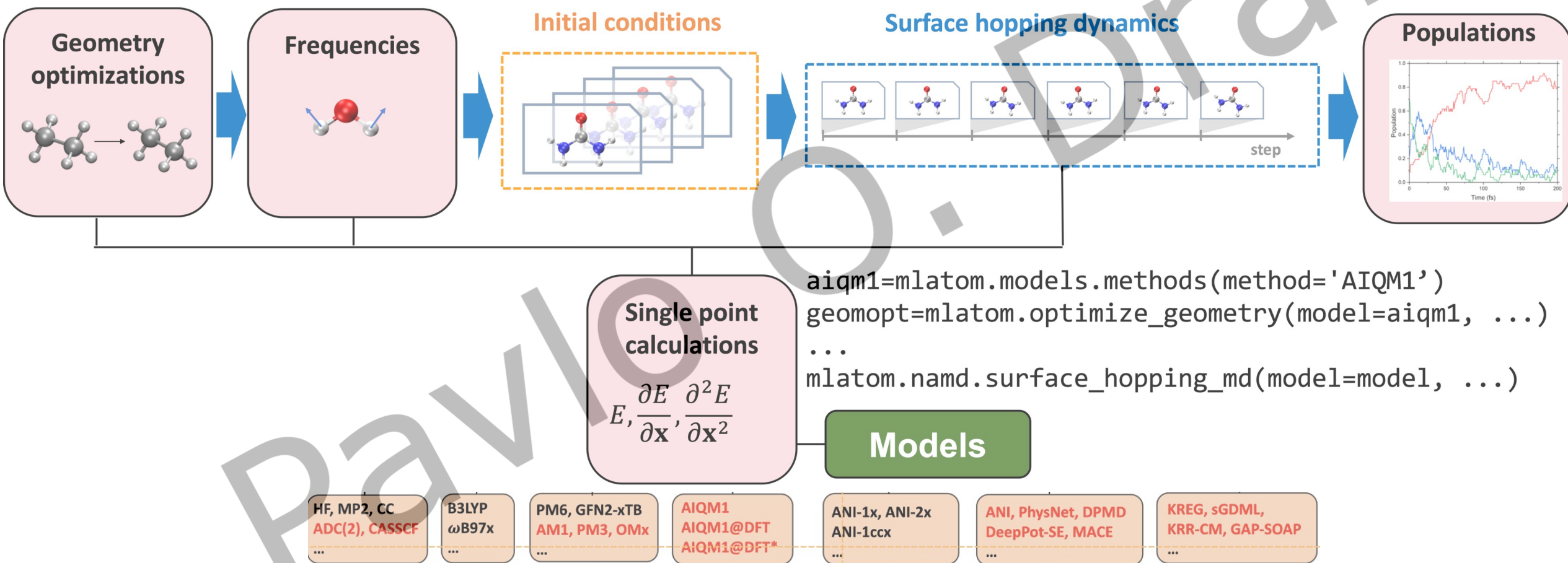
## Abstract

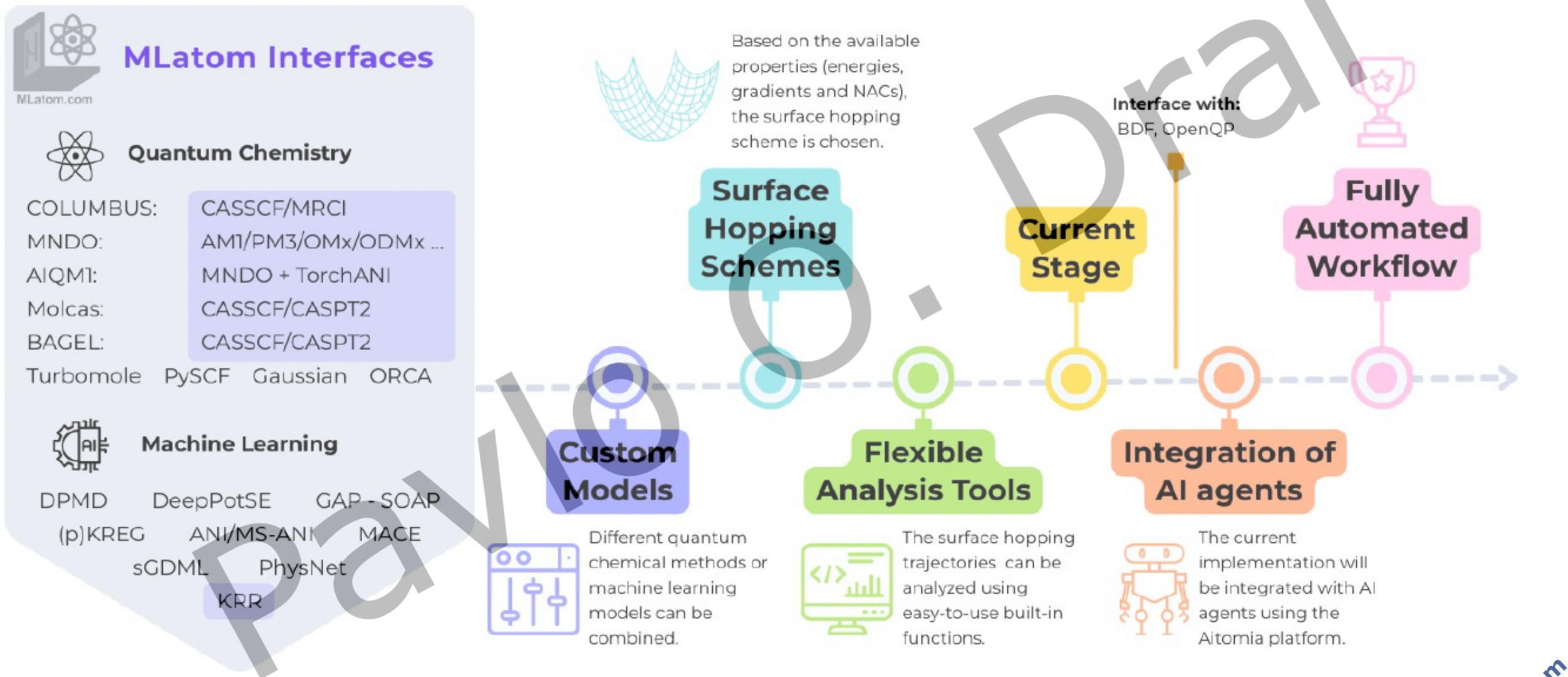
The recent development of machine learning (ML) methods for quantum chemistry has tremendously boosted the efficiency of molecular calculations. In this work, we use ML to enable nonadiabatic molecular dynamics (NAMD) simulations without access to the analytical energy gradients from the underlying electronic structure method. By fine-tuning our foundational model for excited-states, OMNI-P2x, on energies alone, and leveraging automatic differentiability to obtain forces, we eliminate the gradient computation bottleneck that restricts calculations, such as NAMD, to methods with available analytical derivatives. Firstly, we validate the method on the benchmark system, fulvene, demonstrating that gradient-free ML potentials accurately reproduce NAMD populations and dynamics across multiple levels of theory: AIQM1/MRCI, CASSCF, and MRSF-TDDFT. This enables, for the first time, performing dynamics at the QD-NEVPT2 level, where analytical gradients remain unavailable. We further benchmark the protocol on cyclohexadiene photoinduced ring-opening, where gradient-free training on XMS-CASPT2 energies reproduces reference dynamics with high accuracy, and compare them to QD-NEVPT2 results. Finally, we apply the approach to trans-azobenzene, a prototypical molecular photoswitch, by performing fully dimensional simulations of its photoisomerization dynamics at the CASSCF and QD-NEVPT2 levels, establishing the highest-level excited-state simulations of this photoreaction to date.



