



**Lecture: ML potentials I**

**Pavlo O. Dral**  
Xiamen University, P.R. China

Visiting Professor in  
Nicolaus Copernicus University, Poland

10 September 2024

Please check the link  
[and register on XACS cloud – optional]



**before lunch break**



Questionary

# Pavlo O. Dral

## AI in computational chemistry



Professor | **Outstanding Youth (Overseas)**

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

Research Areas: **artificial intelligence, quantum chemistry, dynamics, excited states, semi-empirical methods**

2024-Present: Nicolaus Copernicus University, Visiting Professor

2021-Present: Xiamen University, Full Professor

2019-2021: Xiamen University, Associate Professor

2013-2019: Max-Planck-Institut für Kohlenforschung, Postdoc

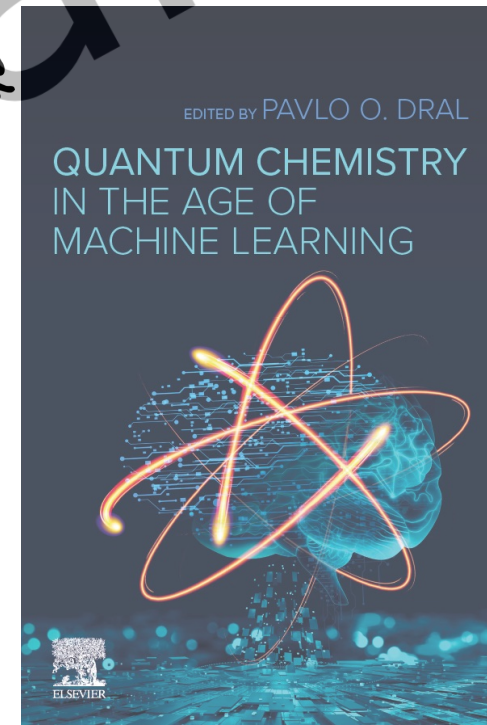
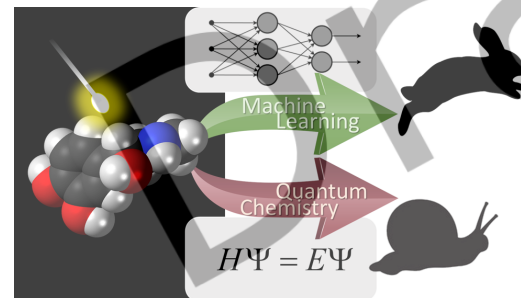
2010-2013: University of Erlangen-Nürnberg, M.Sc. & Ph.D.

2008-2010: University of Erlangen-Nürnberg, M.Sc.

2004-2010: National Technical University of Ukraine "KPI", B.Sc. & M.Sc.

### Research Interests:

Our research transforms chemical physics simulations by developing novel AI methods and providing software and cloud computing platforms.



### ✓ Selected papers:

- AI platform:** *J. Chem. Theory Comput.* **2024**, 20, 1193
- AI-quantum dynamics:** *Nat. Commun.* **2022**, 13, 1930
- AI-quantum mechanics:** *Nat. Commun.* **2021**, 12, 7022
- AI-excited states:** *Nat. Rev. Chem.* **2021**, 5, 388
- AI force fields:** *Chem. Sci.* **2021**, 12: 14396

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

# Looking for talented group members!



2023



**国家自然科学基金委员会**  
National Natural Science Foundation of China



**嘉庚创新实验室**  
TAN KAH KEE INNOVATION LABORATORY

2024





**国家自然科学基金委员会**  
National Natural Science Foundation of China



**嘉庚创新实验室**  
TAN KAH KEE INNOVATION LABORATORY

2023 – in Warsaw

2024 – in Xiamen



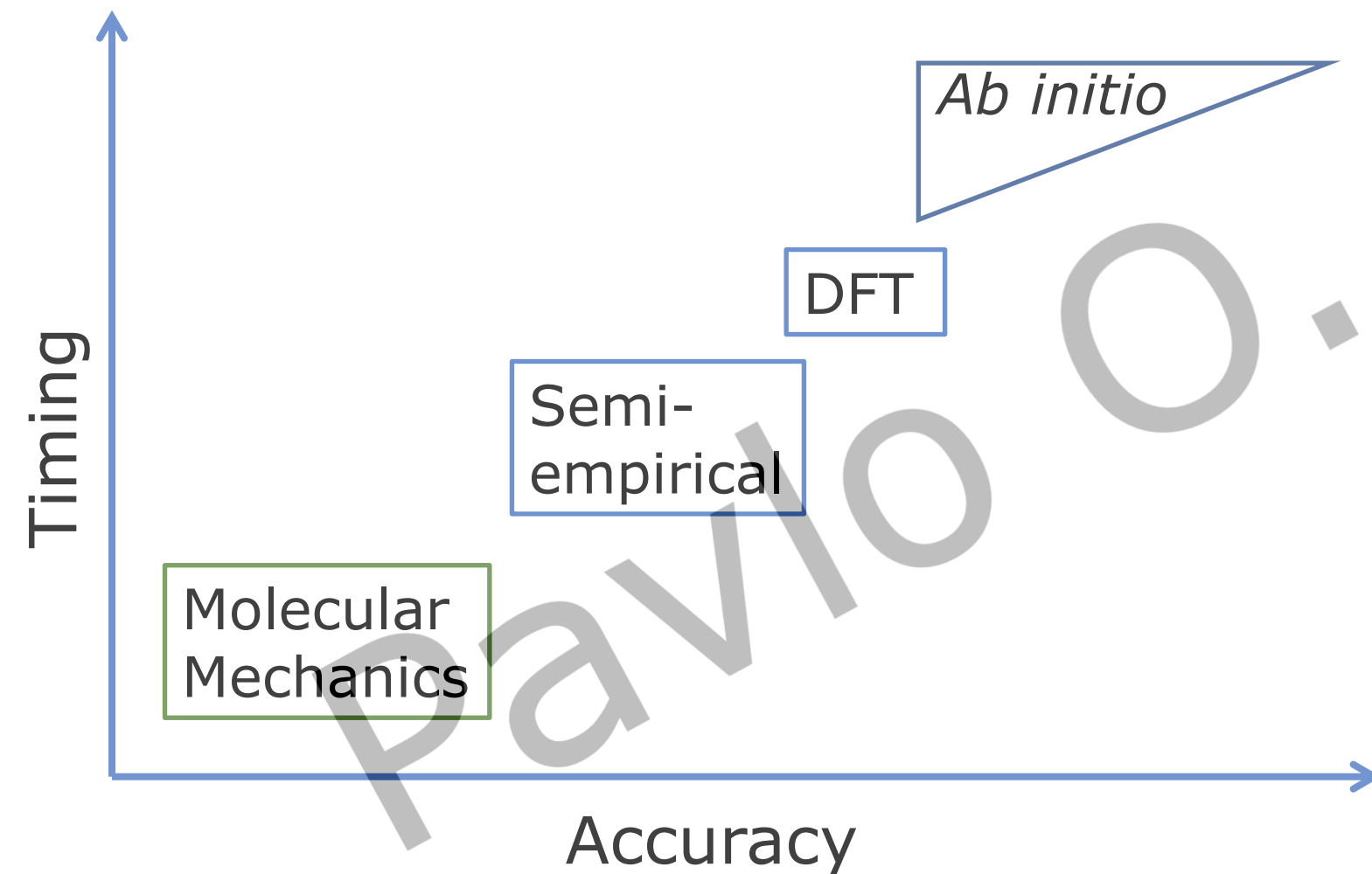


Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, *11*, 2336

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

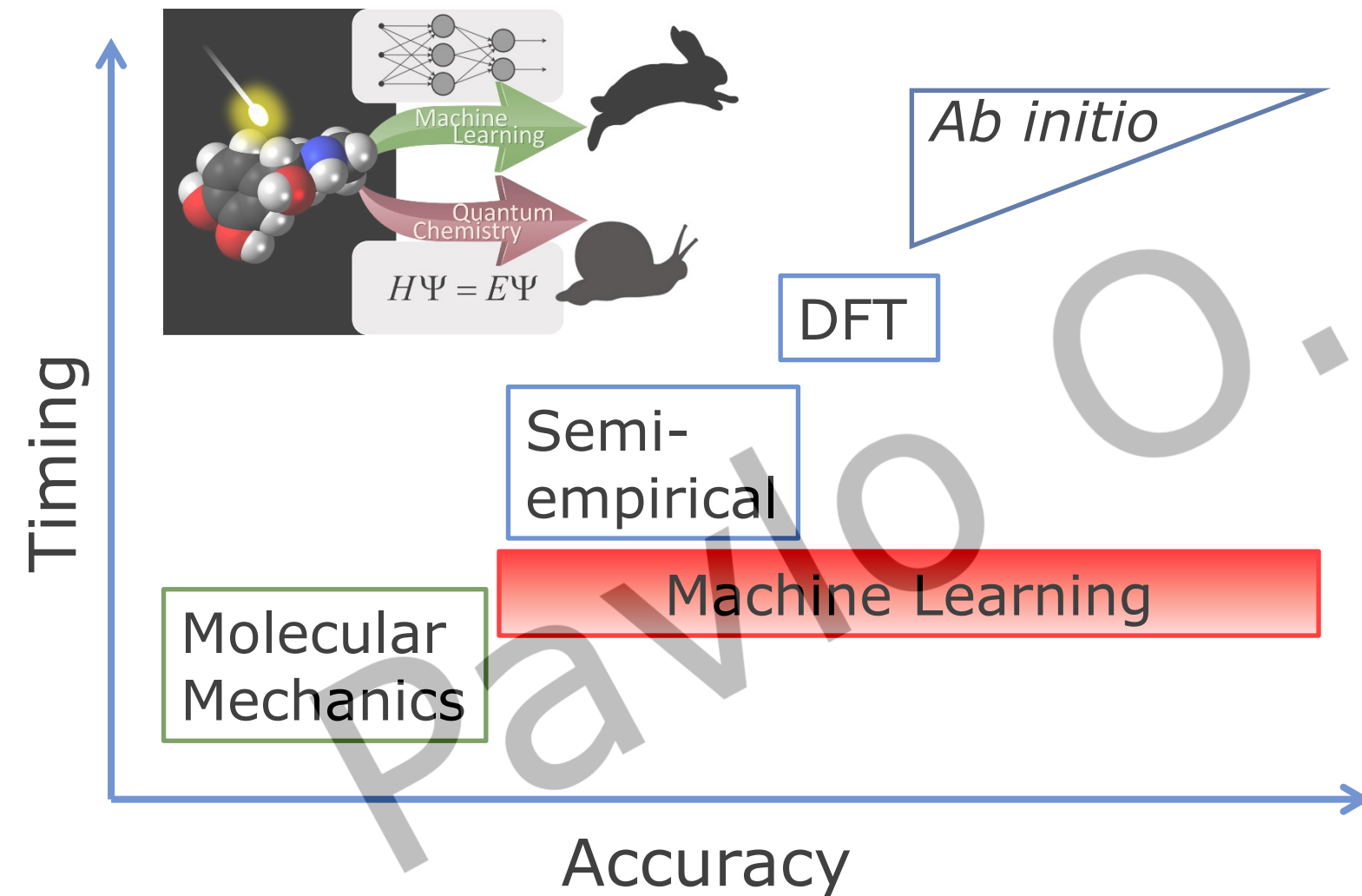
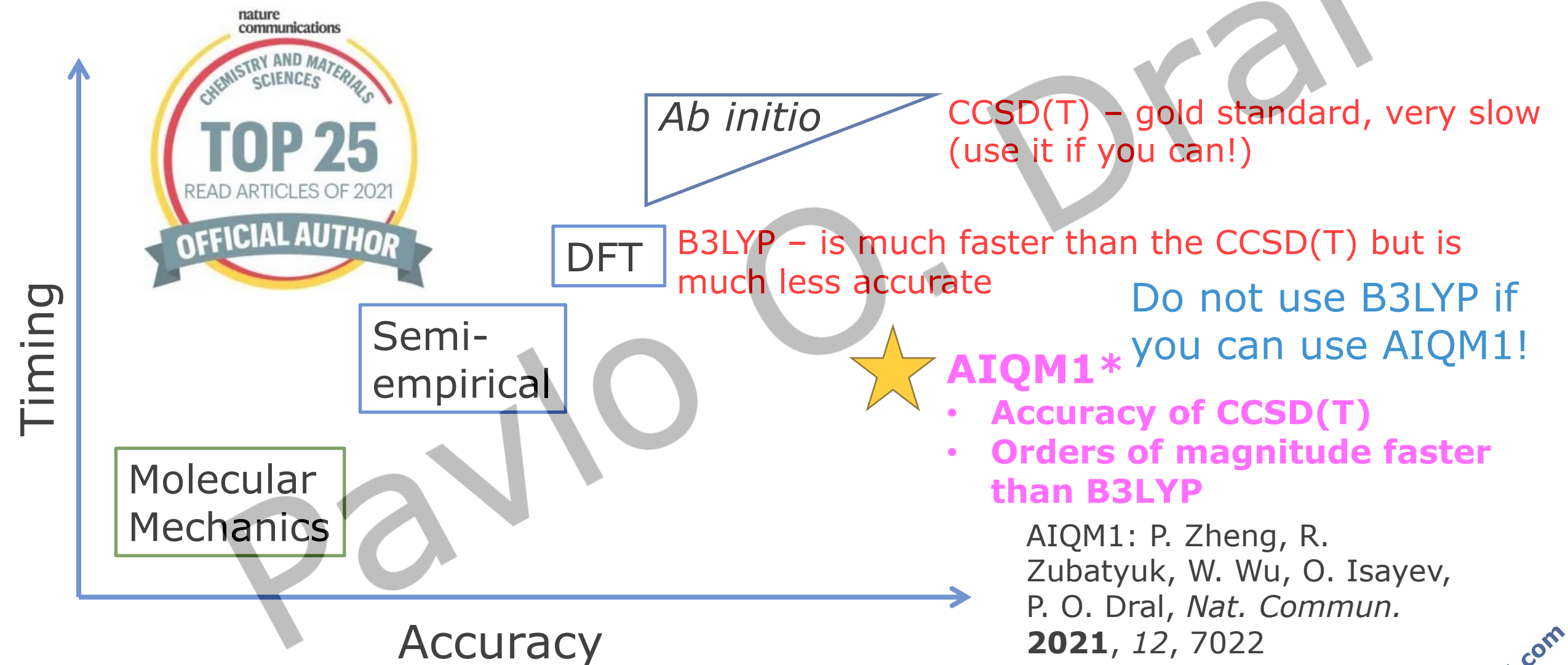


Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336



\*CHNO elements only – extensions on the way



P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

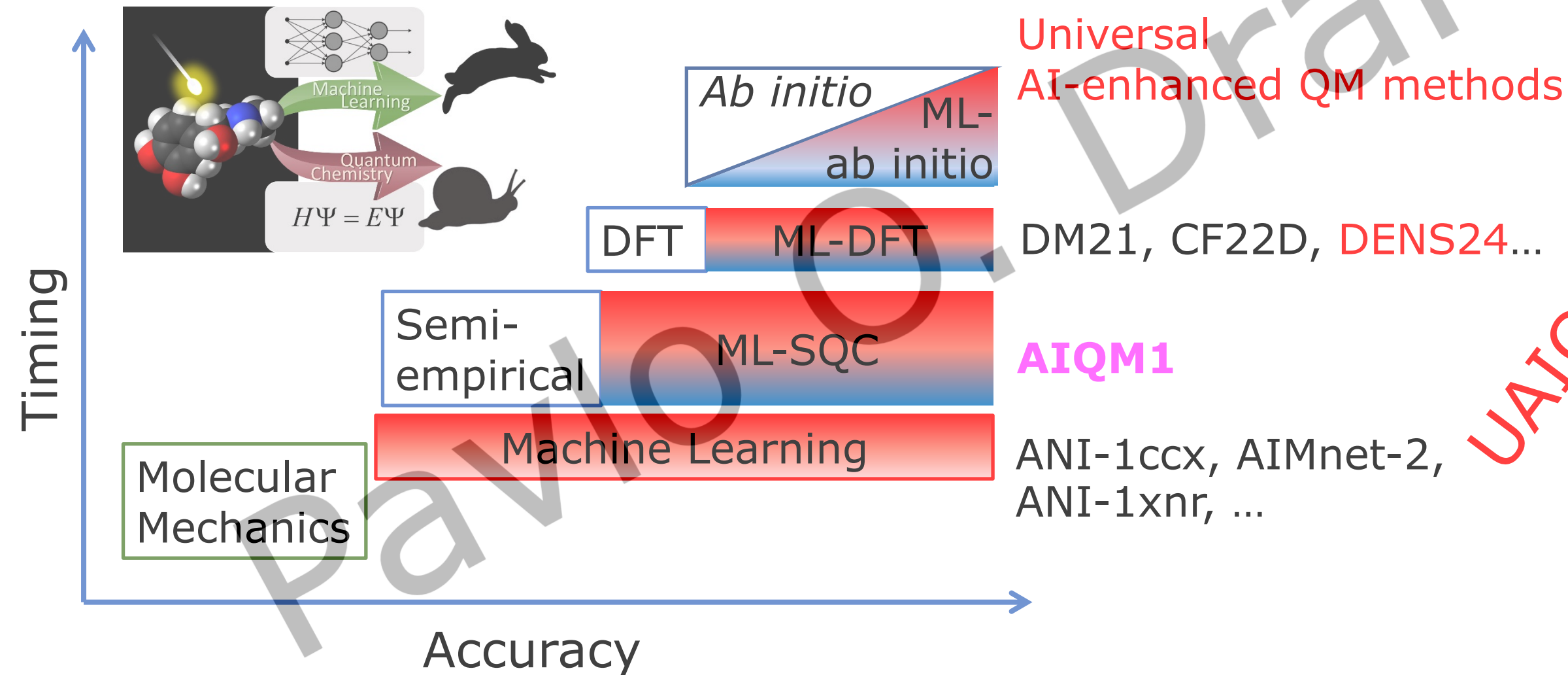
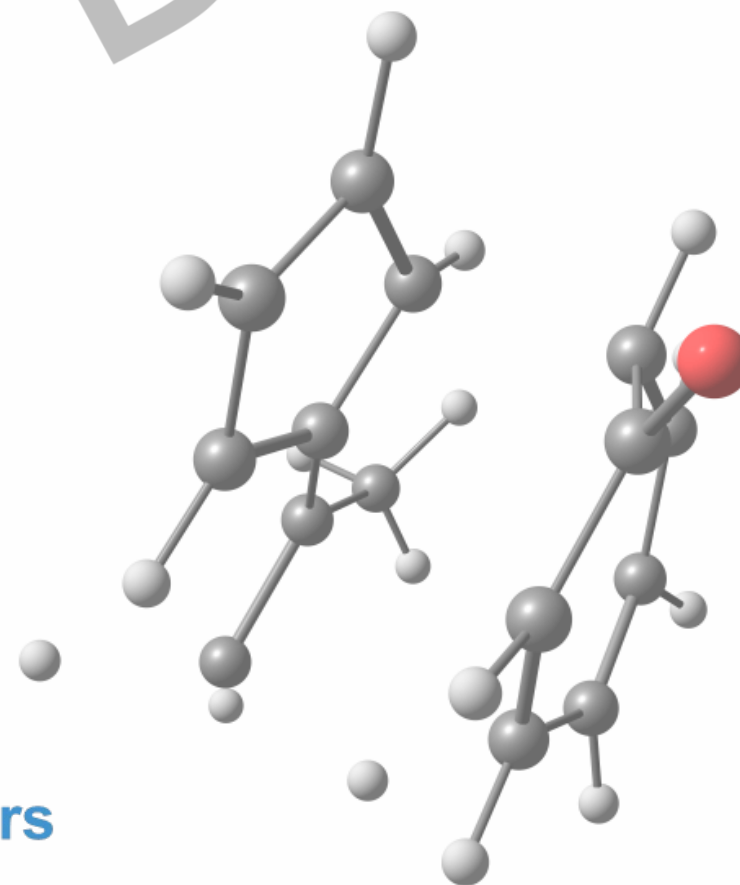
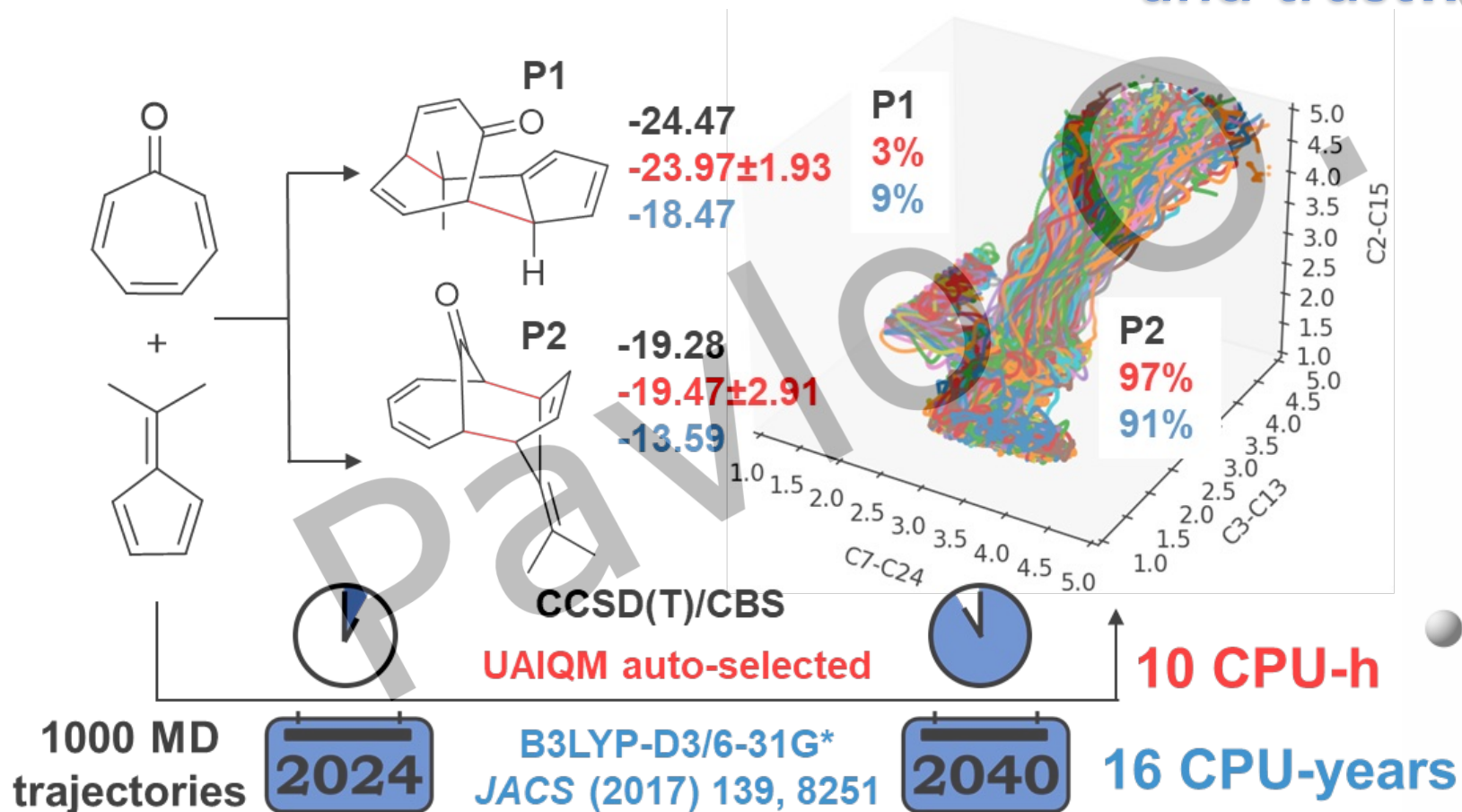


Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

*No reason to do non-ML computational chemistry!*

*AI methods: easy to use, also online*

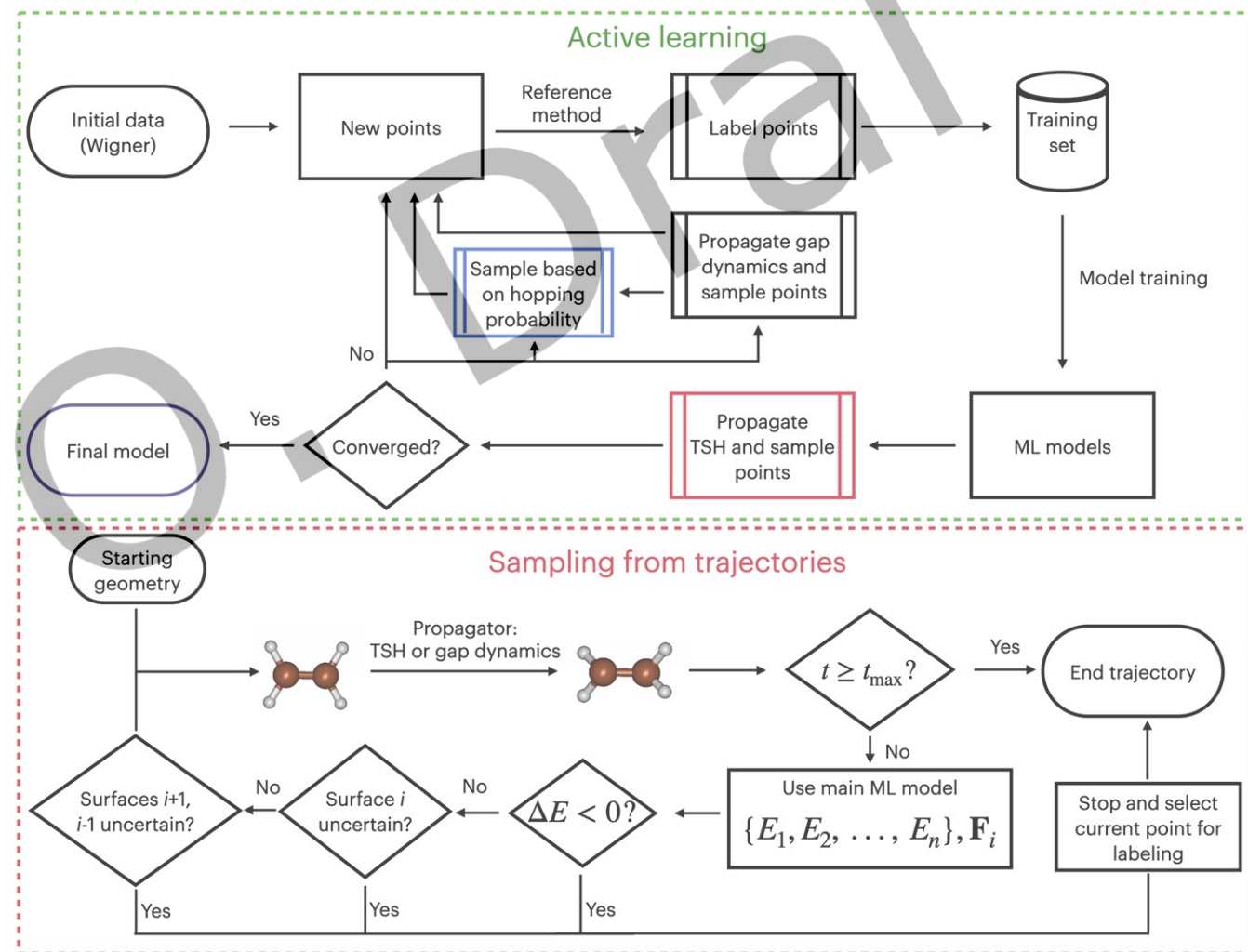
*AI methods are faster, more accurate, and trustworthy*



8 years of research!