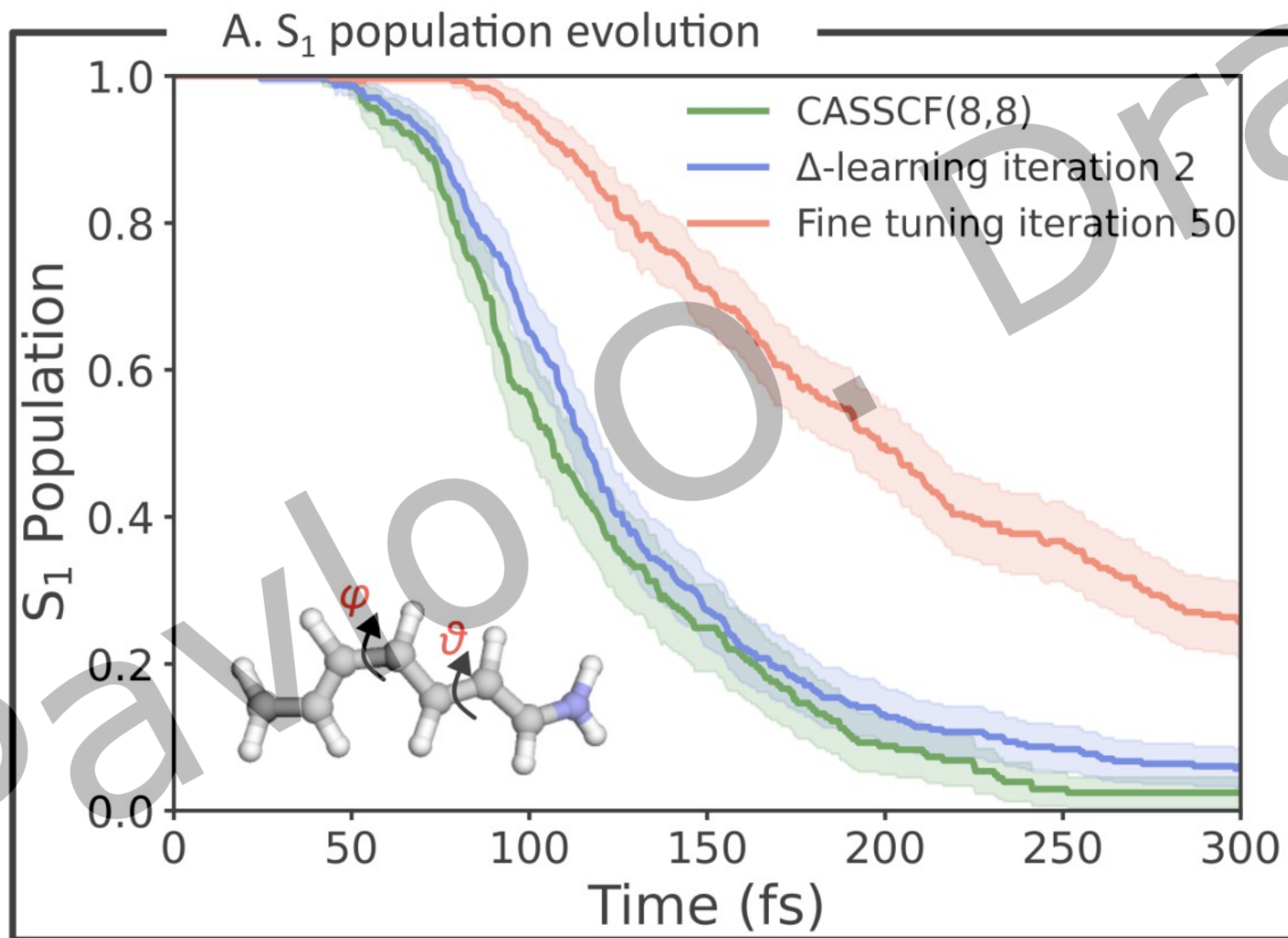








**Solution for  
molecules with  
many  
conformers**

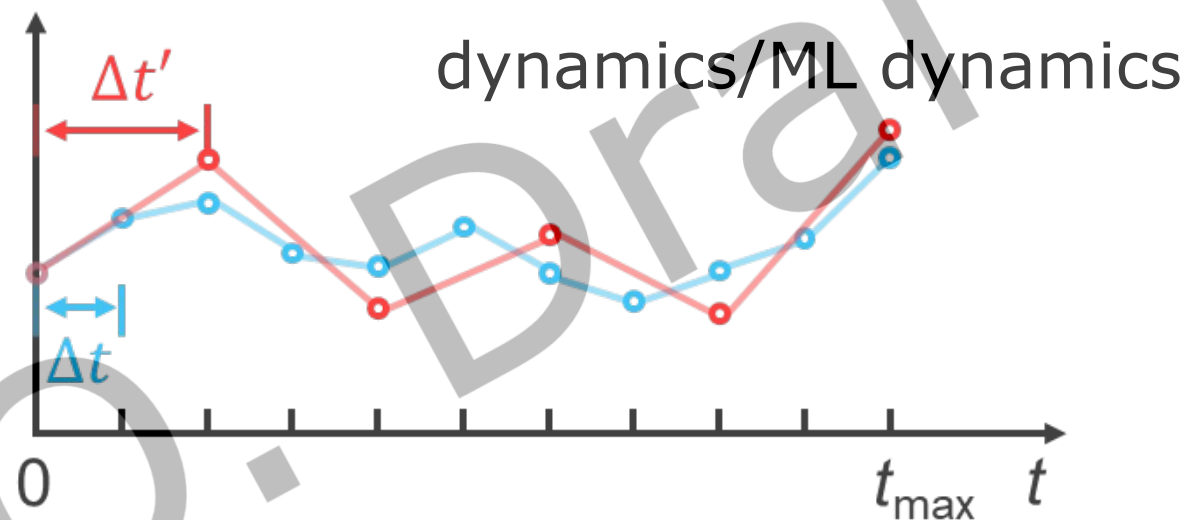




# Can we do better?

dynamics propagation is

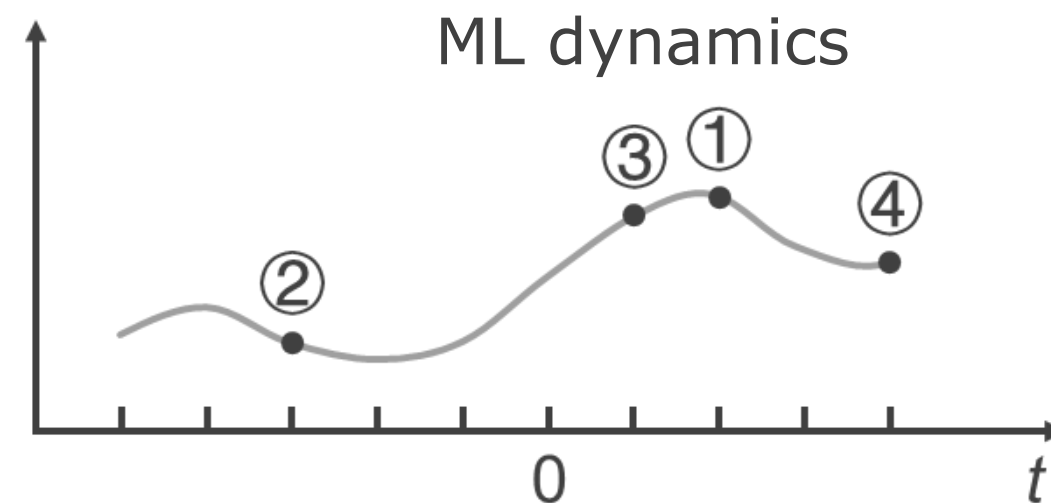
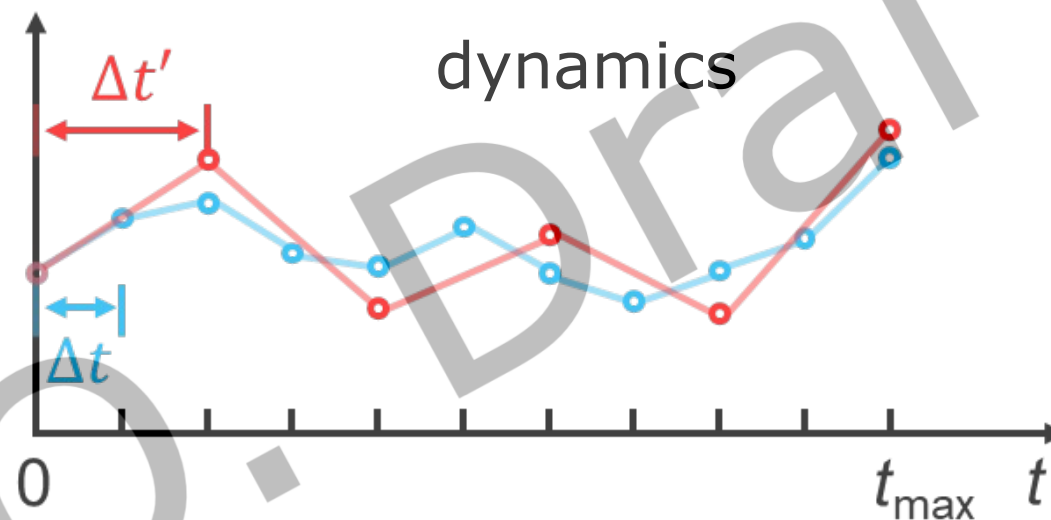
- computationally expensive
- recursive (iterative)



# Can we do better?

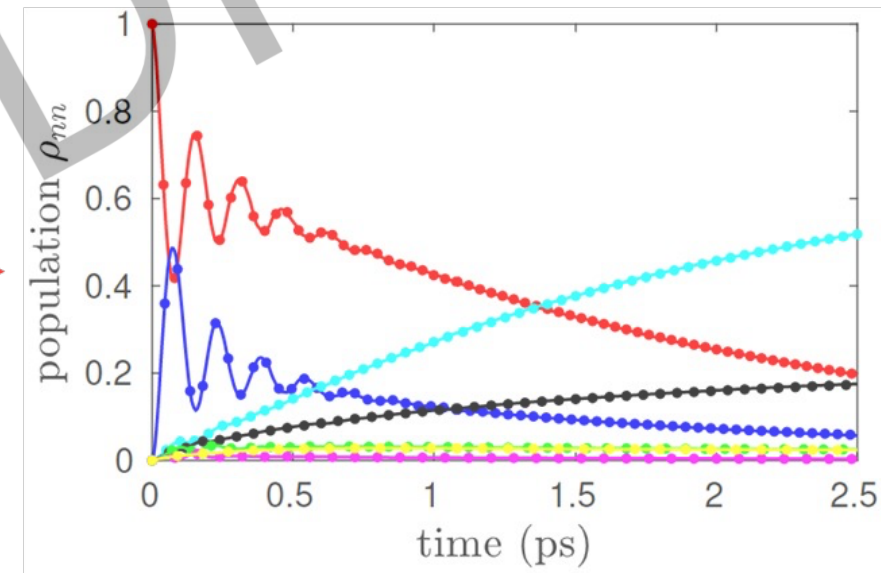
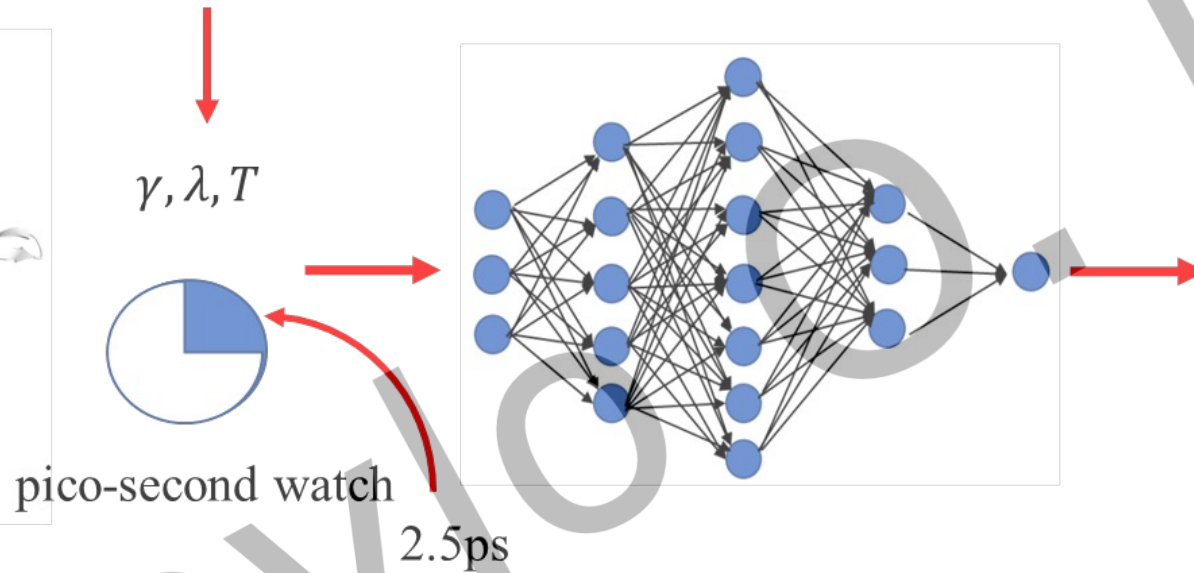
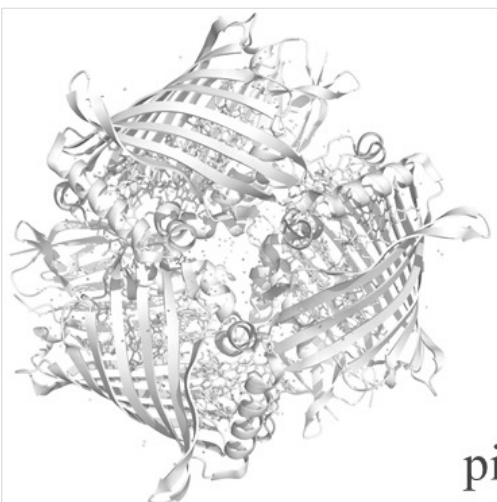
dynamics propagation is

- computationally expensive
- ~~recursive~~ (iterative)



$\gamma$  = characteristic frequency  
 $\lambda$  = reorganization energy  
 $T$  = temperature

PDB code: 3ENI

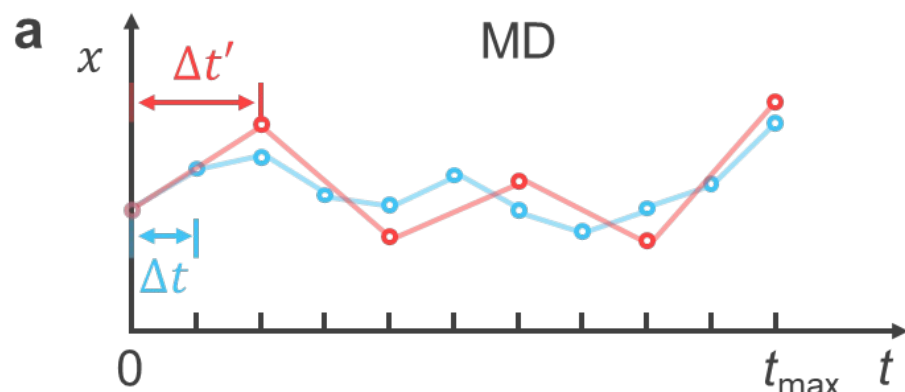


**Dots: Reference**  
**Line: AI-QD**

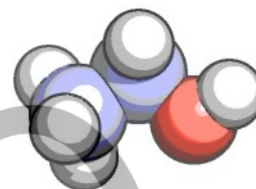
## 7-sites Fenna-Matthews-Olson (FMO) complex

A. Ullah, P. O. Dral. Predicting the future of excitation energy transfer in light-harvesting complex with artificial intelligence-based quantum dynamics. *Nat. Commun.* **2022**, 13, 1930

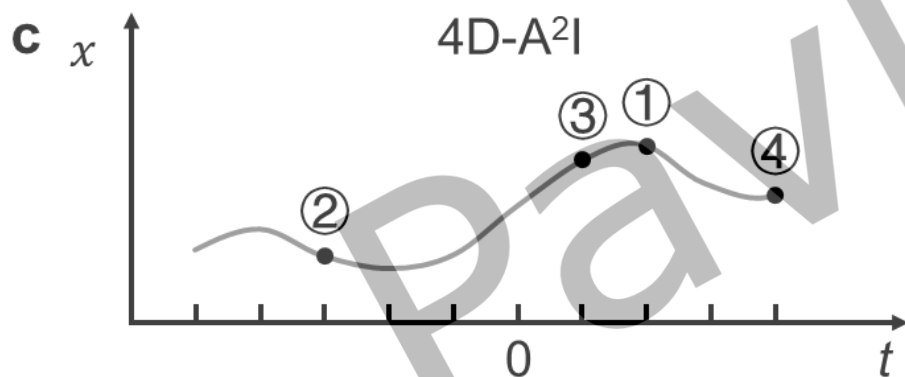
We introduced a radically different way to do MD, 100x faster than with ML potentials  
我们提出了一种全然不同的分子动力学方法，其速度比机器学习势快 100 倍。



Reference MD

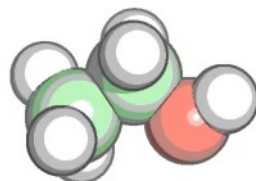


Almost everyone thought it was impossible due to the vast phase space of multi-atomic systems  
几乎所有人都认为这是不可能的，因为多原子体系具有巨大的相空间



Directly  
AI-generated MD

0.0 fs

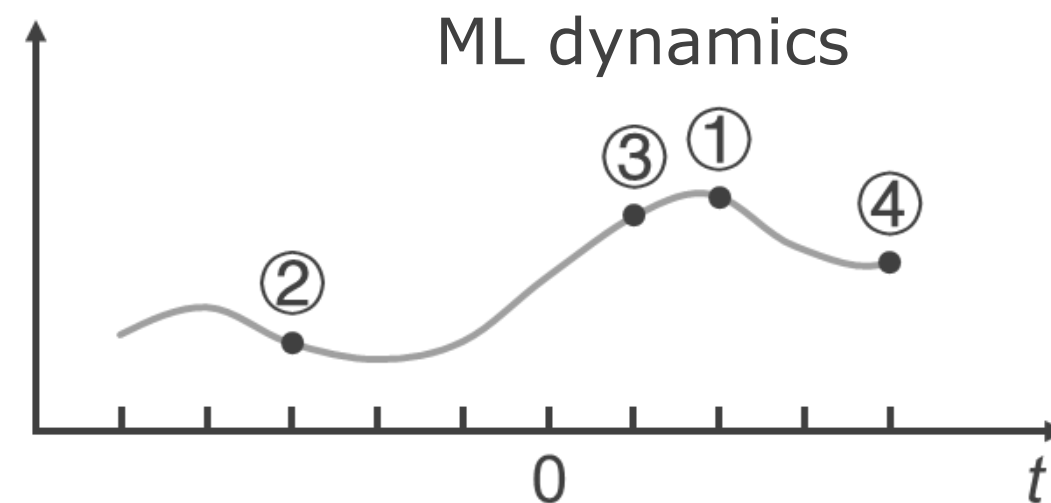
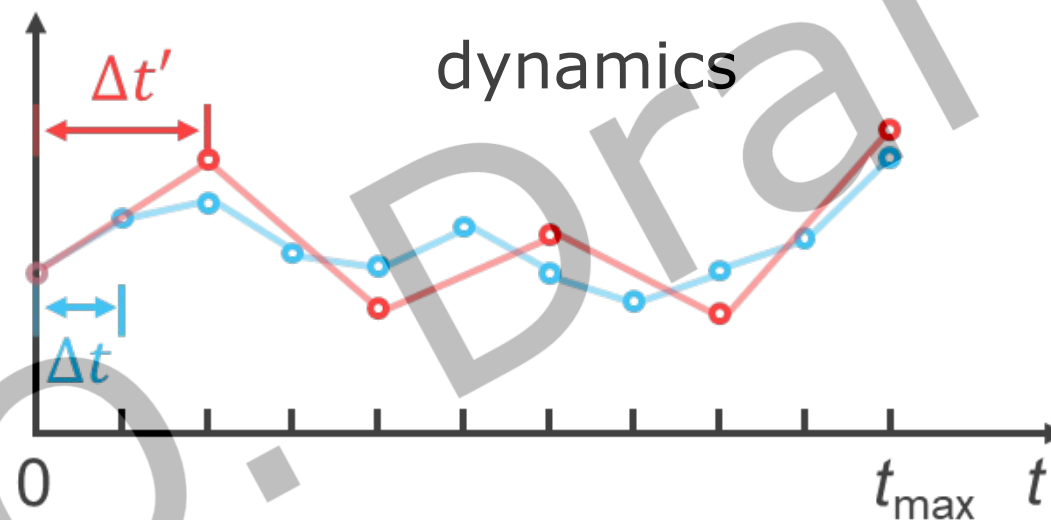


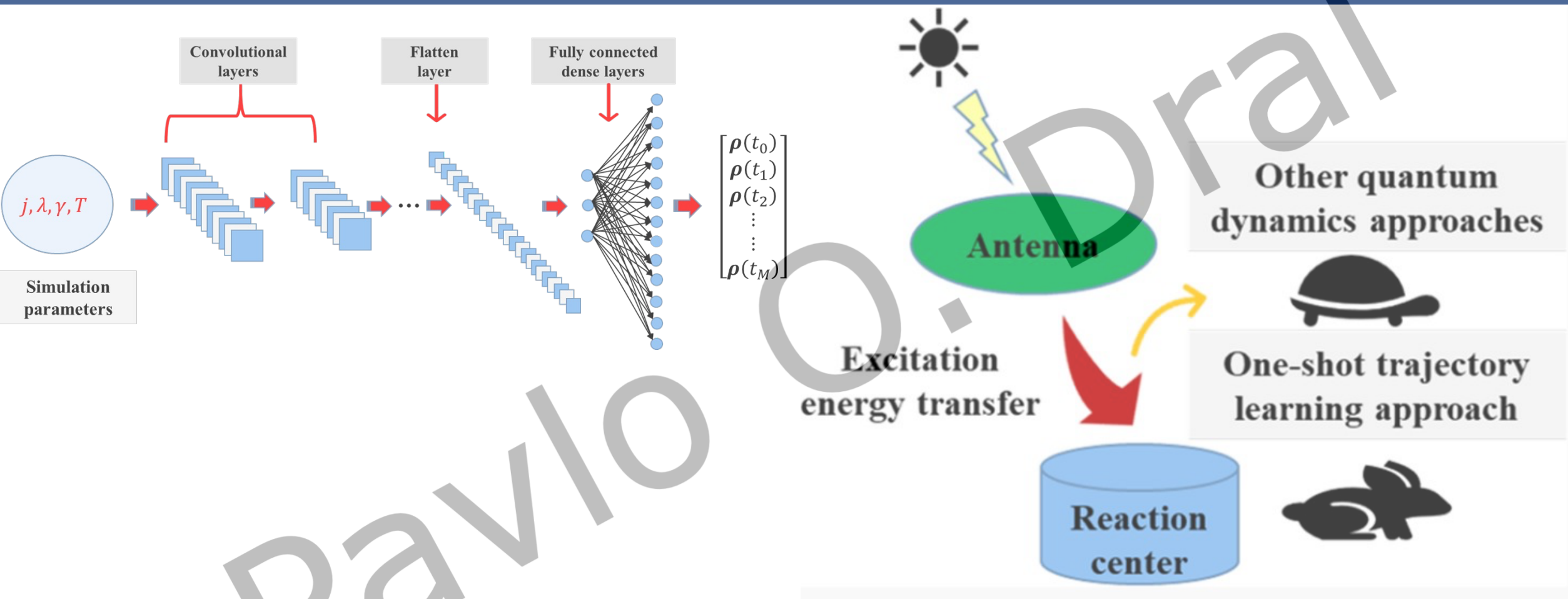


# Can we do even better?

dynamics propagation is

- computationally expensive
- ~~recursive~~ (iterative)





- **10 ps long dynamics in just 70 ms**
- good for massive simulation in parameter space





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CHEMICAL  
SOCIETY

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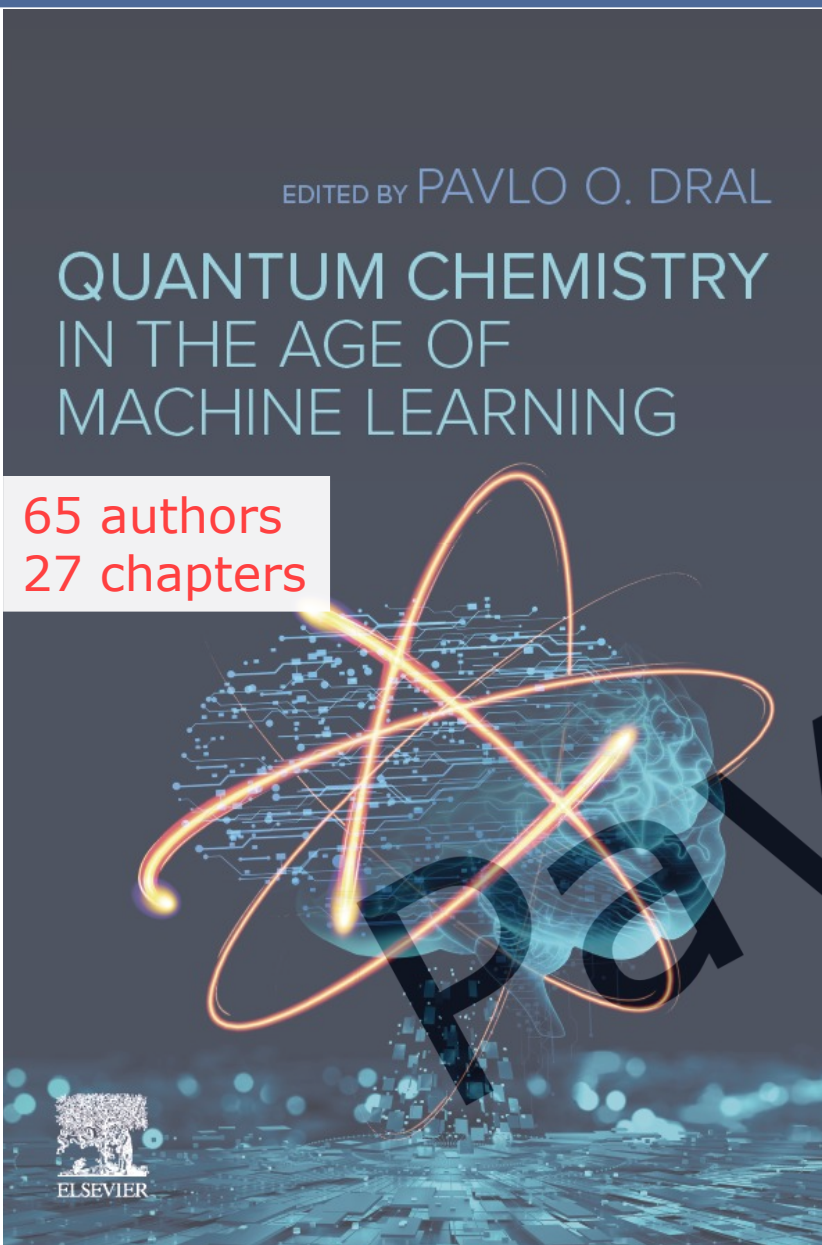
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2025 – Torun, Poland