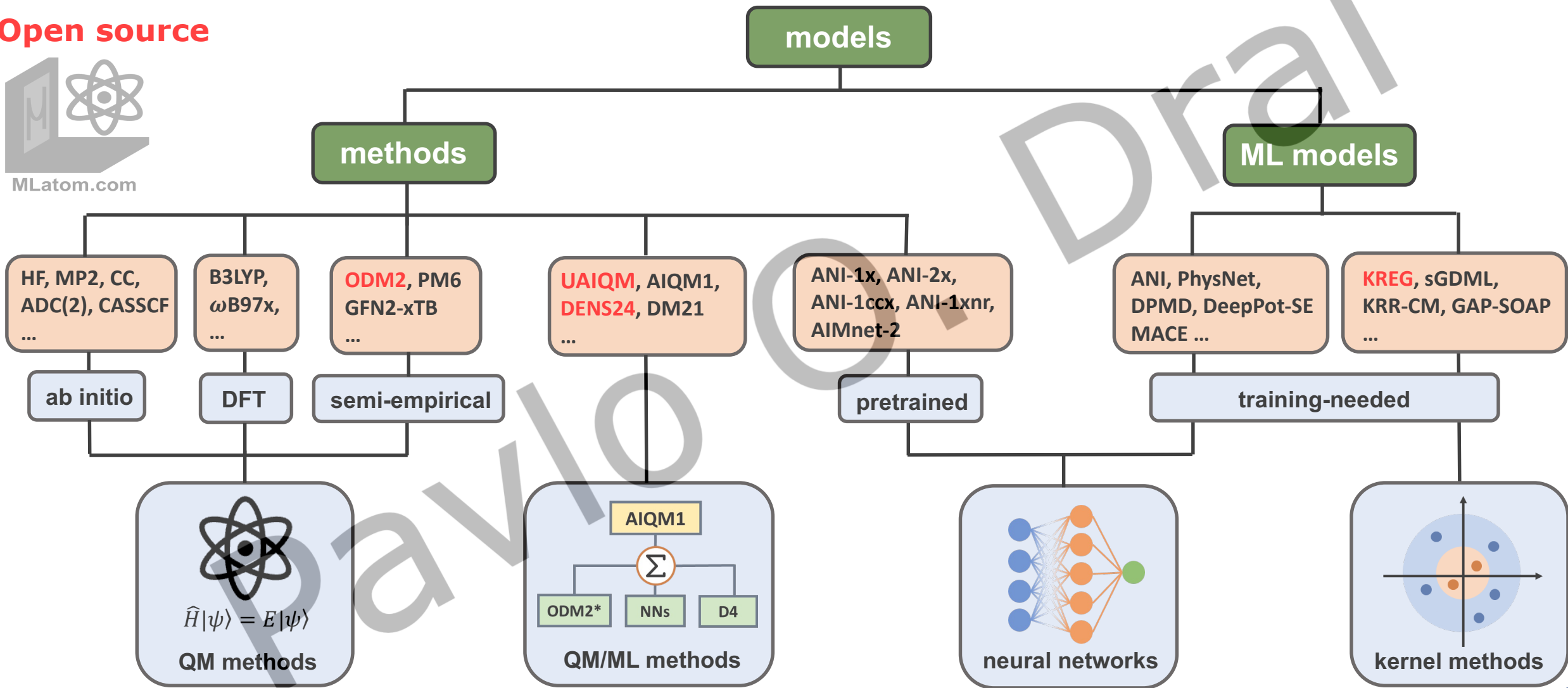
**Within days, can get precise ML-NAMD results on a single GPU and <16 CPU threads\***

\*from scratch for simple molecules



M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

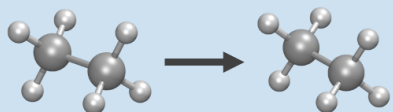
## Open source



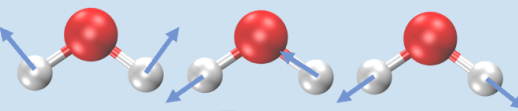
Single point calculations

Energies, forces, Hessian matrix...

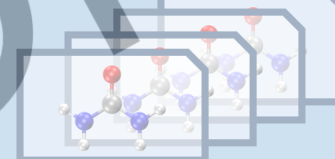
Geometry optimizations



Frequency calculations



Molecular dynamics

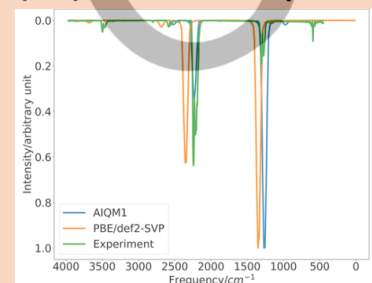


Thermochemistry  
calculations

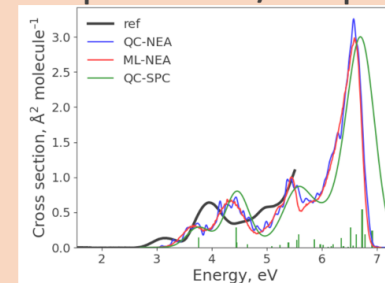
Heat of formation:

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

(Ro)vibrational spectra



One-photon UV/vis spectra



Properties &  
spectra

Simulations

Quantum dissipative  
dynamics

Two-photon  
absorption spectra

# Machine Learning



# Simulations




**MLatom**

# Quantum Chemistry


# Data

## Cloud Computing

 Job Submitter

 Terminal

 File Manager

 Job Manager

 Jupyter Lab

## Software

 Download

## Learning

 Courses

## Courses

### Testimonials:

*"Dr. Dral offers a fantastic introduction to the concepts around machine learning in chemistry!"*

For teachers: Your course can be here!

### 计算化学与人工智能

计算化学和人工智能迷你课程

作者: Pavlo O.Dral

发布日期: 2024年6月14日

Go

### Computational chemistry & AI

Hands-on course on computational chemistry and artificial intelligence (AI) by Pavlo O. Dral

Go

Recommend them further!

## Top review from the United States

 Patrick Faith

★★★★★ **Singularly Brilliant**

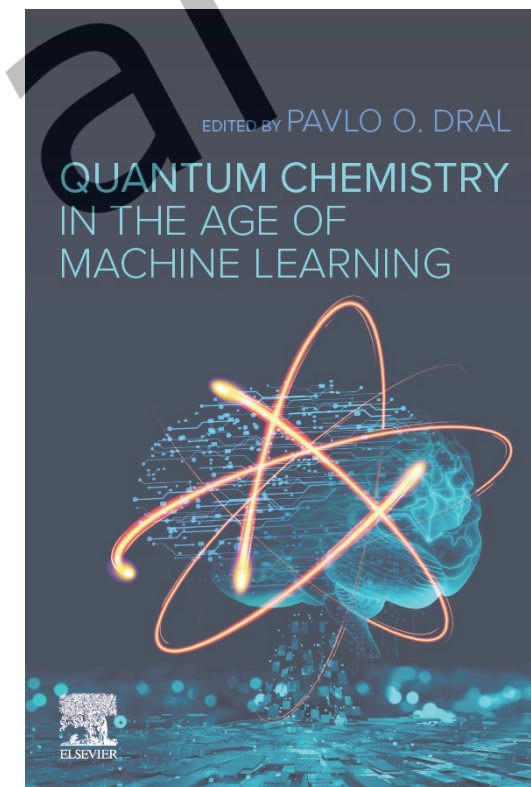
Reviewed in the United States on October 6, 2022

**Verified Purchase**

I bought this book really with no expectation, it's kind of expensive so was a bit of a stretch buy with really no reviews. Recieved it yesterday and quickly jumped through it, the last book I read that had this level of brilliance was Wheeler's Gravity, but **this book has much more of a team spirit reminding me of the early days of quantum mechanics.** Only modern book that it reminds me of is Wolframs recent "new kind of ..." but **this book is completely practical, team based, and I think a signifacly better path.**

I noticed a lot of research lead by the Xiamen area, connected to europe with a bit of usa work. Represents a major shift in in leadership of science, is a book every one should atleast attempt to glance through to "get" what is now going on.

27 chapters  
65 authors!





P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

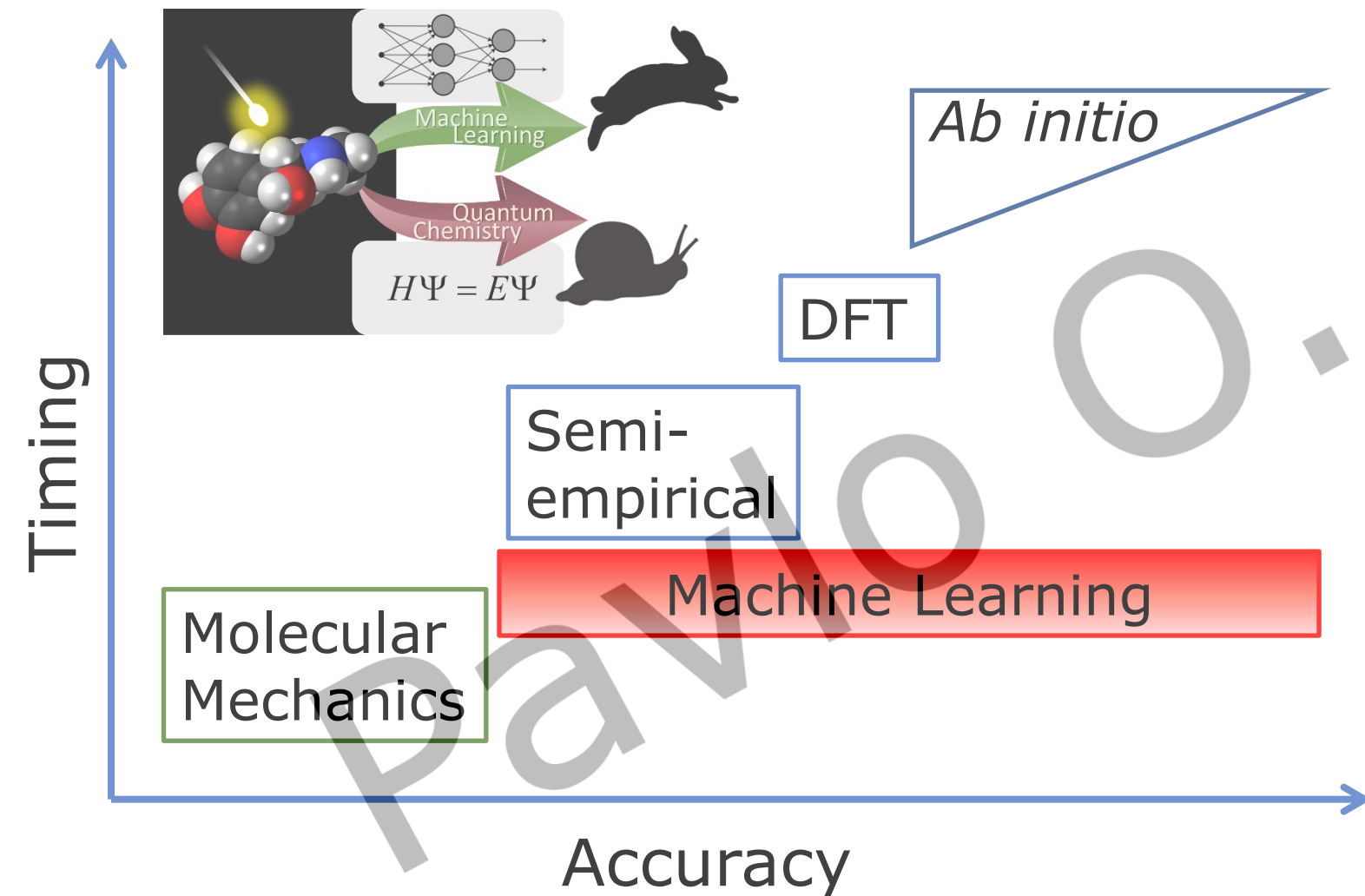


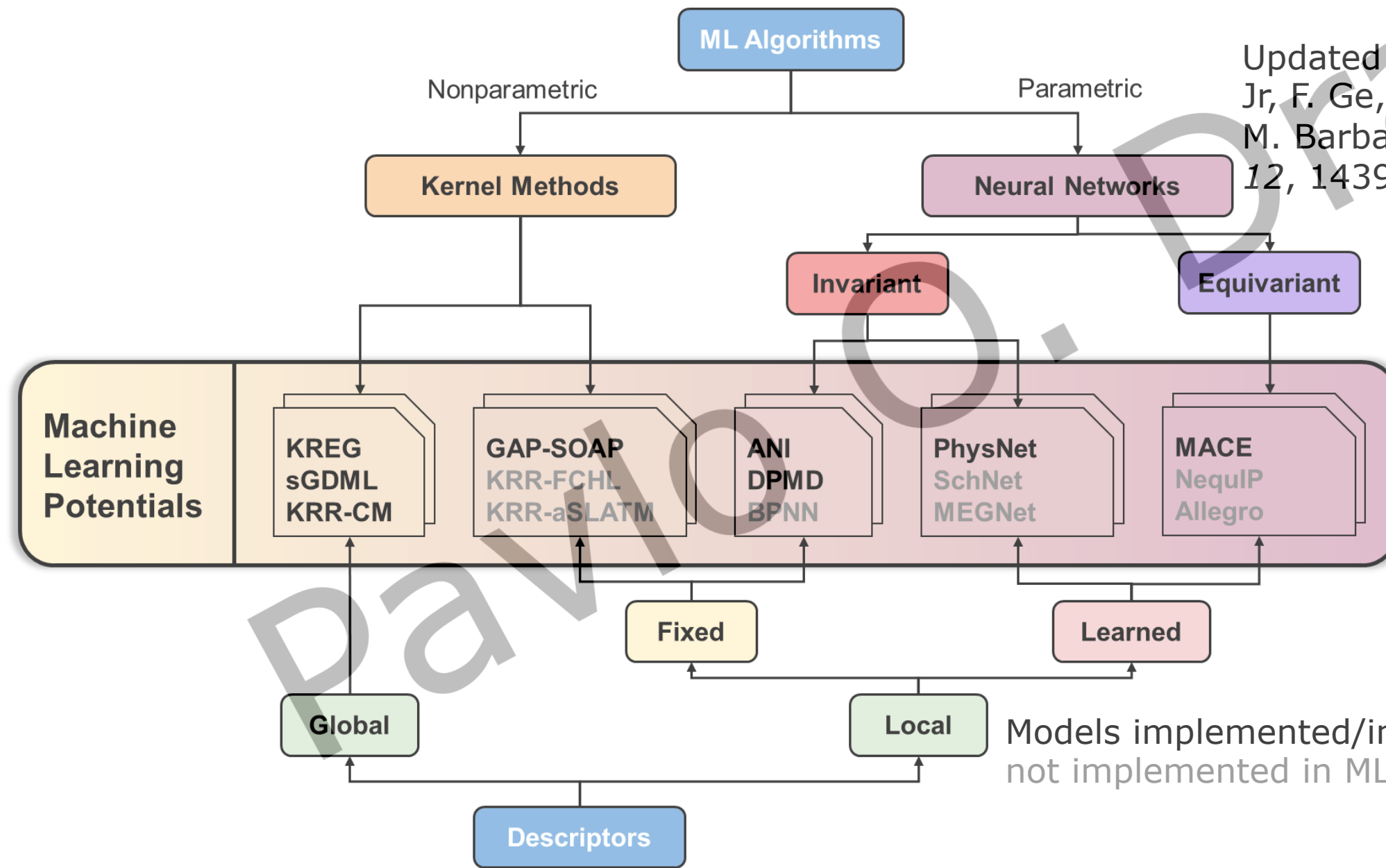
Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

$$E = f(\mathbf{R})$$

$$F_{A,d} = - \frac{\partial E}{\partial x_{A,d}} = - \frac{\partial E_{\text{machine learning potential}}(\mathbf{x})}{\partial x_{A,d}}$$



Updated based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, *12*, 14396–14413



Models implemented/interfaced in **MLatom**  
not implemented in MLatom

# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$

# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$

Use this function for making new predictions given just  $\{x'\}$

# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$

training set

train

ML model

Use this function for making new predictions given just  $\{x'\}$

# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$

training set

train

ML model

Use this function for making new predictions given just  $\{x'\}$

- Data
- Choice of x (descriptor)
- Choice of y (labels)
- Fitting function (ML algorithm, ML model)
- Optimization of ML model parameters



# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$   
training set  $\rightarrow$  train  $\rightarrow$  ML model

Use this function for making new predictions given just  $\{x'\}$

- Data
- Choice of x (descriptor)
- Choice of y (labels)
- Fitting function (**ML algorithm**, ML model)
- Optimization of ML model parameters

# ***ML algorithms***

Pavlo O. Dral

- Various types of neural networks (NN), **deep learning**
- Gaussian processes (GP)
- Kernel ridge regression (KRR)
- Support vector machines (SVMs) & support vector regression (SVR)
- Linear regression!
- Decision trees
- k-Nearest neighbor algorithm
- and many more...

Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Kernel ridge regression (KRR)

$$f(\mathbf{x}_i; \boldsymbol{\alpha}) = \sum_{j=1}^{N_{tr}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j)$$

'nonparametric model'

Examples: KREG, KRR-CM

Neural networks (NN)

Number of parameters is fixed

$$f(\mathbf{x}; \boldsymbol{\beta}) = \beta_0 + \beta_1 Z_1 + \beta_2 Z_2 + \dots$$

$$Z_m = \sigma(\alpha_{m0} + \alpha_{m1} x_1 + \alpha_{m2} x_2 + \dots)$$

'parametric model'

Examples: ANI-1ccx, AIQM1

# ***Parametric vs nonparametric algorithms***

Pavlo

Dr. Dral

$f(x; \text{parameters})$

Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Number of parameters is fixed: parametric model

Neural networks are also parametric models

Kernel ridge regression (KRR)

$$f(\mathbf{x}_i; \mathbf{p}) = \sum_{j=1}^{N_{\text{tr}}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j; \mathbf{b})$$

Number of parameters depends on number of training points:  
nonparametric model, e.g. KRR

- Various types of neural networks (NN), **deep learning**
- Gaussian processes (GP)
- Kernel ridge regression (KRR)
- Support vector machines (SVMs) & support vector regression (SVR)
- Linear regression!
- Decision trees
- k-Nearest neighbor algorithm
- and many more...

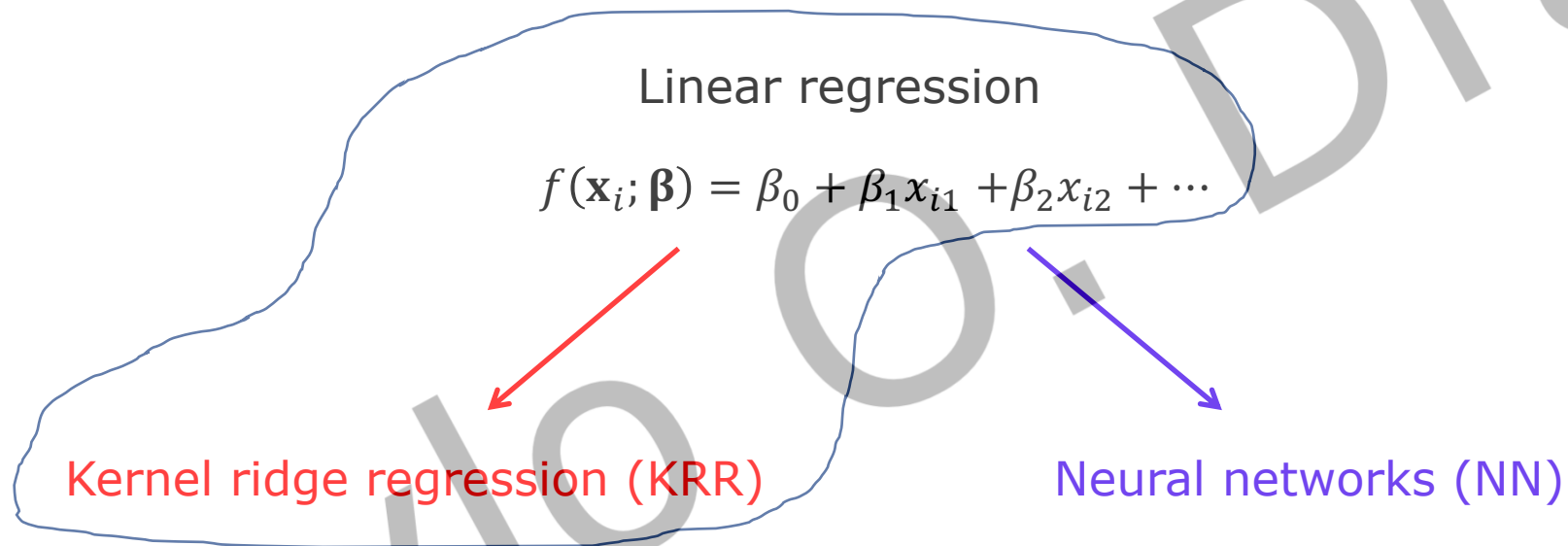
Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Kernel ridge regression (KRR)

Neural networks (NN)





Multiple linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \sum_{j=1}^p \beta_j x_{ij}$$

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \mathbf{x}_i^T \boldsymbol{\beta}$$

**How to find the coefficients  $\beta_j$ ?**

Multiple linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \sum_{j=1}^p \beta_j x_{ij}$$

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \mathbf{x}_i^T \boldsymbol{\beta}$$

We can find the coefficients  $\boldsymbol{\beta}$  using the method of least squares, where coefficients are fit to get the minimum residual sum of squares (RSS) with respect to the training set with  $N_{tr}$  reference values  $\mathbf{y}$ :

$$\arg \min_{\boldsymbol{\beta}} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \boldsymbol{\beta}) - y_i)^2$$

$$\arg \min_{\boldsymbol{\beta}} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \boldsymbol{\beta}) - y_i)^2$$

$$L(\boldsymbol{\beta}) = \sum_{i=1}^{N_{tr}} (\mathbf{x}_i^T \boldsymbol{\beta} - y_i)^2$$

$$L(\boldsymbol{\beta}) = (\mathbf{X}\boldsymbol{\beta} - \mathbf{y})^T (\mathbf{X}\boldsymbol{\beta} - \mathbf{y})$$

$$\mathbf{X} = \begin{pmatrix} x_{11} & \cdots & x_{1p} \\ \vdots & \ddots & \vdots \\ x_{N_{tr}1} & \cdots & x_{N_{tr}p} \end{pmatrix}$$

$$\frac{\partial L(\boldsymbol{\beta})}{\partial \boldsymbol{\beta}} = 2\mathbf{X}^T (\mathbf{X}\boldsymbol{\beta} - \mathbf{y}) = \mathbf{0}$$

# Linear regression

$$\mathbf{X}^T \mathbf{X} \boldsymbol{\beta} = \mathbf{X}^T \mathbf{y}$$

$$\boldsymbol{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

Linear regression has an analytical solution!

While it is very advantageous, it assumes that the data follow the linear distribution, which is often not the case

Pavlo

**Task 4: Fit linear model  $E = aR$  on a training set with 20 points sampled along  $H_2$  dissociation curve (energies  $E$  in Hartree at FCI/aug-cc-pV6Z; internuclear distances  $R$  in Angstrom)**

**Calculate  $R^2$  and residual sum of squares (RSS)**

**Task 5: Fit linear regression with intercept  $b$ ?**

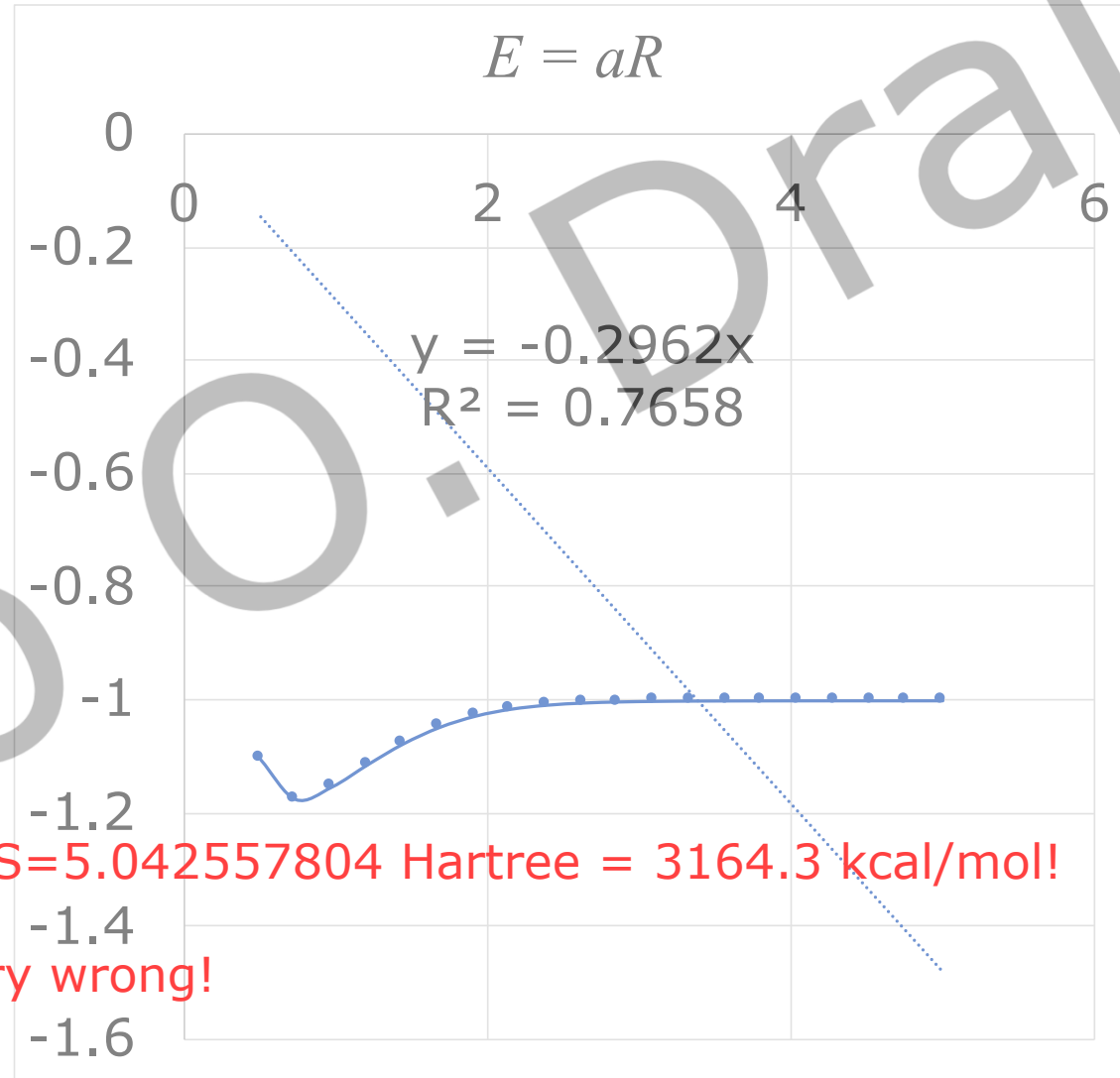
**Calculate  $R^2$  and residual sum of squares (RSS)**

$$x_{i1} = R$$

$$E = aR + b$$
$$RSS = \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \boldsymbol{\beta}) - y_i)^2$$

# Linear regression

H<sub>2</sub> dissociation curve



**Task 5: Fit linear regression with intercept  $b$ ?**

**Calculate  $R^2$  and residual sum of squares (RSS)**

$$E = aR + b$$

$$\text{RSS} = \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \boldsymbol{\beta}) - y_i)^2$$

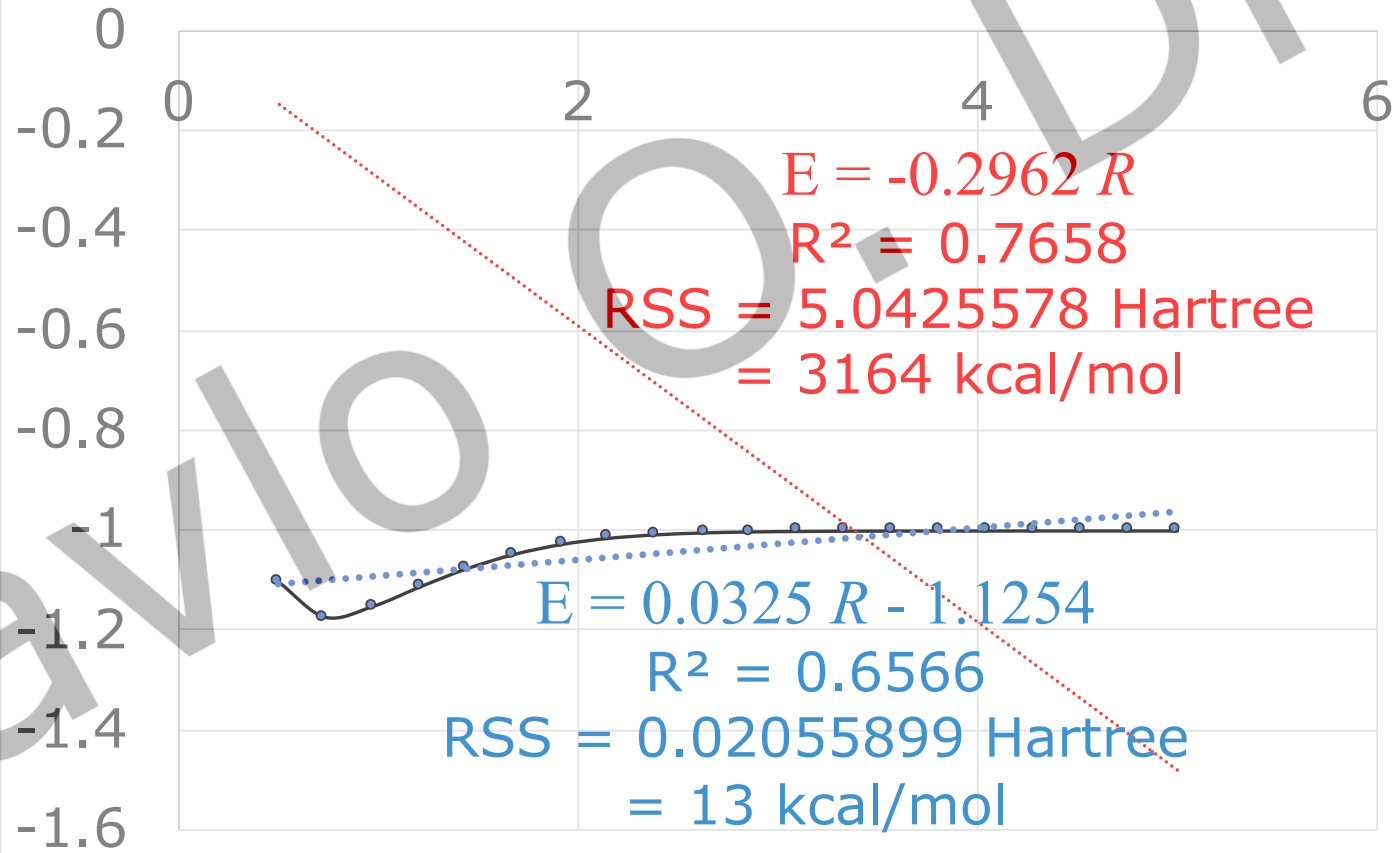


# Linear regression

Much better,  
but still qualitatively wrong!