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# Practical Guide to AI & ML Simulations

Pavlo O. Dral

Xiamen University, China

Nicolaus Copernicus University in Toruń, Poland

Aitomistic, Shenzhen, China

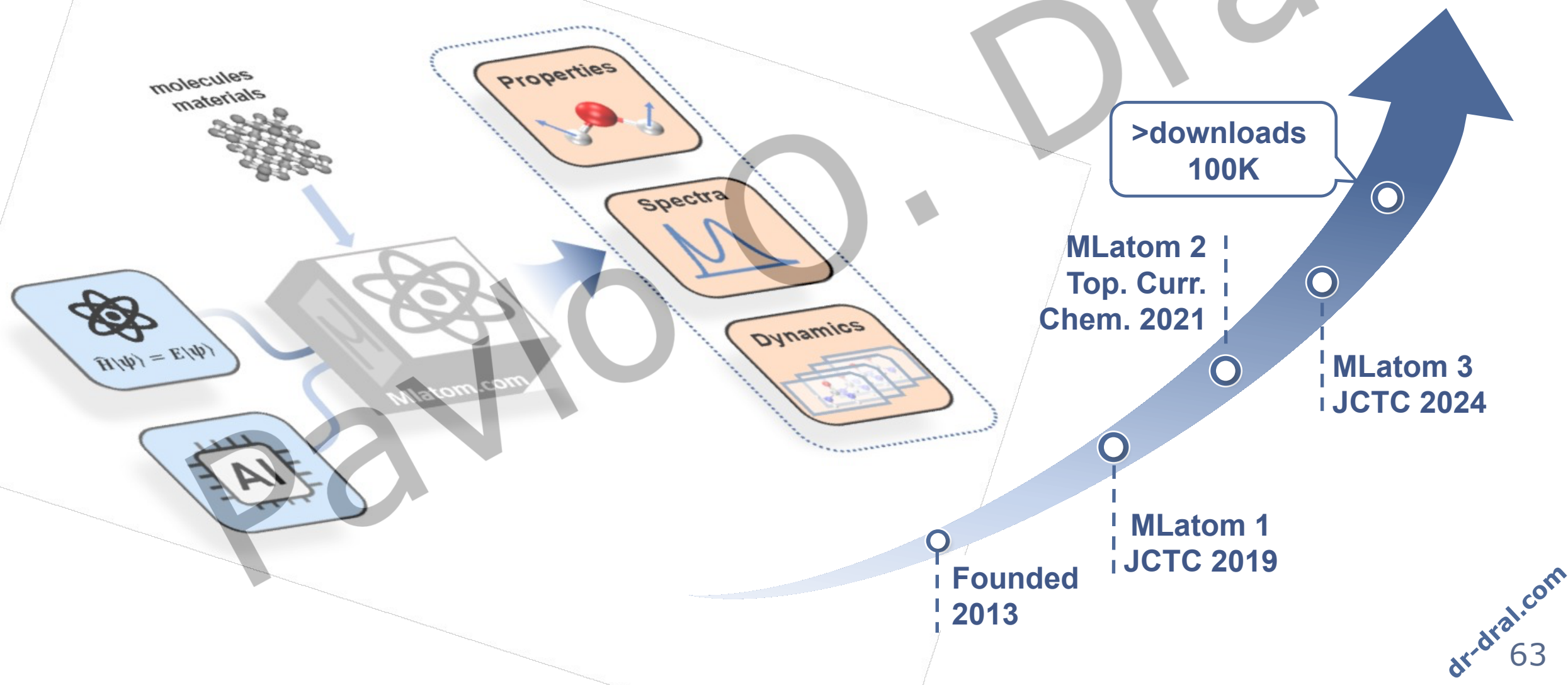
9 September 2026

dr-dral.com

# Our AI ecosystem for computational chemistry

We develop MLatom – an open-source software ecosystem integrating AI and quantum chemistry

<https://github.com/dralgroup/mlatom>





 **add interface to read "energy" and "gradient" from BDF**

#38 opened on Apr 9 by ilcpm • Review required

Quantum chemical programs:



Gaussian



ORCA

PySCF

Turbomole

COLUMBUS

## MLQD

[MLQD: A. Ullah, P. O. Dral.  
*Comput. Phys. Commun.* **2024**,  
294, 108940]

Semi-empirical quantum chemical programs:



SPARROW

MNDO

Machine learning programs:

PhysNet

GAP

TORCHANI



SGML

hyperopt

Dynamics and other atomistic simulation:

ASE



NX  
newtonx.org



# Machine Learning

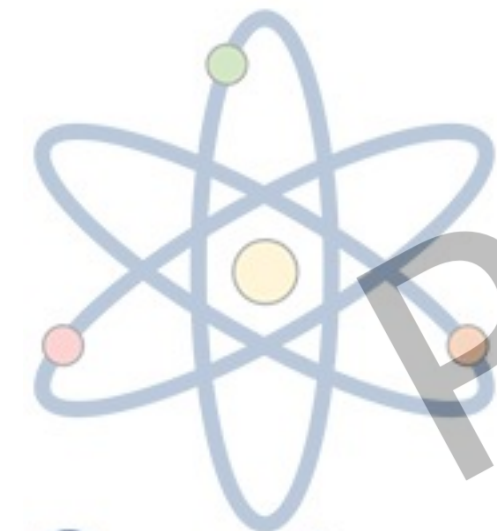


# Simulations

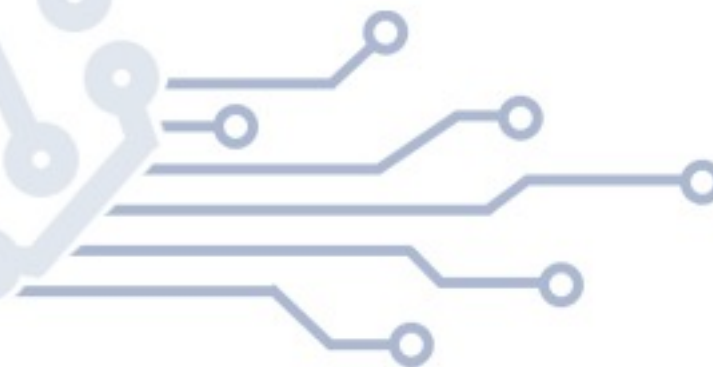


# Mlatom

>100000 downloads

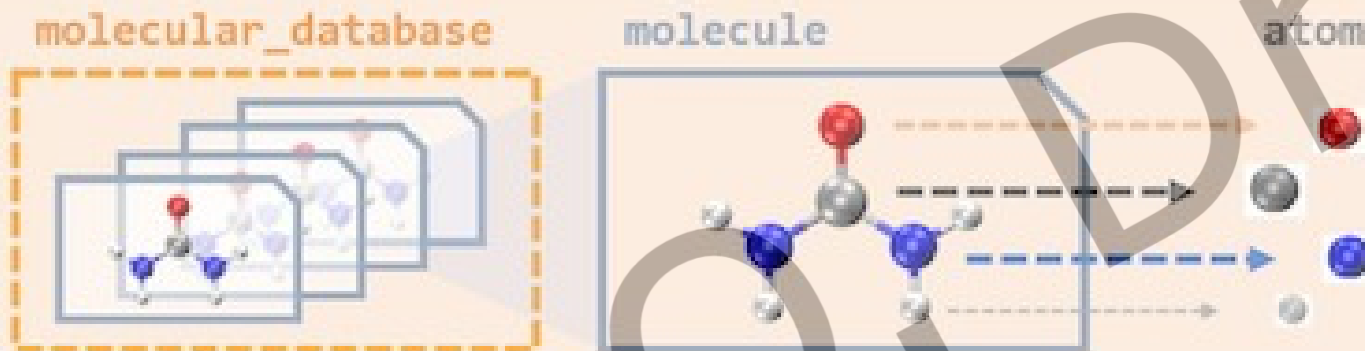


# Quantum Chemistry



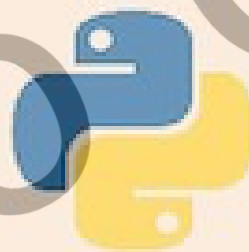
# Data

## ➤ Data in different formats and types **V3**



## ➤ Python API **V3**

```
import mlatom as ml
```

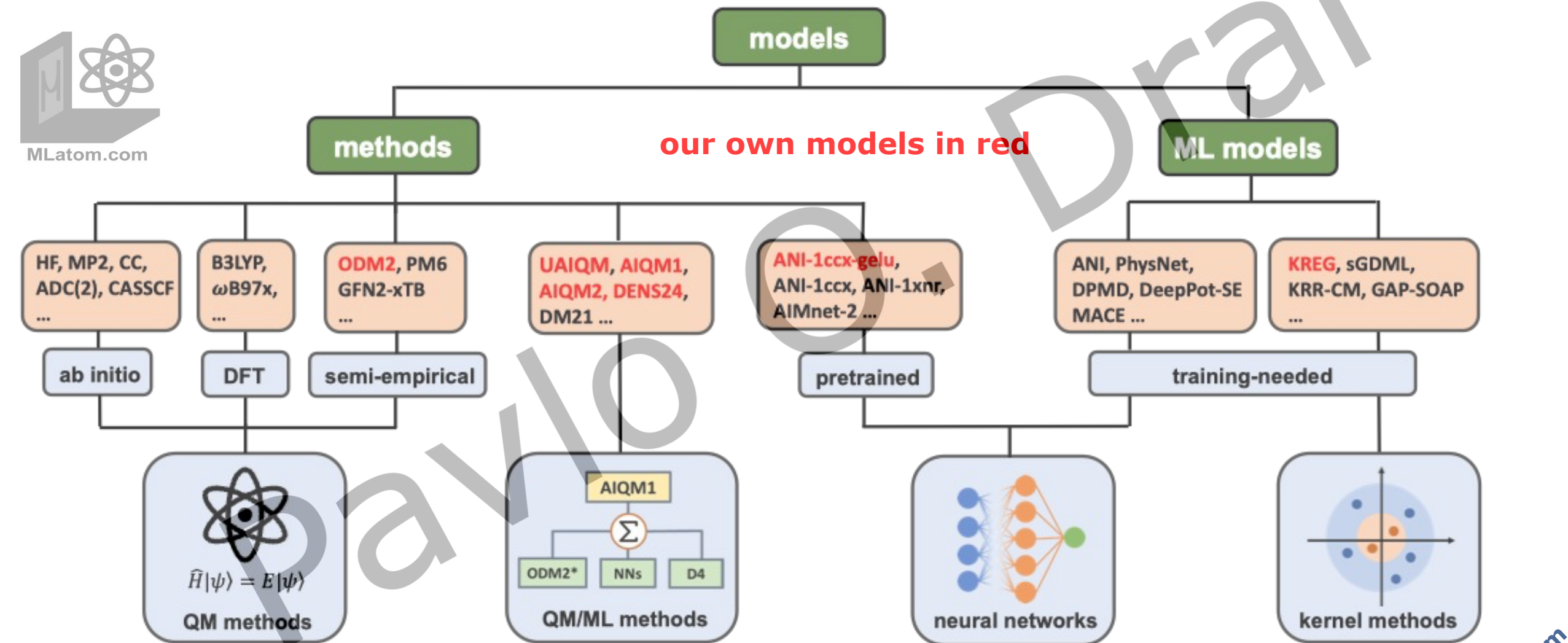


## Input file

```
ANI-1ccx
geomopt
xyzfile=init.xyz
optxyz=opt.xyz
```

## ➤ Command line

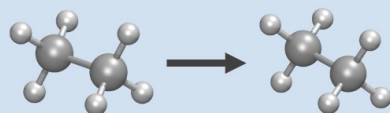
```
> $mlatom ANI-1ccx geomopt xyzfile=init.xyz optxyz=opt.xyz
```



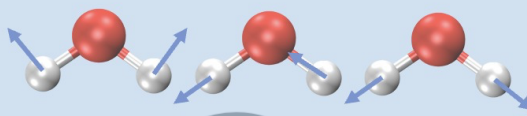
Single point calculations

Energies, forces, Hessian matrix...