$$E = aR$$

$$E = aR + b$$



**Q: What about linear regression with intercept  $b$ ?**

$$E = aR + b$$

It is equivalent to mapping function  $R \rightarrow (R, 1)$ ,  $\Phi((R)) = (R, 1)$ , where  $\Phi$  maps from  $p$ -dimensional input space into  $p^d$ -dimensional **feature space**

Now we can solve multiple linear regression with two variables

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_1 x_{i1} + \beta_2 x_{i2} = \beta_1 R + \beta_2 1 = aR + b$$

$$x_{i1} = R_i$$

$$x_{i2} = 1$$

$$\beta_1 = a$$

$$\beta_2 = b$$

**Any ideas how to get the dissociation curve shape right?**

Pavlo O. Dral

**Q: Any ideas how to get the dissociation curve shape right?**

We can use mapping  $R \rightarrow (R^{-6}, R^{-12}, 1)$  inspired by Lennard-Jones potential, this allows us to treat data set  $(E, R)$  nonlinear in input space  $(R)$  using vectors in feature space  $(R^{-6}, R^{-12}, 1)$

Now we can solve multiple linear regression with three variables:

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} = \beta_1 R_i^{-6} + \beta_2 R_i^{-12} + \beta_3 1 = a R_i^{-6} + b R_i^{-12} + c$$

$$x_{i1} = R_i^{-6}$$

$$x_{i2} = R_i^{-12}$$

$$x_{i3} = 1$$

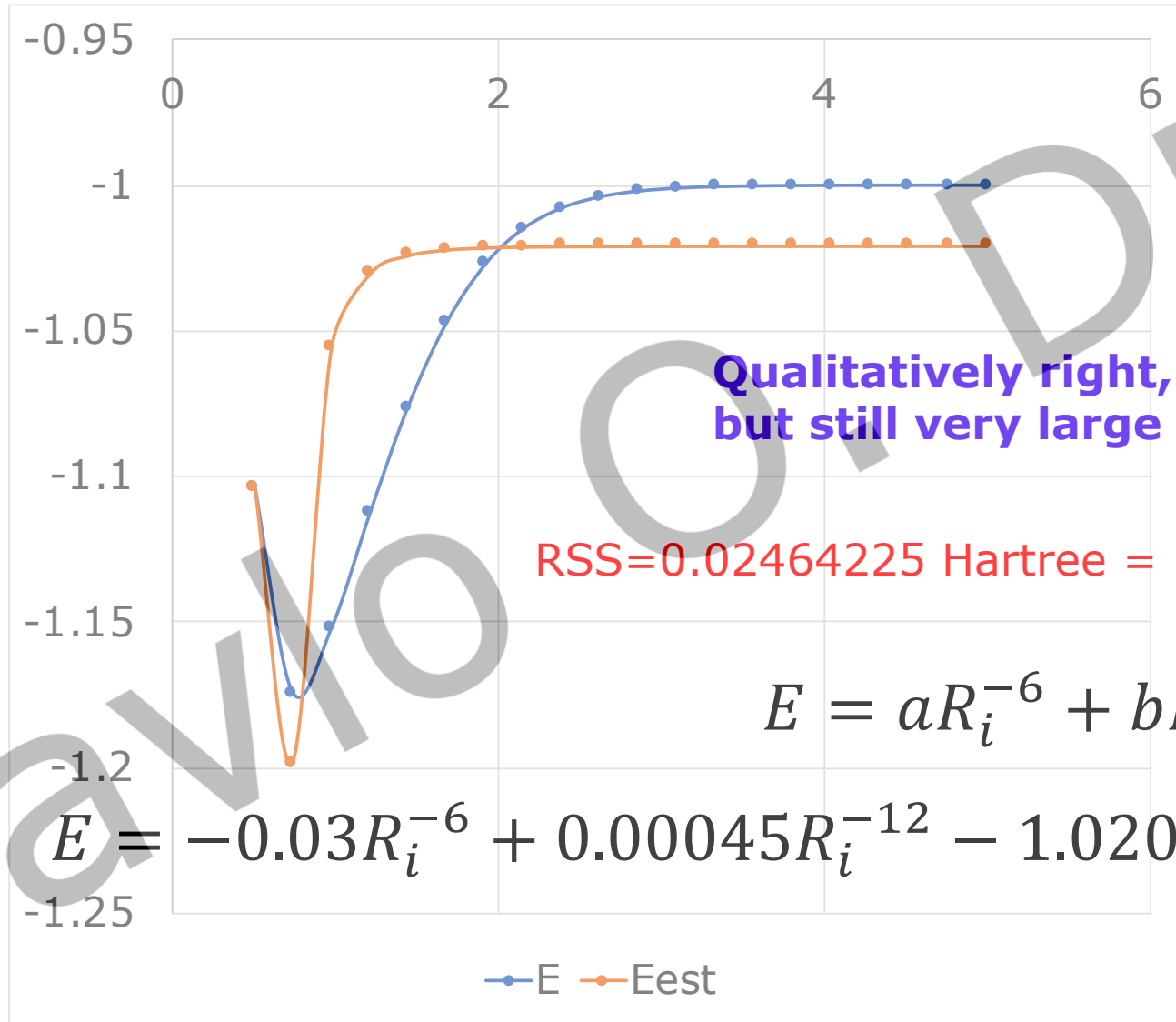
$$\beta_1 = a$$

$$\beta_2 = b$$

$$\beta_3 = c$$

Max Pinheiro Jr, P. O. Dral, Kernel methods.  
In *Quantum Chemistry in the Age of Machine Learning*,  
P. O. Dral, Ed. Elsevier: **2023**.  
Paperback ISBN: 9780323900492

# Linear regression



**Can we extend it to more variables and make it more flexible?**

Yes! We can go to infinite number of variables!

How?

Using a kernel trick

Pavlo O. Dral

Let's rewrite the linear regression equation by representing the regression coefficients via a sum over all training points:

$$f(\mathbf{x}'; \boldsymbol{\beta}) = \sum_{j=1}^p \beta_j x'_j$$

$$\beta_j = \sum_{i=1}^{N_{tr}} \alpha_i x_{ij}$$

$$f(\mathbf{x}') = \sum_{j=1}^p \left( \sum_{i=1}^{N_{tr}} \alpha_i x_{ij} \right) x'_j = \sum_{i=1}^{N_{tr}} \alpha_i \sum_{j=1}^p x_{ij} x'_j = \sum_{i=1}^{N_{tr}} \alpha_i \mathbf{x}_i^T \mathbf{x}'$$

$$\mathbf{x}_i^T \mathbf{x}' = \langle \mathbf{x}_i, \mathbf{x}' \rangle$$

Dot-product = inner product = scalar product  
of two vectors

As we have seen before, we can map vectors  $\mathbf{x}$  and  $\mathbf{x}'$  from  $p$ -dimensional input space into  $p^d$ -dimensional feature space using mapping function  $\Phi$ :

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i \Phi(\mathbf{x}_i)^T \Phi(\mathbf{x}')$$

In previous examples we knew the mapping function and explicit forms of vectors in the feature space. But all we need is a dot-product between vectors in the feature space, not their explicit forms. Such dot-product is called **kernel** denoted  $k(\mathbf{x}_i, \mathbf{x}')$  and it is calculated using vectors in the input space (not feature space!):

$$k(\mathbf{x}_i, \mathbf{x}') = \Phi(\mathbf{x}_i)^T \Phi(\mathbf{x}')$$

The kernel trick is substitution of the calculation of the dot-product using explicit representations of vectors in the feature space by using a kernel function:

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$



$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

This is a kernel-based machine learning function.

Kernel trick allows us to use tools of linear regression for data nonlinear in the input space by converting variables into (higher dimensional) feature space.

**Q: How to find the regression coefficients  $\alpha$ ?**

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

This is a kernel-based machine learning function.

Kernel trick allows us to use tools of linear regression for data nonlinear in the input space by converting variables into (higher dimensional) feature space.

We can find the coefficients  $\alpha$  using the method of least squares, where coefficients are fit to get the minimum residual sum of squares (RSS) with respect to the training set with  $N_{tr}$  reference values  $\mathbf{y}$ :

$$\arg \min_{\alpha} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \alpha) - y_i)^2$$

$$\arg \min_{\alpha} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \alpha) - y_i)^2$$

$$L(\beta) = \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \alpha) - y_i)^2$$

$$f(\mathbf{x}_i) = \sum_{j=1}^{N_{tr}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j)$$

$$L(\beta) = \sum_{i=1}^{N_{tr}} \left( \sum_{j=1}^{N_{tr}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j) - y_i \right)^2$$

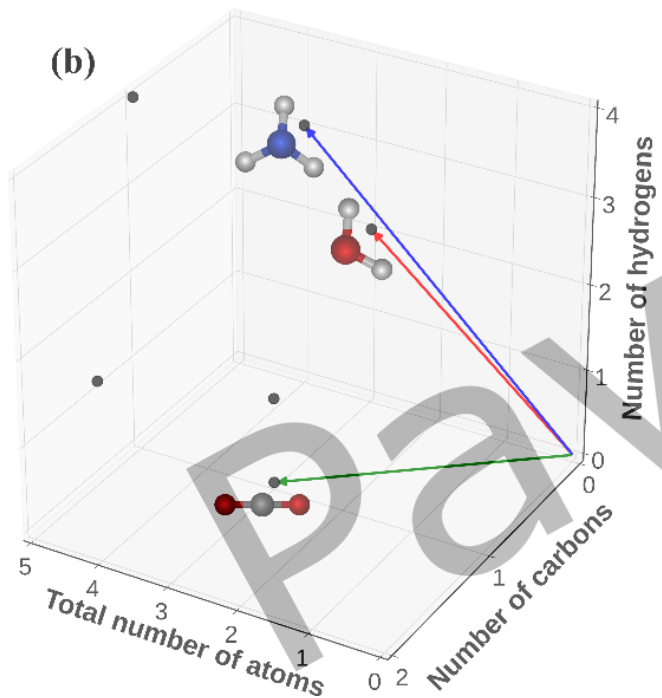
$$L(\beta) = (\mathbf{K}\alpha - \mathbf{y})^T (\mathbf{K}\alpha - \mathbf{y})$$

$$\mathbf{K} = \begin{pmatrix} k(\mathbf{x}_1, \mathbf{x}_1) & \cdots & k(\mathbf{x}_1, \mathbf{x}_{N_{tr}}) \\ \vdots & \ddots & \vdots \\ k(\mathbf{x}_{N_{tr}}, \mathbf{x}_1) & \cdots & k(\mathbf{x}_{N_{tr}}, \mathbf{x}_{N_{tr}}) \end{pmatrix}$$

Kernel matrix

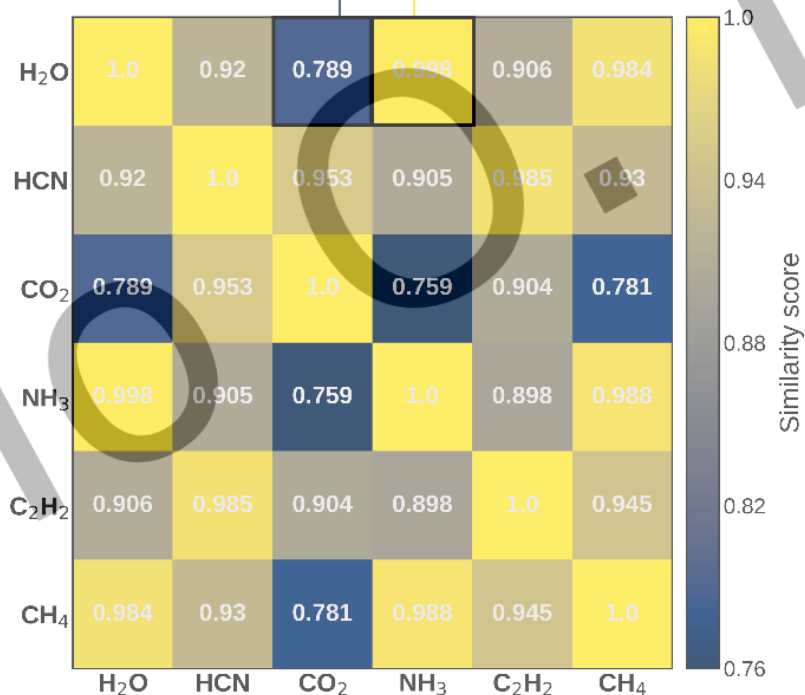
(a) Molecule num\_atoms num\_carbons num\_hydrogens

Molecule	num_atoms	num_carbons	num_hydrogens
H2O	3	0	2
HCN	3	1	1
CO2	3	1	0
NH3	4	0	3
C2H2	4	2	2
CH4	5	1	4



(c)

$$K_{ij} = \langle \text{H}_2\text{O}, \text{NH}_3 \rangle$$



Kernel matrix  
(matrix measuring similarities  
between points – here cosine)

$$\cos \theta = \mathbf{a} \cdot \mathbf{b} / (\|\mathbf{a}\| \|\mathbf{b}\|)$$

$$\arg \min_{\alpha} \sum_{i=1}^{N} (f(\mathbf{x}_i; \alpha) - y_i)^2$$

$$L(\beta) = (\mathbf{K}\alpha - \mathbf{y})^T (\mathbf{K}\alpha - \mathbf{y})$$

$$\frac{\partial L(\alpha)}{\partial \alpha} = 2\mathbf{K}^T (\mathbf{K}\alpha - \mathbf{y}) = 2\mathbf{K}(\mathbf{K}\alpha - \mathbf{y}) = 2\mathbf{K}\mathbf{K}\alpha - 2\mathbf{K}\mathbf{y} = 0$$

$$\mathbf{K}\mathbf{K}\alpha = \mathbf{K}\mathbf{y}$$

$$\mathbf{K}^{-1}\mathbf{K}\mathbf{K}\alpha = \mathbf{K}^{-1}\mathbf{K}\mathbf{y}$$

$$\mathbf{K}\alpha = \mathbf{y}$$

$$\alpha = \mathbf{K}^{-1}\mathbf{y}$$

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

This is a kernel-based machine learning function.

If the kernel function is itself a dot-product:

$$k(\mathbf{x}_i, \mathbf{x}') = \mathbf{x}_i^T \mathbf{x}'$$

The expression becomes equivalent to the linear regression as we have seen above and that is why such a dot-product kernel is also called "linear kernel":

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i \mathbf{x}_i^T \mathbf{x}' = \sum_{j=1}^p \beta_j x'_j$$

$$\beta_j = \sum_{i=1}^{N_{tr}} \alpha_i x_{ij}$$

$$\beta_j = \sum_{i=1}^{N_{tr}} \alpha_i x_{ij}$$

**One can get the same  $\beta_j$  for infinite combinations of  $\alpha_i$**

**Some solutions will have very large  $\alpha_i$  with opposite signs trying to compensate each other**

$$\beta = \sum_{i=1}^{N_{tr}=20} \alpha_i x_i = \sum_{i=1}^{N_{tr}=20} \alpha_i R_i$$

$$\beta = (-0.2962 \cdot 2) \cdot 0.5 + \sum_{i=2}^{N_{tr}=20} 0x_i$$

$$\beta = 198476910439 \cdot 0.5 - 19847691043.95924 \cdot 5 + \sum_{i=3}^{N_{tr}=20} 0x_i$$

$$f(x; \beta) = \beta x = -0.2962x$$

$$\alpha = K^{-1}y$$

**Prone to overfitting,  
numerically unstable  
Often, K is not invertible  
matrix**

Ridge regression – belongs to shrinkage methods (useful for feature importance analysis)

$$\arg \min_{\beta} (\mathbf{X}\beta - \mathbf{y})^T (\mathbf{X}\beta - \mathbf{y}) + \lambda \beta^T \beta$$

$$\beta = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^T \mathbf{y}$$

Another example of shrinkage method is the lasso

$$\arg \min_{\beta} (\mathbf{X}\beta - \mathbf{y})^T (\mathbf{X}\beta - \mathbf{y}) + \lambda \sum_{i=1}^p |\beta_i|$$

Identity matrix

$$\mathbf{I} = \begin{pmatrix} 1 & \cdots & 0 \\ \vdots & \ddots & \vdots \\ 0 & \cdots & 1 \end{pmatrix}$$

Kernel ridge regression (KRR)

$$\arg \min_{\alpha} (\mathbf{K}\alpha - \mathbf{y})^T (\mathbf{K}\alpha - \mathbf{y}) + \lambda \alpha^T \mathbf{K}\alpha$$

$$\alpha = (\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{y}$$

Coefficient magnitude is forced to shrunk with larger  $\lambda$  in these methods  
 $\lambda$  is nonnegative regularization hyperparameter, smoothens function and makes solution numerically more stable.



$$f(\mathbf{x}') = \sum_{i=1}^{N_{\text{tr}}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

This is a kernel-based machine learning function.

One of the popular kernel functions is the Gaussian kernel function:

$$k(\mathbf{x}_i, \mathbf{x}_j) = \exp\left(-\frac{1}{2\sigma^2} \sum_s^{N_x} (x_{i,s} - x_{j,s})^2\right)$$

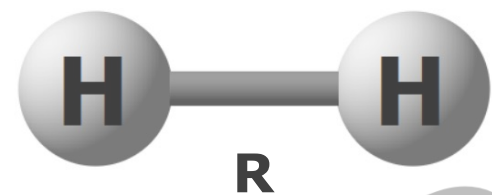
It maps vectors  $\mathbf{x}$  from  $N_x$ -dimensional input space into infinite-dimensional feature space.

$\sigma$  is a positive hyperparameter defining the length scale of the Gaussian function.

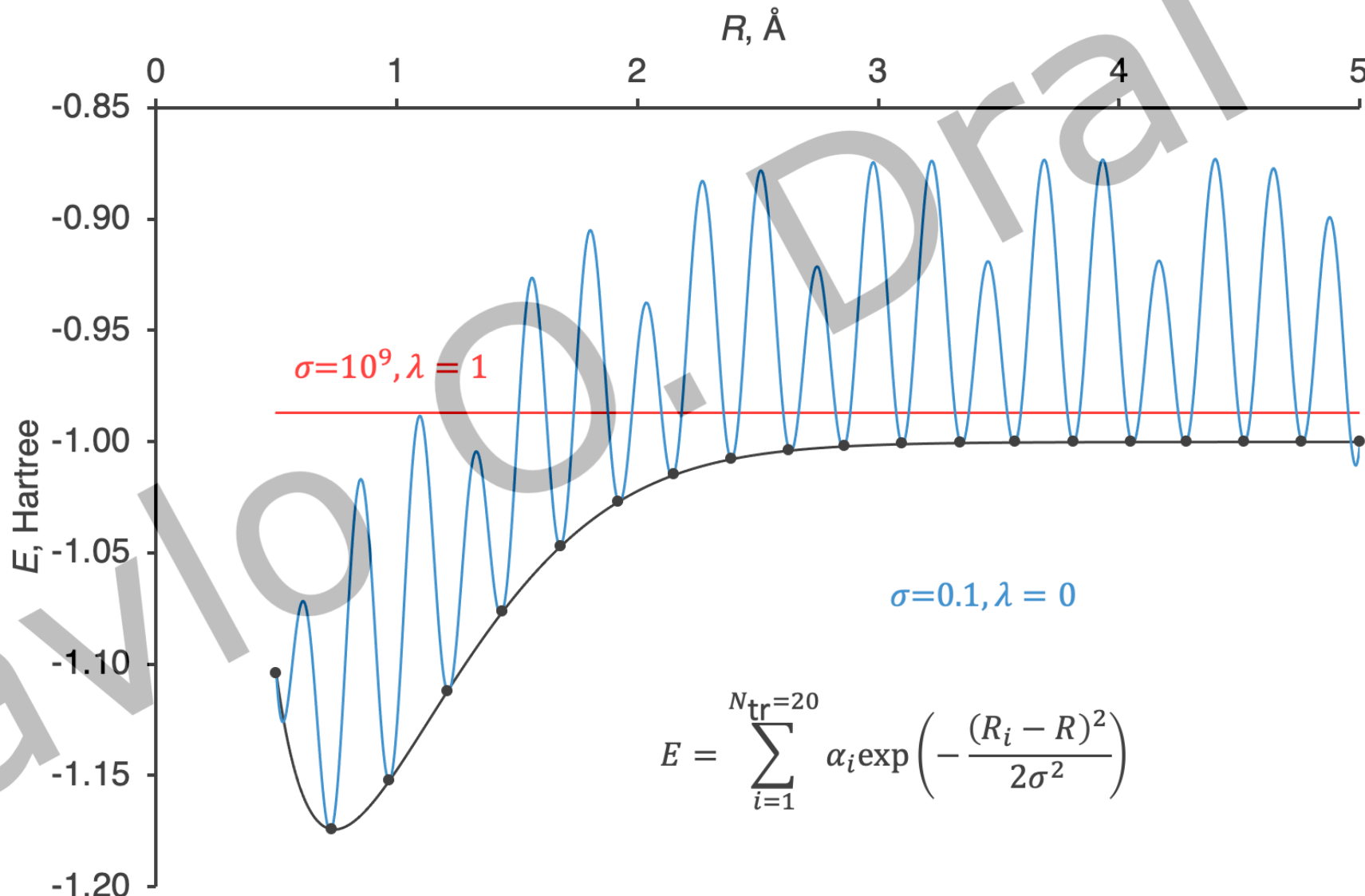
Many ML algorithms can give you practically ideal predictions for their own training set

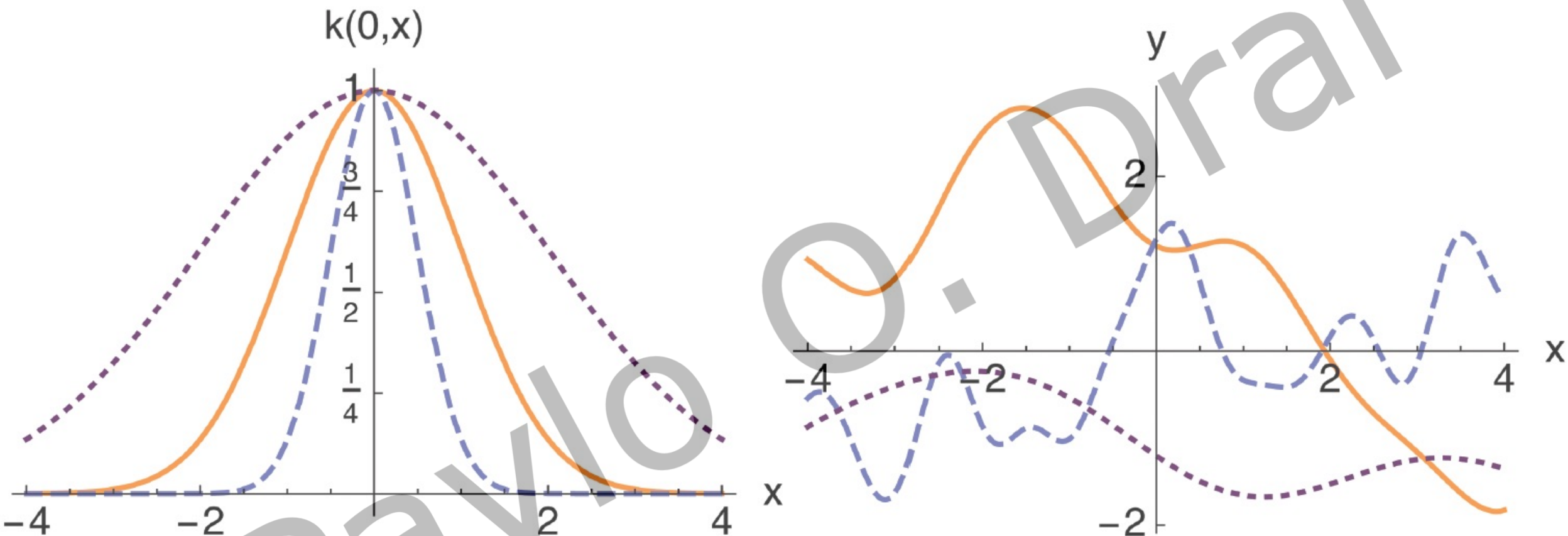
**Underfitting**  
Low variance  
High bias

H<sub>2</sub> dissociation curve



**Overfitting**  
High variance  
Low bias





(b) Gaussian kernel with  $\sigma = 0.5, 1, 2$  (dashed, solid, dotted lines).

Take KRR with Gaussian kernel

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i \exp\left(-\frac{1}{2\sigma^2} \sum_s^{N_x} (x_{i,s} - x'_s)^2\right)$$

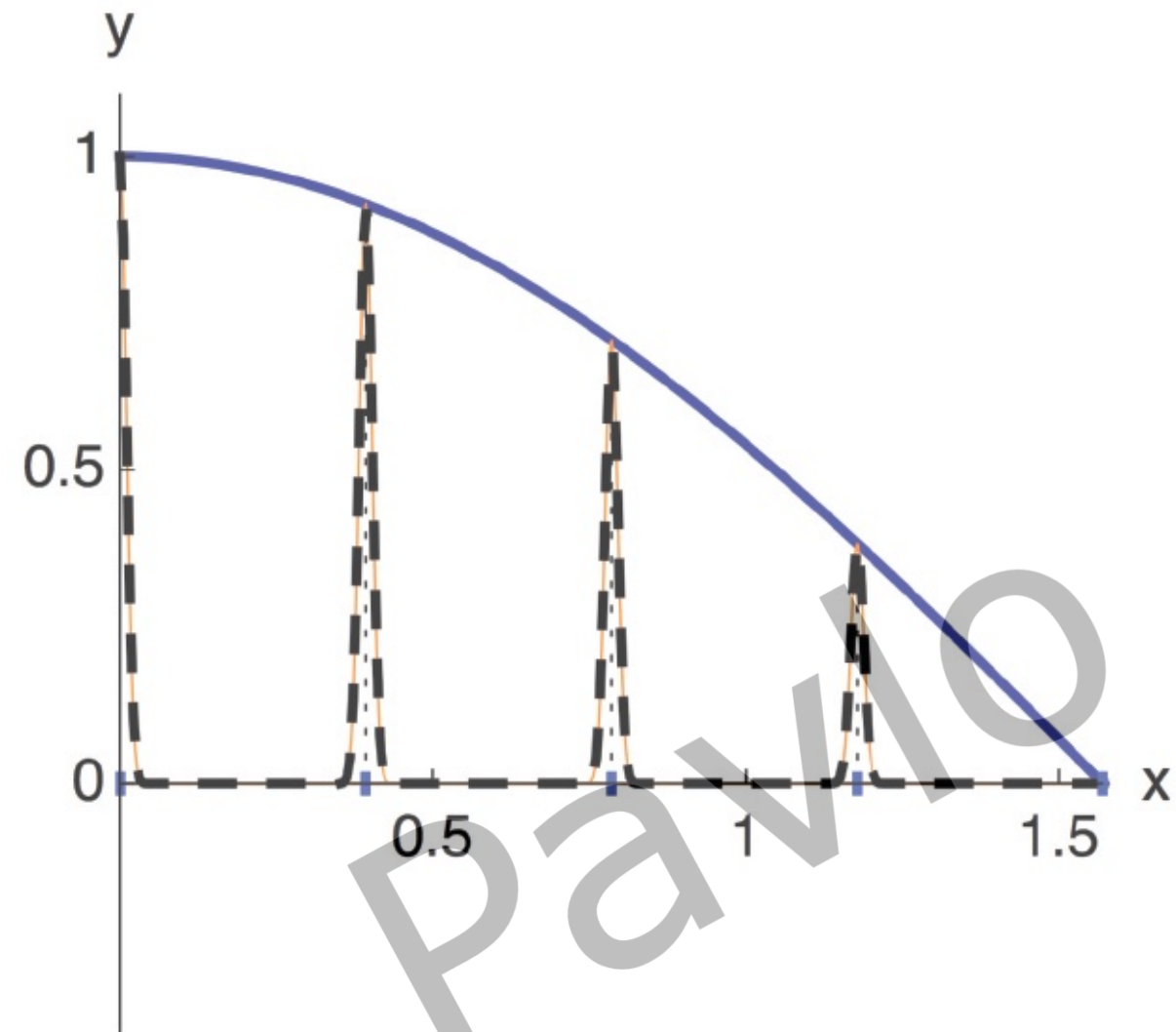
and consider what happens for very small  $\sigma \rightarrow 0$ :

$$f(\mathbf{x}') = \begin{cases} \alpha_i, & \text{for } \mathbf{x}' = \mathbf{x}_i \\ 0, & \text{for } \mathbf{x}' \neq \mathbf{x}_i \end{cases}$$