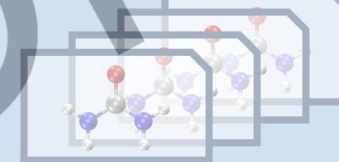
Geometry optimizations



Frequency calculations



Molecular dynamics



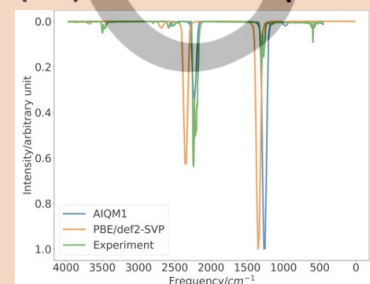
+PIMD coming soon...

Thermochemistry calculations

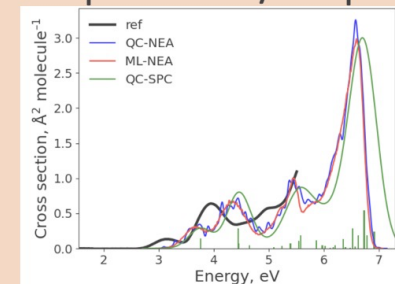
Heat of formation:

$$\Delta H_{\text{at},T} = \left[ \sum_A H_T(A) \right] - H_T$$

(Ro)vibrational spectra



One-photon UV/vis spectra



Properties & spectra

Simulations

Quantum dissipative dynamics

Two-photon absorption spectra



github.com/dralgroup/mlatom

README Code of conduct Contributing Apache-2.0 license Security

## Local installation

The easiest way to run MLatom is [online](#) — no installation. To install locally:

```
python3 -m pip install -U mlatom
```

That pulls in the required dependencies automatically (including the PyTorch/TorchANI and geometry-optimization backends the **AIQM2** quick start uses). A few features need one more package — install only what you use:

| for                                                                                   | install                                                    |
|---------------------------------------------------------------------------------------|------------------------------------------------------------|
| PySCF methods: single point, TDDFT, frequencies and thermochemistry, densities, Raman | <code>pip install pyscf</code>                             |
| AIMNet2 models                                                                        | <code>pip install aimnet</code>                            |
| format conversions (SMILES to xyz and back)                                           | <code>conda install -c conda-forge openbabel</code>        |
| the ASE interface (ASE optimizers and dynamics) and MACE                              | <code>pip install ase</code>                               |
| KREG_API backend ( <code>ml_program='MLatomF'</code> needs none of this)              | <code>conda install -c conda-forge mkl mkl-service</code>  |
| MDtrajNet / MLTPA / DMC / hyperparameter search                                       | <code>pip install e3nn / rdkit / pyvibdm / hyperopt</code> |

AIQM2 additionally needs the DFT-D4 program, installed via conda:

```
conda install -c conda-forge 'dftd4==3.6.0' # the version MLatom is tested against
export dftd4bin=$(which dftd4) # point MLatom at the dftd4 executable
```

See the [installation guide](#) for the full dependency list and other methods.

## Quick start

Optimize the geometry of a water molecule with **AIQM2** — an AI-enhanced quantum-mechanical method (native to MLatom, CHNO elements) that reaches beyond-DFT accuracy at semi-empirical cost:

```
import mlatom as ml

mol = ml.data.molecule.from_xyz_string('''
O  0.00000  0.00000  0.11779
H  0.00000  0.75545 -0.47116
H  0.00000 -0.75545 -0.47116
''')

aiqm2 = ml.methods(method='AIQM2')
opt = ml.optimize_geometry(model=aiqm2, initial_molecule=mol).optimized_molecule
print(opt.energy) # optimized energy in hartree (= -76.3838)
```

The same calculation as an input file ( `geomopt.inp` , with `init.xyz` holding the geometry):

```
AIQM2          # method
geomopt        # task: geometry optimization
xyzfile=init.xyz # input geometry
optxyz=opt.xyz  # output geometry
```

```
mlatom geomopt.inp
```

Prefer zero setup? Run these online on the [Aitomistic Hub](#) or [Aitomistic Lab@XMU](#) (both powered by [Protomia](#)) — no installation needed.

<https://github.com/dralgroup/mlatom>

<https://github.com/dralgroup/mlatom-skills>

B3LYP/6-31G\*

geomopt

xyz GFN2-xTB

geomopt

xyz MP2/cc-pVDZ

geomopt

xyz ANI-1ccx

geomopt

xyz MLmodelType=ANI MLmodelIn=ani.pt

geomopt

xyz UAIQM  
geomopt  
xyzfile='5'

|   |               |               |               |
|---|---------------|---------------|---------------|
| C | 0.0000000000  | 0.0000000000  | 0.0000000000  |
| H | 1.0870000000  | 0.0000000000  | 0.0000000000  |
| H | -0.3623333220 | -1.0248334322 | -0.0000000000 |
| H | -0.3623333220 | 0.5124167161  | -0.8875317869 |
| H | -0.3623333220 | 0.5124167161  | 0.8875317869  |
| , |               |               |               |

atom.xmu.edu.cn/course

## Courses

Browse available courses and enroll to start learning.

### Computational chemistry and AI

Course materials for the part of the course "Computational Chemistry and AI" taught by Prof. Pavlo O. Dral in the Spring semester of 2026

[View](#)

atom.xmu.edu.cn/courses/comchem\_2026/qc\_intro.html#lecture1-ml

### Computational chemistry & AI

Search docs

#### TABLE OF CONTENT

- Preface
- Introduction to quantum chemistry and machine learning
  - Slides
  - Quantum chemical levels
  - Finally, your first ML calculations!
  - Homework
  - Further reading
- Why does DFT work? Improving upon it
- Universal machine learning models
- Machine learning basics
- ML for PES
- Transfer learning

matter how well you do ML but if it is trained on a bad QM method, it will also be bad ("garbage in - garbage out").

Without further ado, let's jump into our first ML calculations with Task 1.3 (in in-class exercises, you have 10 min, send answers on WeChat and after everyone finishes, write down the answer on a blackboard):

#### Exercise ML-basics-1.

Calculate the bond length in H<sub>2</sub> molecule with machine learning (ML) (see instructions below).

Write down:

- Bond length you obtained.
- How much time it took to complete the calculations.

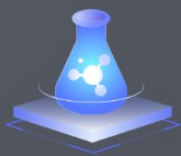
The MLatom input file ([h2\\_opt\\_kreg.inp](#)) is pretty self-explanatory:

```
geomopt          # Request geometry optimization
MLmodelType=KREG # with the ML model of the KREG type
MLmodelIn=energies.unf # in energies.unf file
XYZfile='
2

H      0.000      0.000      0.000
H      0.000      0.000      0.800
'
optXYZ=eq_KREG.xyz # optimized geometry output
```

You also need to upload the file with ML model [energies.unf](#) to the same folder as the input file.

← → ↻ ⚙ atom.xmu.edu.cn/login



**Aitomistic Lab@XMU**  
A Platform for Aitomistic Simulations at Xiamen University

**Welcome Back!**  
Sign in to your account to continue.

\* Email Address/Username  
  
Please input your email address or Username

\* Password  
  
Please input your password!

By signing up or logging in, you consent to Aitomistic Lab@XMU's [Terms of Use](#).

☒ Remember my account

[Don't have an account? Sign Up](#) [Forgot password?](#)

<https://atom.xmu.edu.cn> (available for free when registered with an academic email)

<https://aitomistic.xyz> (registration and basic use free for everyone)



## Create your account

Sign up to get started.

\* Username

Username cannot be empty

\* First Name

\* Last Name

\* E-mail

Please register with your academic e-mail address.  
Accounts from other addresses can be verified only manually and may be declined.

\* Verification Code

Get Code

Haven't received the code? Please check your spam folder or try registering with a different email address.

\* Password

\* Confirm Password

\* Phone Number

\* Region

\* Institute

By signing up or logging in, you consent to Aitomistic Lab@XMU's [Terms of Use](#), [Privacy Policy](#) and [Cookie Policy](#).

Register

<https://atom.xmu.edu.cn>  
(available for free  
when registered with an  
academic email)

<https://aitomistic.xyz>  
(registration and basic  
use free for everyone)

You Can Now Use Aitomistic Lab@XMU! Introduction to Aitomistic Lab@XMU with Usage Examples

Dear  ,

thank you for registering at Aitomistic Lab@XMU! We have approved your access and you may start using it by logging in to <https://atom.xmu.edu.cn>.

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Sincerely yours,

Your Team at Aitomistic Lab@XMU

Introducing Aitomistic Lab@XMU



EXPLOR...

☐ Show hidden
 ☐ Only show shared

demo

input

jupyter

python

uvvis

ethanol\_init.xyz

File deleted successfully

Drop your files (xyz, docx, etc.) here to upload

Create new file

Quick Instructions

Create demo folder

Protomia Solution 2.8.2

Hi, I'm Protomia. What can I help you with today?

You can drop any files here to add them to the context, and ask me to analyze or work with them.

20260827-29\_XACS-26

Back/Forward

| Name             |
|------------------|
| 01.png           |
| 20260829_AI.pptx |

Recents

Shared

Favorites

Applic...

Downl...

Movies

Music

Pictures

?

dr-dral.com 76





## calculations

## 16ITA

## 16ITA\_dimer

## core\_structure\_of\_dimer

## opt\_freq\_full\_structure

## UAIQM

16ITA.json

16ITA.json.zip

16ITA\_crystal.xyz

16ITA\_dimer\_ini.xyz

16ITA\_uaiqm\_opted.xyz

2ed9d415-40cf-46fb-k

76de54c7-ccbf-4415-a

calc.py

dftd4.json

dftd4.txt

eb08f3a9-0af6-4a04-a

eb08f3a9-0af6-4a04-a

example.sh

freq.log

gaussian.com

model.json

model\_predict\_kwargs

## opt\_freq\_model

## uvvis

1 714

2

3 C

4 C

5 C

6 H

7 C

8 H

9 C

10 H

11 C

12 C

13 H

14 C

15 H

16 H

17 H

18 C

19 H

20 H

21 H

22 C

23 H

24 C

25 H

26 H

27 H

28 C

29 H

30 H

31 C

2.2112054708775

2.3333751994675

2.2640290150525

2.3412011859445

2.0977040351767

2.0408275352385

1.9896957011252

1.8504990036569

2.0390377502618

1.9102512919706

1.9246634197864

3.1070721463076

4.0340826493530

3.1742041911812

3.0367451126373

0.5645362113050

-0.2543357346174

0.4639530906585

0.4609182286033

2.5132061188163

2.9952651007218

1.1610324792186

0.5519540104443

0.5985391790111

1.3121438634266

3.4370192949536

4.2740641529375

3.3528097176930

3.5424696493984

3.5197793043290

4.7066975749135

5.9136671677824

6.8389174713286

5.9309931689067

6.8729871278169

4.7352491104490

4.7503980176509

3.5099159499065

2.2048200652166

1.3829144679854

1.9816909817246

1.9950606698427

2.7558987735340

1.0130664111894

2.1333103852210

2.2048710966892

1.1884602005229

2.9544875537983

4.6806333070713

3.7144102646041

4.6573806039098

3.8158594339043

5.5701700636147

4.5846008946392

5.7059312080626

6.0444844539798

6.0296102641114

1.6256248254038

-2.6493395676902

-3.3700576203373

-2.6737718246351

-3.2300126972531

-1.2932445696169

-0.7660259784736

-0.5900502224319

0.4860094406428

-1.2586865861668

-0.4942902744449

-1.2130082792365

0.4331621671269

-0.1458940028188

1.2004980915698

0.9354549480634

0.2300612321852

-0.4928895210576

0.7723740268585

0.9396898449164

-4.8845438326538

-5.0947189136245

-5.6201528065277

-5.2880360832700

-5.4170329076792

-6.6971127200995

-5.4082601637649

-4.8094897315349

-6.4397312829619

-3.3404468227055

## TERMINAL

## OUTPUT

(base) dral@login:~\$



Jupyter Lab

No Jupyter instances are currently running.

Start a new instance by launching a new job.

\* CPU Number: 1

\* GPU Number: 0

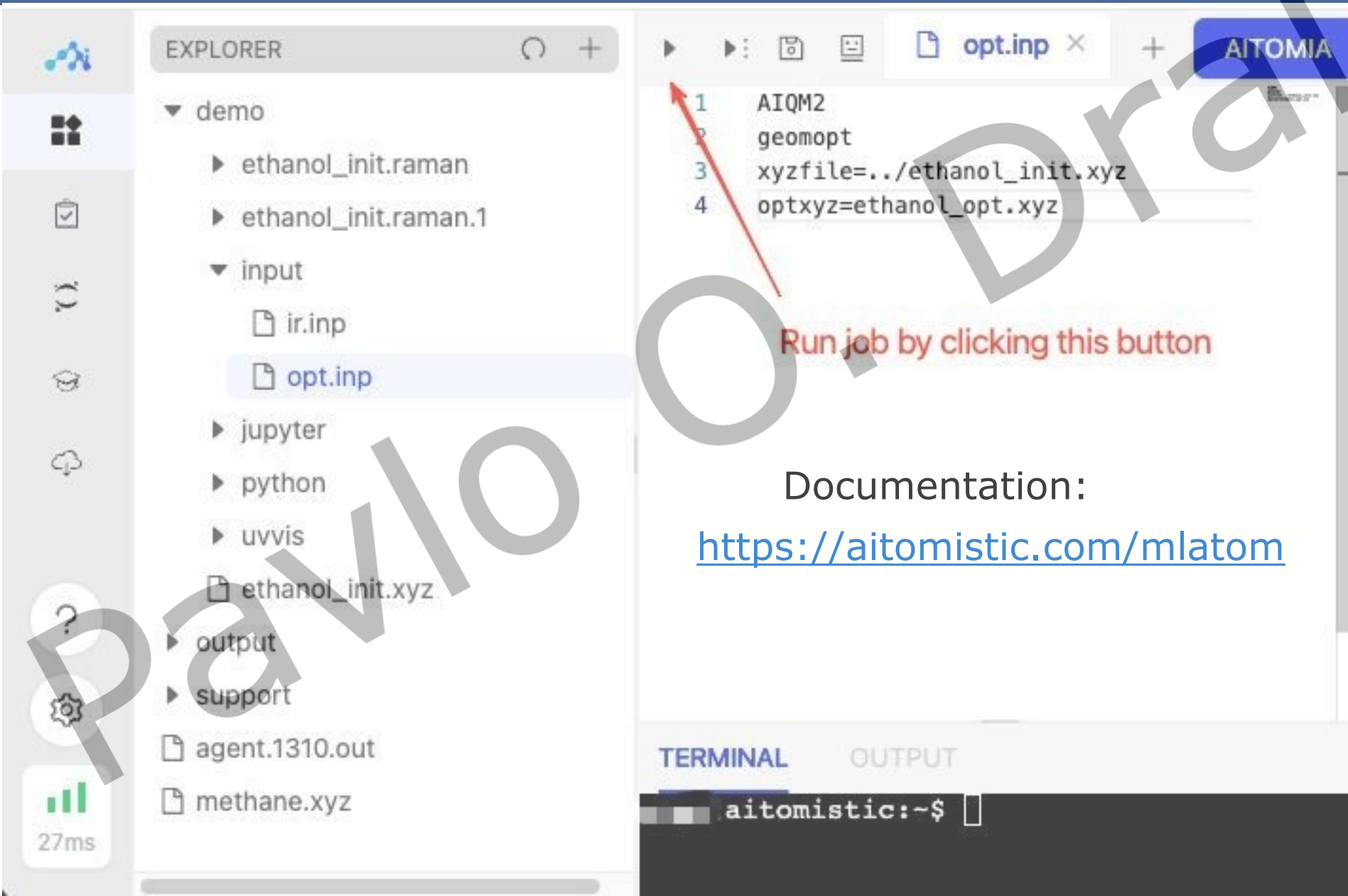
Partition: Default

Node: Select a machine

Launch



89ms



The screenshot displays the MLatom web interface. On the left is the 'EXPLORER' panel showing a file tree with folders like 'demo', 'input', 'output', and 'support', and files like 'ethanol\_init.raman', 'ir.inp', 'opt.inp', 'ethanol\_init.xyz', 'agent.1310.out', and 'methane.xyz'. The 'opt.inp' file is selected. The main panel shows the content of 'opt.inp' with the following text:

```
1 AIQM2
2 geomopt
3 xyzfile=../ethanol_init.xyz
4 optxyz=ethanol_opt.xyz
```

A red arrow points to the first line of the file. Below the editor, there is a 'Run job by clicking this button' text. At the bottom, the 'TERMINAL' panel shows the prompt 'aitomistic:~\$'.

Documentation:

<https://aitomistic.com/mlatom>



EXPLORER

- course
- demo
  - input
  - jupyter
  - python
  - uvvis
  - ethanol.com
  - ethanol\_init.xyz
- demo2
- h2\_test
- test\_kreg
- tests
- Gau-78383.inp

ethanol\_init.xyz ethanol.com

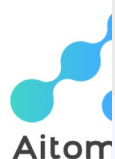
```

~ > demo > ethanol.com
1 #P b3lyp/6-31G* opt
2
3 ethanol
4
5 0 1
6 C -3.902027020 -1.283783760 0.000000000
7 H -3.545372590 -2.292593770 0.000000000
8 H -3.545354180 -0.779385570 -0.873651500
9 H -4.972027020 -1.283770580 0.000000000
10 C -3.388684800 -0.557827490 1.257404970
11 H -3.418258266 0.351059004 0.693533749
12 H -3.412790015 -1.452244073 1.844208100
13 O -2.838512436 0.303845702 2.257269797
14 H -2.541978830 0.592515666 3.123490027
15
16
  
```

Submit with more options

Submit directly (1 CPU)

104ms



Job Name:

Start Time:

End Time:

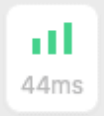
State:

Search



Job list

| Job Name         | CPU Cores | Partition | Node List | Time Start          | Time End            | State    | Actions |
|------------------|-----------|-----------|-----------|---------------------|---------------------|----------|---------|
| jupyter_slurm.sh | 1         | cpu       | b006      | 2025-08-22 10:36:52 | 2025-08-22 18:37:18 | TIMEOUT  |         |
| jupyter_slurm.sh | 1         | cpu       | b006      | 2025-08-21 20:10:31 | 2025-08-22 04:10:47 | TIMEOUT  |         |
| opt.py           | 1         | cpu       | b006      | 2025-08-19 11:16:21 | 2025-08-19 11:16:34 | FAILED   |         |
| opt.py           | 1         | cpu       | b006      | 2025-08-19 11:10:24 | 2025-08-19 11:11:07 | FAILED   |         |
| jupyter_slurm.sh | 1         | cpu       | b006      | 2025-08-14 17:33:23 | 2025-08-15 01:33:32 | TIMEOUT  |         |
| jupyter_slurm.sh | 1         | cpu       | b002      | 2025-08-07 14:37:20 | 2025-08-07 22:37:45 | TIMEOUT  |         |
| jupyter_slurm.sh | 1         | cpu       | b006      | 2025-08-05 10:28:51 | 2025-08-05 18:29:10 | TIMEOUT  |         |
| ir.inp           | 1         | cpu       | b002      | 2025-07-24 22:16:59 | 2025-07-24 22:17:35 | COMPLETE |         |
| mol1.inp         | 1         | cpu       | b002      | 2025-07-24 22:16:42 | 2025-07-24 22:22:09 | COMPLETE |         |
| ir.inp           | 1         | cpu       | b004      | 2025-07-21 19:48:31 | 2025-07-21 19:49:02 | COMPLETE |         |



**EXPLORER**

- ▶ demo
- ▼ promotion
  - ▼ calc
  - ▼ chat
  - ▼ reaction
  - ▼ xyz
    - butadiene.xyz
- ▶ tests
- ▶ tutorials
- ▶ umlip

**Protomia**

Hello, how can I help you?

Drop files or folders here

butadiene.xyz

Press Enter to send message

**Protomia**

Hello, how can I help you?

opt **opt + freq** freq IR Raman TSopt

UV/Vis

butadiene.xyz

Do geometry optimization and frequency analysis for butadiene.xyz using the AIQM2 method.

B3LYP/6-31G\*

**AIQM2**

Do geometry optimization and frequency analysis for butadiene.xyz using the wb97x/def2-svp method.

**Editable block**

**EXPLORER**

- ▶ demo
- ▼ promotion
  - ▼ calc
  - ▼ chat
  - ▼ reaction
- ▼ xyz
  - butadiene.xyz
- ▶ tests
- ▶ tutorials
- ▶ umlip

**Protomia**

Hello, how can I help you?

**Drop multiple folders here**

3 items

**Working directory**

✕ calc ✕ chat ✕ reaction

**Pick one working directory**

29ms



EXPLORER

- demo
- promotion
  - calc
  - chat
  - reaction
- xyz
  - butadiene\_opt\_AIQM2
  - butadiene\_opt\_freq\_AIQM2
    - scr
    - butadiene\_opt\_freq.inp
    - butadiene\_opt\_freq.log
    - butadiene.json
    - butadiene.xyz
  - tests
  - tutorials
  - umlip

28ms

Protomia

Hello, how can I help you?