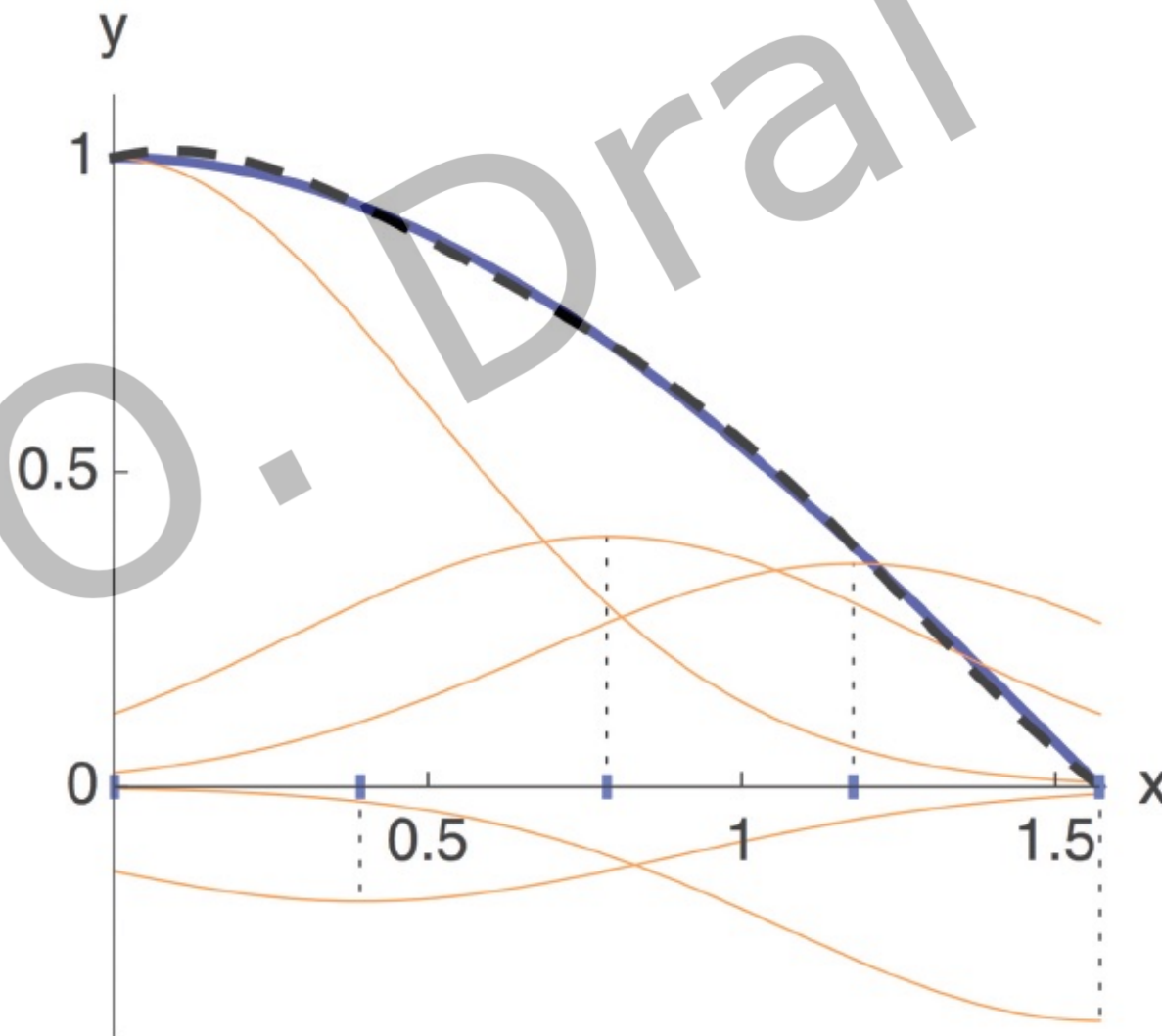
and consider what happens for very large  $\sigma \rightarrow \infty$ :

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i = \text{const}$$

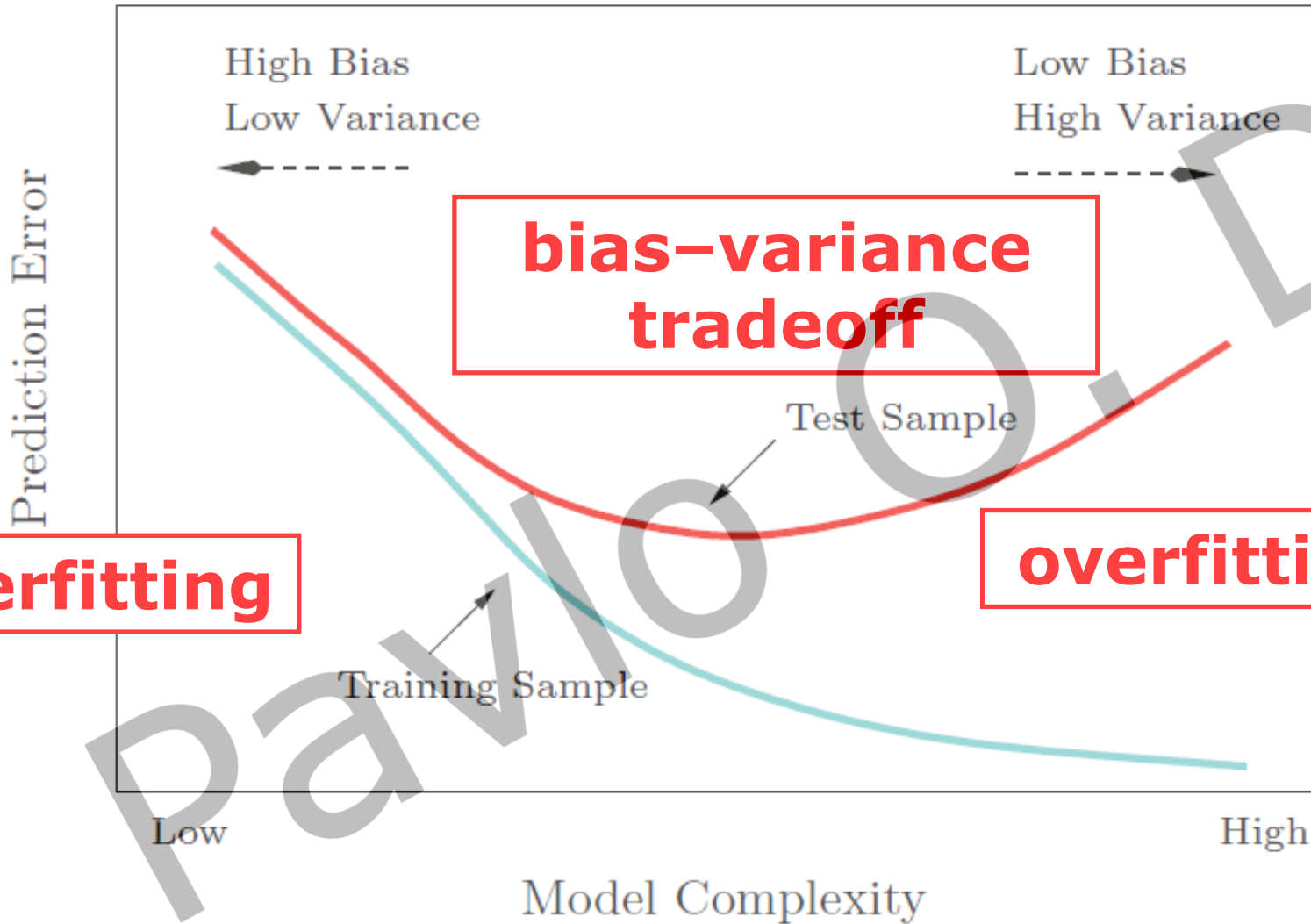
# KRR with Gaussian kernel



(a) Overfitting ( $\sigma = 0.01$ )



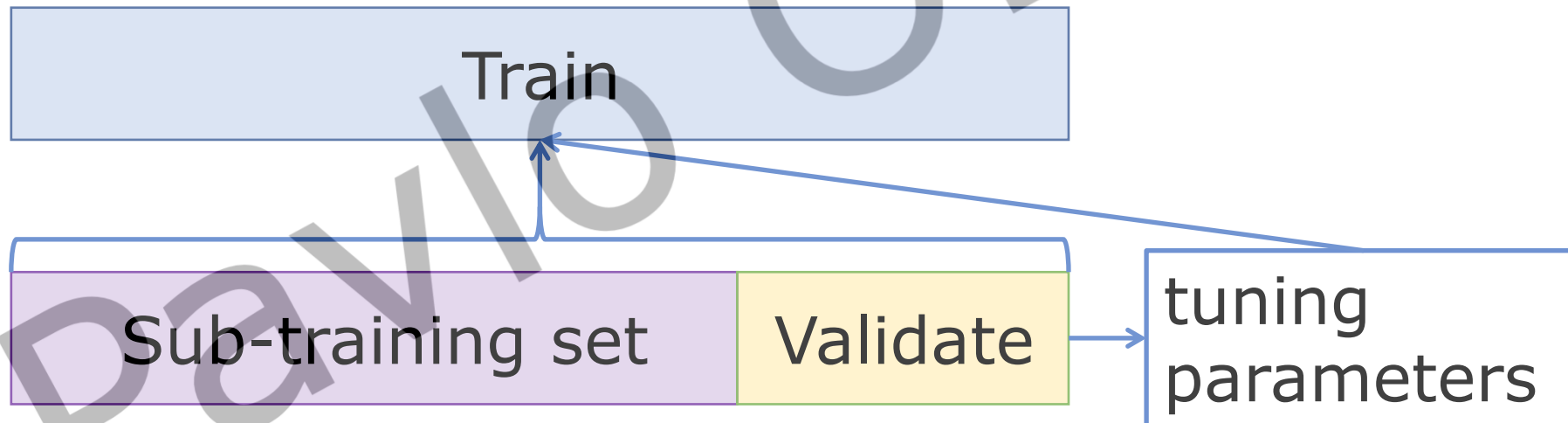
(b) Fitting ( $\sigma = 0.5$ )



**Q: How to choose hyperparameters?**

Pavlo O. Dral

We target minimal error **not** in the training set, but in the validation set for models trained on the sub-training set.

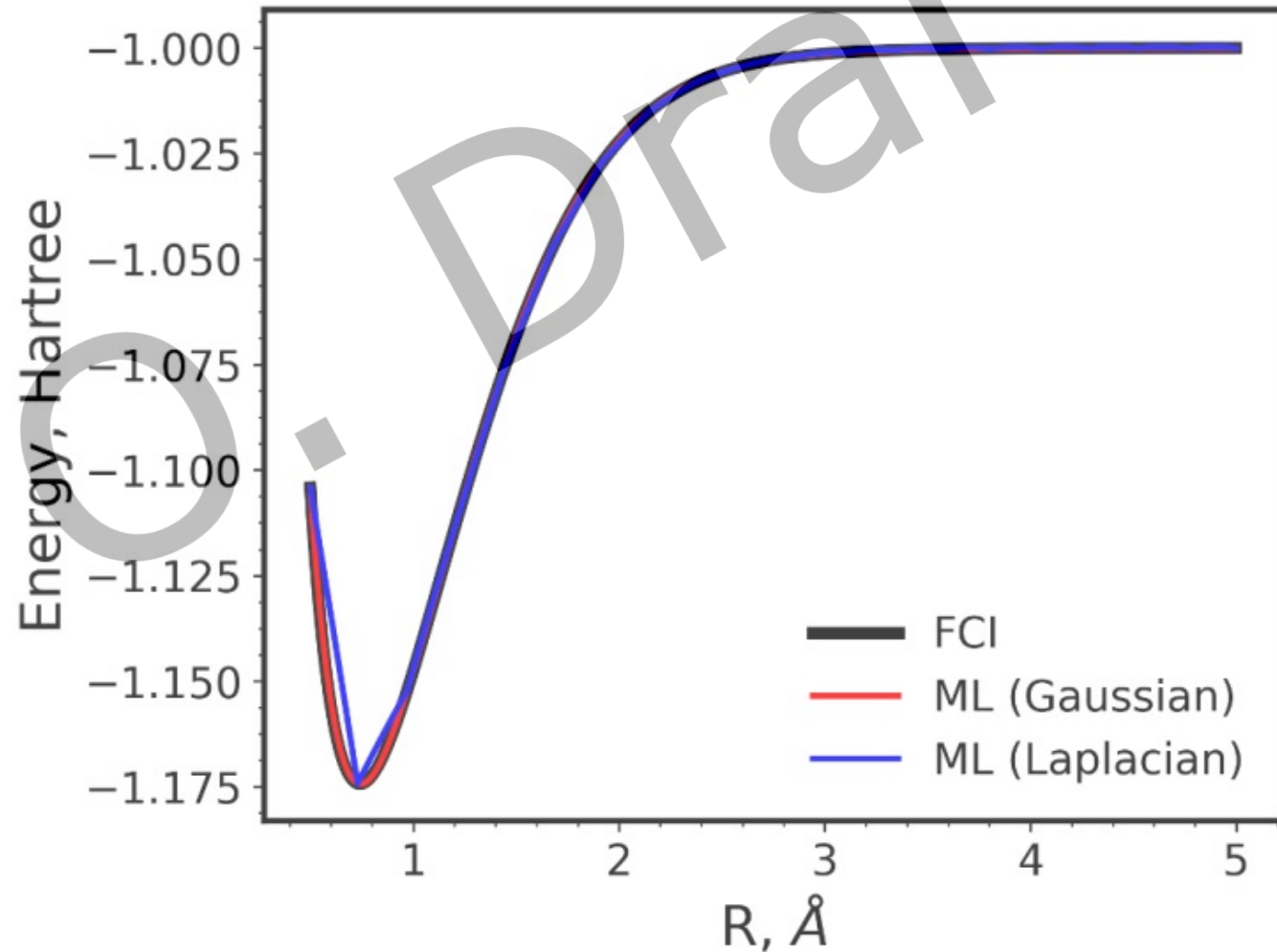




H<sub>2</sub> dissociation curve

Full CI calculations:  
more than 30 min  
for one value of R.