**Drop calculation files here**

butadiene\_opt\_freq.log

analyze this log file

**Ask for analysis**

Protomia

RECENT ACTIVITY

Interpreting instructions

1 item(s)

I've analyzed your log file — this was a successful vibrational frequency and thermochemistry calculation for butadiene using the AIQM2 method. The geometry was read from [butadiene\\_opt\\_freq.xyz](#), and all 24 vibrational frequencies are real and positive, confirming you've found a true energy minimum (no imaginary modes). That's great — it means your structure is stable.

Key results at 298 K:

Press Enter to send message

## Ask about AIQM2

what is AIQM2 and how to use it?



AIQM2 is the second-generation AI-enhanced quantum mechanics method designed for high-speed, chemically accurate simulations of organic reactions — including transition state searches, geometry optimizations, frequency calculations, and reactive dynamics. It combines machine learning with semi-empirical quantum mechanics (using GFN2-xTB as its baseline) to deliver accuracy competitive with DFT at a fraction of the computational cost. AIQM2 also supports property predictions like dipole moments for IR spectrum generation.

To use AIQM2 in MLatom, input file or Python script.

### Introduction

Example 1: Input File for Single Point Energy + Geometry Optimization

```
AIQM2
xyzfile=ts.xyz
geomopt
ts=True
optxyz=ts_opt.xyz
```

### Example usage

+ promotion

Press Enter to send message



### Example 2: Frequency Calculation & IR Spectrum (Input File)

After geometry optimization, calculate frequencies and generate IR spectrum:

```
ir
AIQM2
xyzfile=opt.xyz
```

### Example usage

Frequencies are automatically scaled by 0.962 for better agreement with experiment.

'acetone', etc.).

Sources:

- [http://mlatom.com/docs/tutorial\\_aiqm2.rst.html](http://mlatom.com/docs/tutorial_aiqm2.rst.html) (tutorial\_aiqm2.rst)
- [http://mlatom.com/docs/tutorial\\_aiqm2.html](http://mlatom.com/docs/tutorial_aiqm2.html) (Explore Diels-Alder reactions with AIQM2)
- [http://mlatom.com/docs/tutorial\\_ts.html](http://mlatom.com/docs/tutorial_ts.html) (Choosing the method or model)
- [http://mlatom.com/docs/tutorial\\_ts.rst.html](http://mlatom.com/docs/tutorial_ts.rst.html) (tutorial\_ts.rst)
- [http://mlatom.com/docs/tutorial\\_uaiqm.rst.html](http://mlatom.com/docs/tutorial_uaiqm.rst.html) (tutorial\_uaiqm.rst)

### Sources and further reading

+ promotion

Press Enter to send message



**Protomia**

Chat History **Check chat history**

what is AIQM2 and how to use it  
3 messages 2026-03-19 16:52:33

analyze this log file  
3 messages 2026-03-17 21:32:27

analyze this log file  
3 messages 2026-03-17 21:29:26

Do geometry optimization and frequency analysis f...  
20 messages 2026-03-17 21:23:29

what's the reaction energy for DA reaction between ...  
28 messages 2026-03-11 10:04:13

for Diels-Alder reaction between ethylene and 1,3-b...  
32 messages 2026-03-11 09:55:42

Do geometry optimization and frequency analysis f...  
16 messages 2026-03-11 09:42:34

**Restart from previous chat**

+ promotion

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34ms

**Protomia**

what is AIQM2 and how to use it?

**RECENT ACTIVITY**  
Interpreting instructions  
1 item(s)

AIQM2 is the second-generation AI-enhanced quantum mechanics method designed for high-speed, chemically accurate simulations of organic reactions — including transition state searches, geometry optimizations, frequency calculations, and reactive dynamics. It combines machine learning with semi-

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**Protomia**

File name: what\_is\_AIQM2\_and\_how\_to\_use\_it.md

Select a folder to save the conversation:

Current Selected Path:  
/aitomistic/hub/home/chenyxx/promotion/chat

- ▶ demo
- ▼ promotion
  - ▶ calc
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**Protomia**

what\_is\_AIQM2\_and\_how\_to\_use\_it.md

Word ... > promotion > chat > what\_is\_AIQM2\_and\_how\_to\_use\_it.md

Wrap

```
1 # what is AIQM2 and how to use it
2
3 **Export Date:** 2026/3/20 15:02:30
4
5 ---
6
7 ## User
8
9 what is AIQM2 and how to use it?
10
11
12 ---
13
14 ## Assistant
15
16 Interpreting instructions
17
18
19 ---
20
21 ## Assistant
22
23 AIQM2 is the second-generation AI-enhanced quantum mechanics method designed for high-speed, chemically accurate simulations of organic reactions – including transition state searches, geometry optimizations, frequency calculations, and reactive dynamics. It combines machine learning with semi-empirical quantum mechanics (using GFN2-xTB as its baseline) to deliver accuracy competitive with DFT at a fraction of the computational
```



|                   |                                                                                     |
|-------------------|-------------------------------------------------------------------------------------|
| 25 Apr 2026 20:56 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 18:20 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 18:14 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 17:21 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 15:20 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 15:17 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 15:14 |  |
| 25 Apr 2026 14:19 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 14:19 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 12:38 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 09:44 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 09:31 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 09:31 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 07:49 | Pavlo O. Dral                                                                       |
| 25 Apr 2026 07:47 | Protomia (Pavlo O. Dral)                                                            |
| 25 Apr 2026 00:02 | Protomia (Pavlo O. Dral)                                                            |
| 24 Apr 2026 23:49 | Protomia (Pavlo O. Dral)                                                            |
| 24 Apr 2026 23:36 | Protomia (Pavlo O. Dral)                                                            |

Show me the electrostatic-potential (ESP) map for water using HF/STO-3G via PySCF — render the 3D ESP-coloured surface as a .png.

 [ESP\\_PYSCF\\_164807](#)

 copy

>  Ran 2 actions

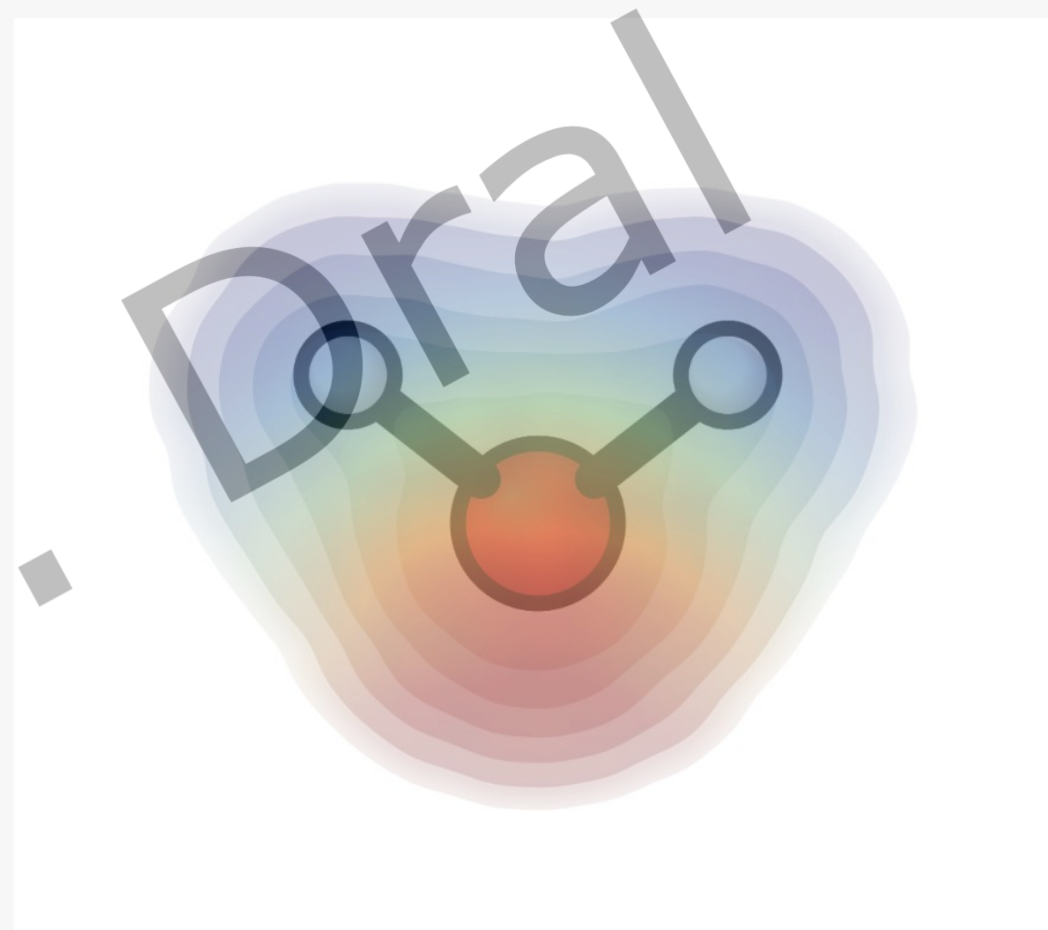
I'll run the HF/STO-3G ESP calculation for water, which will also render the ESP-coloured surface automatically.

>  Ran 1 action

The HF/STO-3G single-point for water has been submitted to the compute cluster (job 183770) — from the resulting wavefunction, the ESP grid will be generated and the ESP-coloured surface rendered to a PNG automatically.

It's a tiny system so it should be quick. I'll report back with the rendered surface and the energy once it finishes — feel free to keep chatting in the meantime.

The HF/STO-3G calculation for water is done — here's the ESP map rendered from the resulting wavefunction:



### Key results (HF/STO-3G, ground state):

- **Total electronic energy: -74.964449 Hartree**
- HOMO: -10.637 eV
- LUMO: 16.190 eV
- HOMO-LUMO gap: 26.827 eV

Show me the HOMO surface picture (3D rendered orbital, .png) for water using HF/STO-3G via PySCF.

 [HOMOLUMO\\_VIZ\\_164807](#)

 copy

>  Ran 2 actions

I'll run an HF/STO-3G single-point on water to generate wavefunctions and render the HOMO orbital surface automatically.

>  Ran 1 action

The water HOMO visualization job has been submitted to the cluster (Slurm job 183766 ) — it's running an HF/STO-3G single-point calculation, after which the HOMO (and LUMO) orbital surface is rendered to a PNG automatically.

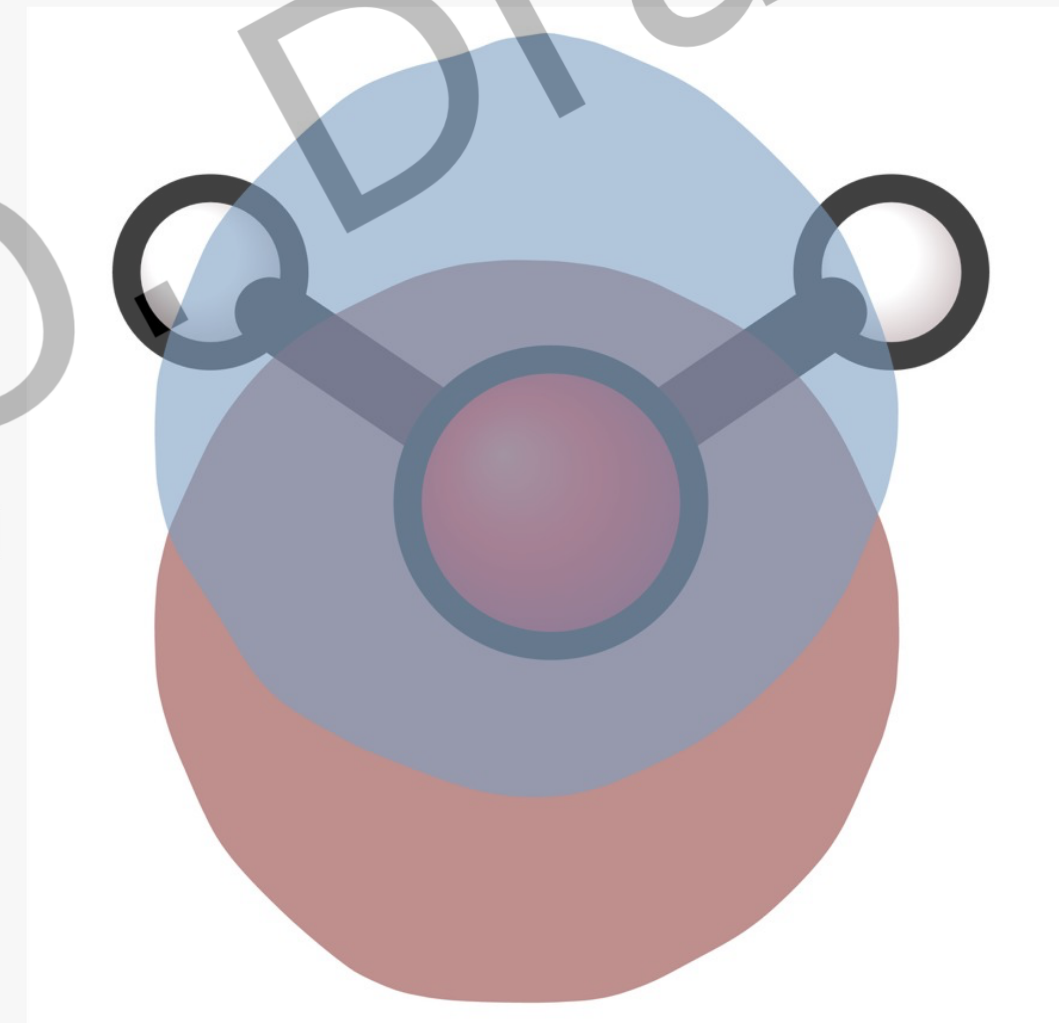
As soon as it finishes — typically a couple of minutes for a small molecule like this — I'll hand back the rendered HOMO surface picture right here in chat. Feel free to keep chatting in the meantime; I'll report the result the moment it's ready.

The water calculation finished — here's your HOMO surface picture!



Below is the 3D-rendered HOMO isosurface from the HF/STO-3G calculation on water (and the LUMO as a bonus). You can see the

Below is the 3D-rendered HOMO isosurface from the HF/STO-3G calculation on water (and the LUMO as a bonus). You can see the classic lone-pair-shaped orbital on the oxygen, which is exactly what you expect as the HOMO of H<sub>2</sub>O.



Compute the UV-Vis absorption spectrum of triphenylamine: optimize the geometry with AIQM3, then run OMNI-P2x for the excited states.

📁 OMNIP2X\_164807

> ✓ Ran 2 actions

I'll set up a multi-stage calculation: AIQM3 geometry optimization first, then an OMNI-P2x excited-state (UV-Vis) single point on the optimized geometry, with everything saved to your working folder.

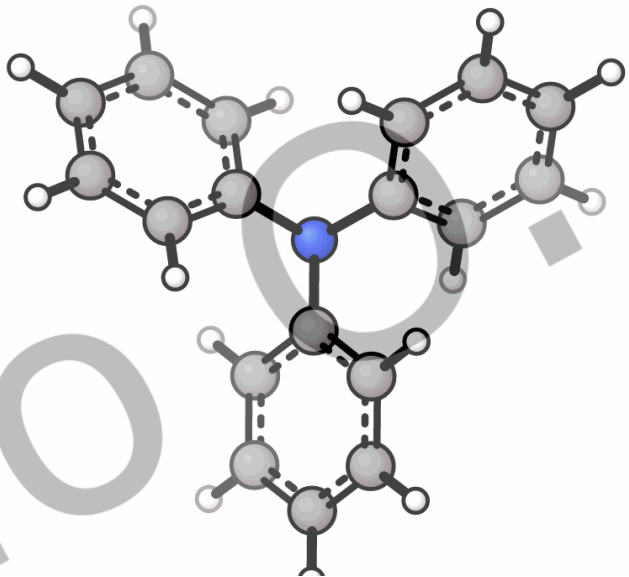
> ✓ Ran 1 action

Your calculation is submitted and running on the cluster. Here's the plan that's queued:

- 1. **Stage 1 — Geometry optimization** of triphenylamine with **AIQM3**
- 2. **Stage 2 — Excited states (UV-Vis)** with **OMNI-P2x** on the optimized geometry

Files in your working folder:

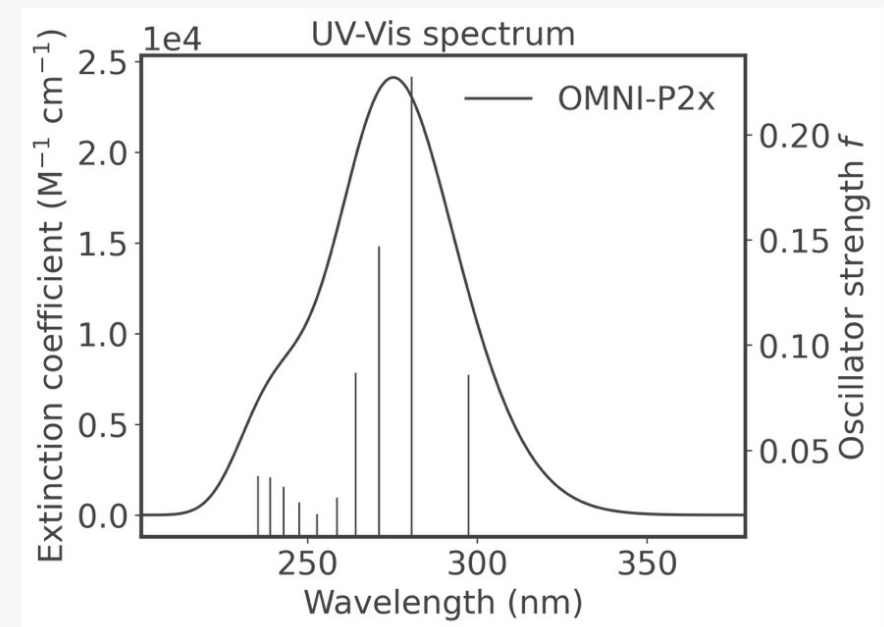
- Result bundle: [triphenylamine\\_opt\\_AIQM3\\_9040fbb9](#)
- Optimized geometry: [triphenylamine\\_opt\\_AIQM3.xyz](#)
- Structure rotation:



Protomia Solution 2.8.2

|                |        |       |       |
|----------------|--------|-------|-------|
| S <sub>2</sub> | 4.4181 | 280.6 | 0.227 |
| S <sub>3</sub> | 4.5736 | 271.1 | 0.147 |
| S <sub>4</sub> | 4.6934 | 264.2 | 0.087 |
| S <sub>5</sub> | 4.7945 | 258.6 | 0.027 |

Spectrum:





Charge Mult

0 1



## Protomia Solution 2.6.0

keep chatting in the meantime!

The NEB search and transition-state refinement have finished! Here are the key results for the ammonia umbrella-inversion TS:

### Transition-state geometry (AIQM3 optimized):

- Total electronic energy:  $-56.4985$  Hartree
- The TS is confirmed with exactly one imaginary frequency:  $-871.1 \text{ cm}^{-1}$
- Forward barrier from reactant:  $6.1 \text{ kcal/mol}$
- Dipole moment:  $0.0 \text{ D}$  (planar  $D_{3h}$  symmetry at the saddle point)

Optimized TS structure: [NH3\\_umbrella\\_inversion\\_tsopt\\_AIQM3.xyz](#)

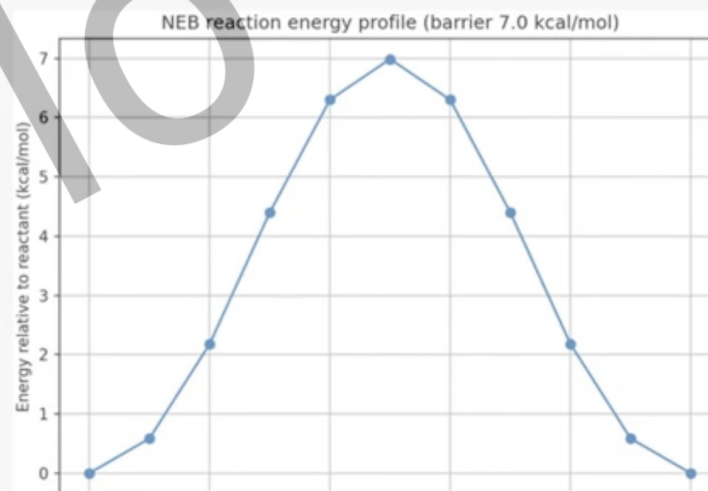
### TS Verification:

Imaginary frequencies: 1 ( $-871.1 \text{ cm}^{-1}$ ) ✓

Normal mode match: umbrella inversion (H atoms move through the N plane) ✓

Verdict: VALID TS

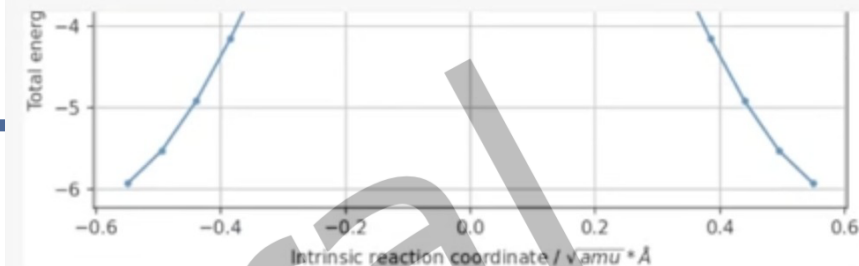
### NEB energy profile (reactant → TS → product):



+ NH3\_umbrella\_inversion\_tsop...

Press Enter to send message

## Protomia Solution 2.6.0



The IRC confirms that the transition state connects the two pyramidal minima (reactant and product) through the planar  $D_{3h}$  saddle point, exactly as expected for the ammonia umbrella inversion.

### IRC trajectory files:

- Full path (both directions): [NH3\\_umbrella\\_inversion IRC\\_full.xyz](#)
- Backward (toward reactant): [backward.xyz](#)
- Forward (toward product): [forward.xyz](#)

Full IRC bundle: [NH3\\_umbrella\\_inversion IRC AIQM3 bundle](#)

### Summary of the complete characterization:

| Property            | Value                                       |
|---------------------|---------------------------------------------|
| TS energy           | $-56.4985$ Hartree                          |
| Imaginary frequency | $-871.1 \text{ cm}^{-1}$                    |
| Forward barrier     | $6.1 \text{ kcal/mol}$                      |
| TS symmetry         | Planar $D_{3h}$ (dipole = $0.0 \text{ D}$ ) |
| IRC connectivity    | Reactant ↔ TS ↔ Product ✓                   |

The umbrella inversion of  $\text{NH}_3$  is now fully characterized: the NEB located the planar transition state, the frequency calculation confirmed it is a first-order saddle point with the correct umbrella-inversion normal mode, and the IRC verified that it smoothly connects the two equivalent pyramidal minima.

copy

+ NH3\_umbrella\_inversion IRC...

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