ML trained on 20 points needs  
less than 1 sec.  
for hundreds of other points



# 5-fold Cross-validation

Training set with randomly shuffled items

Pavlo O. Dral

# 5-fold Cross-validation

Training set with randomly shuffled items

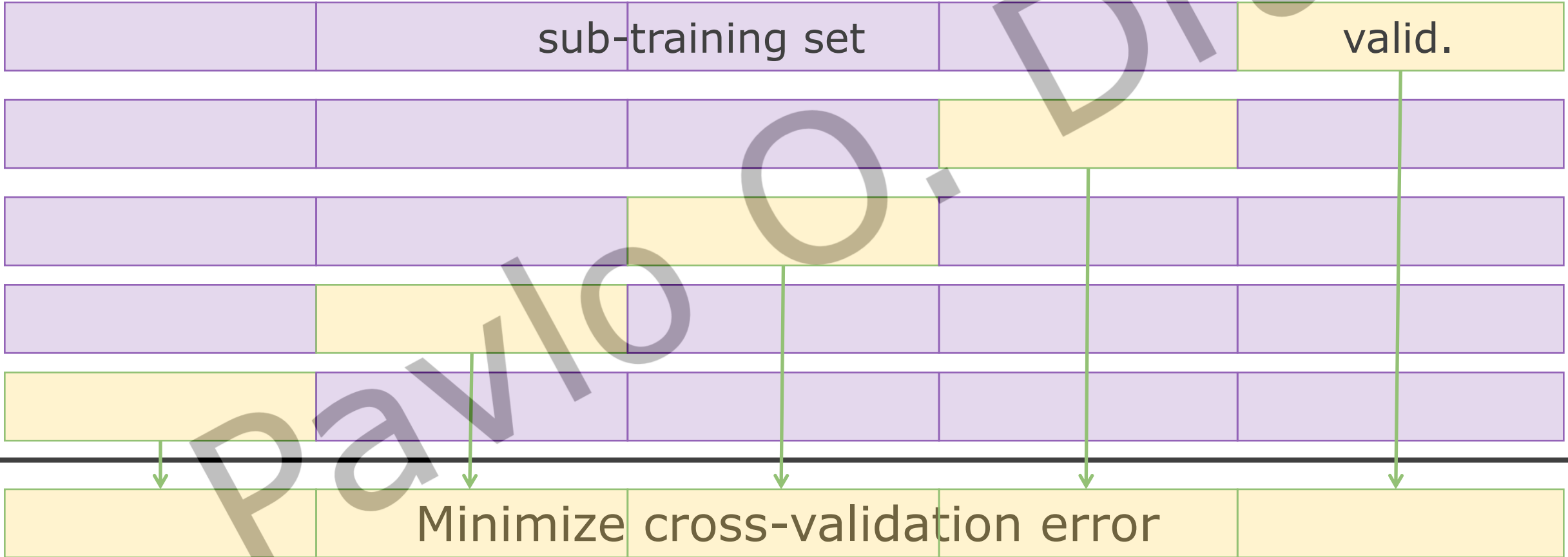
sub-training set

valid.

Pavlo O. Dral

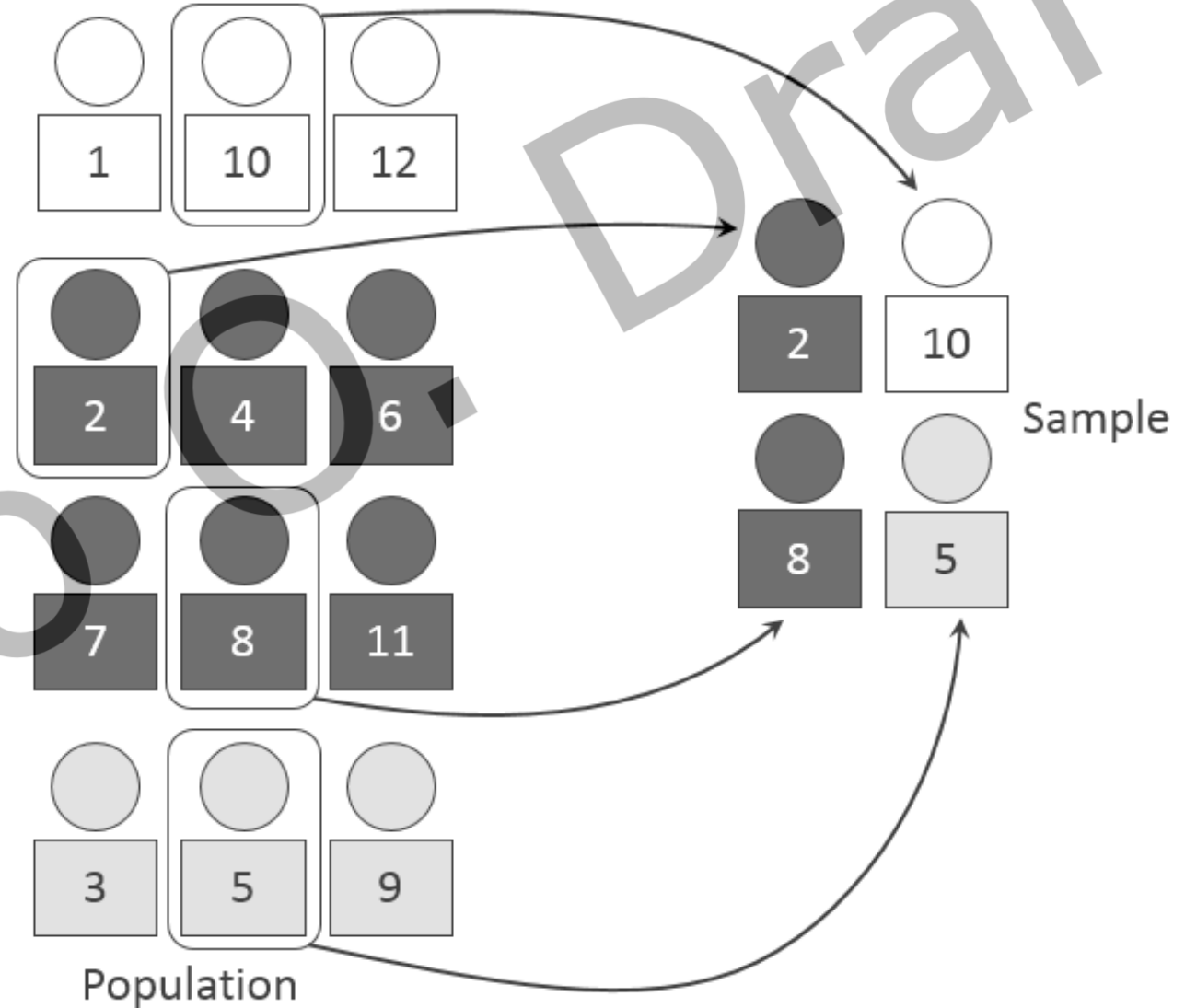
# 5-fold Cross-validation

Training set with randomly shuffled items



Random sampling for model selection is not always a good idea

Sometimes, **stratification** is preferable



Pavlo

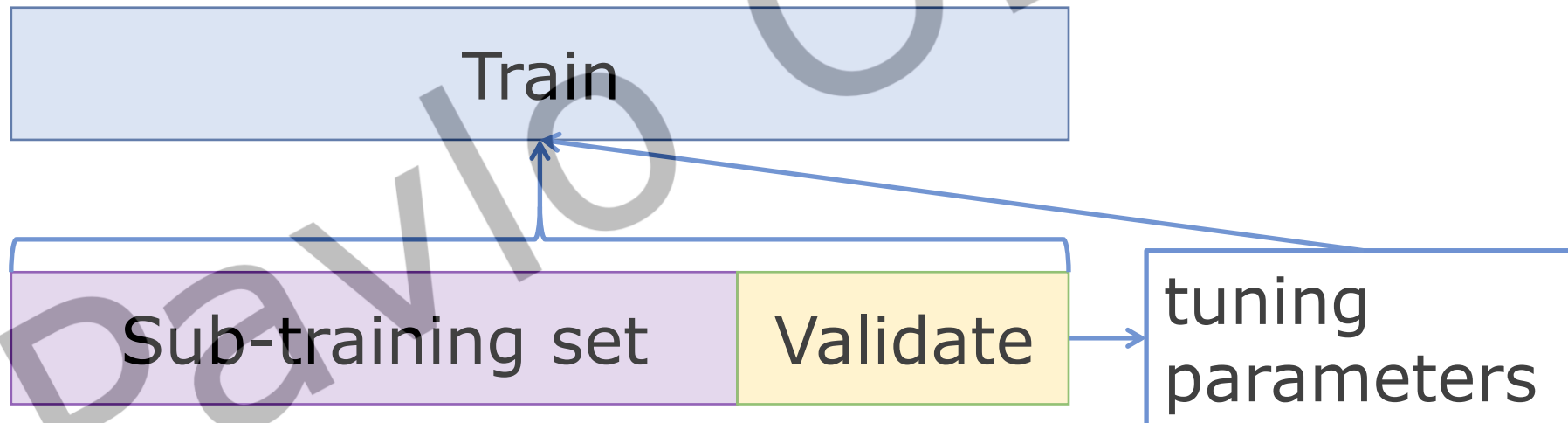
Drail

# ***Model Evaluation***

***(estimation of the generalization error)***

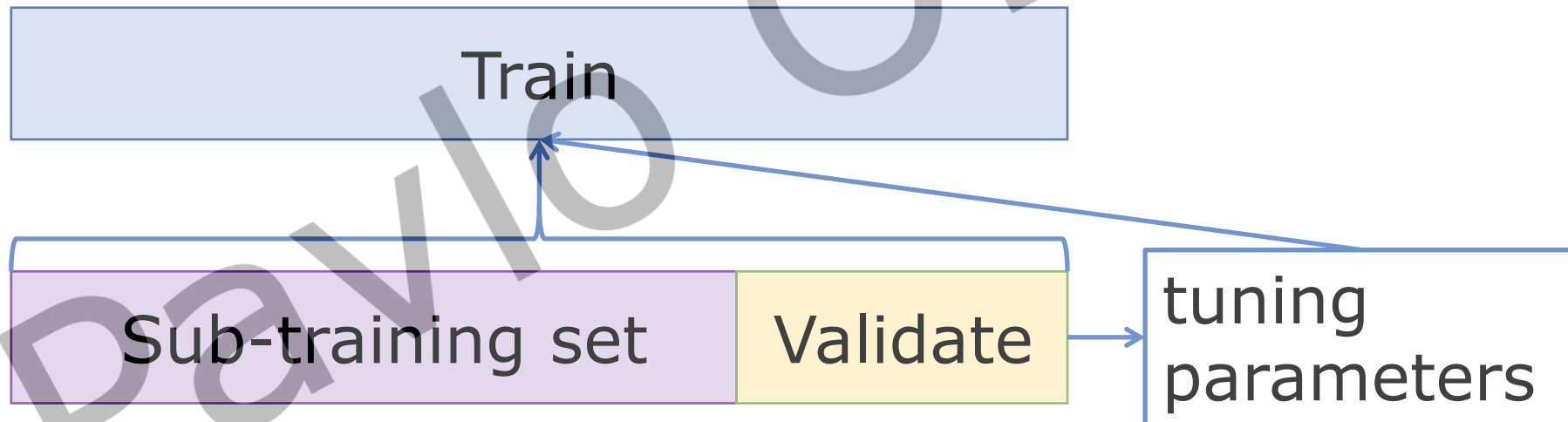
Pavlo O. Dral

- Often ML error for its own training set is close to zero



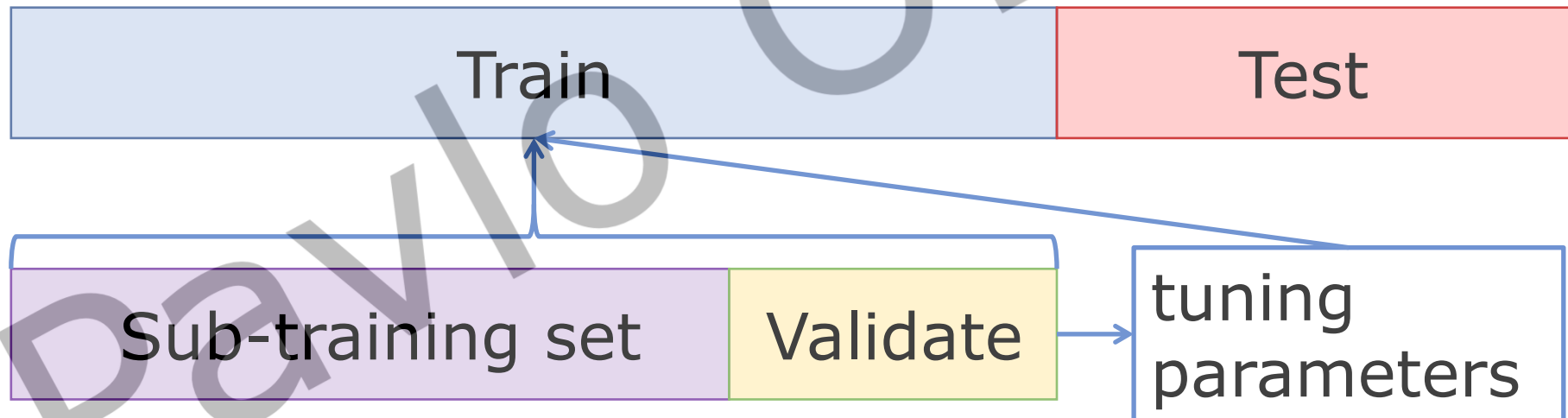
# ML: Error Estimation

- Often ML error for its own training set is close to zero
- Using errors in the validation set would be also incorrect, because their minimization is a part of the training process



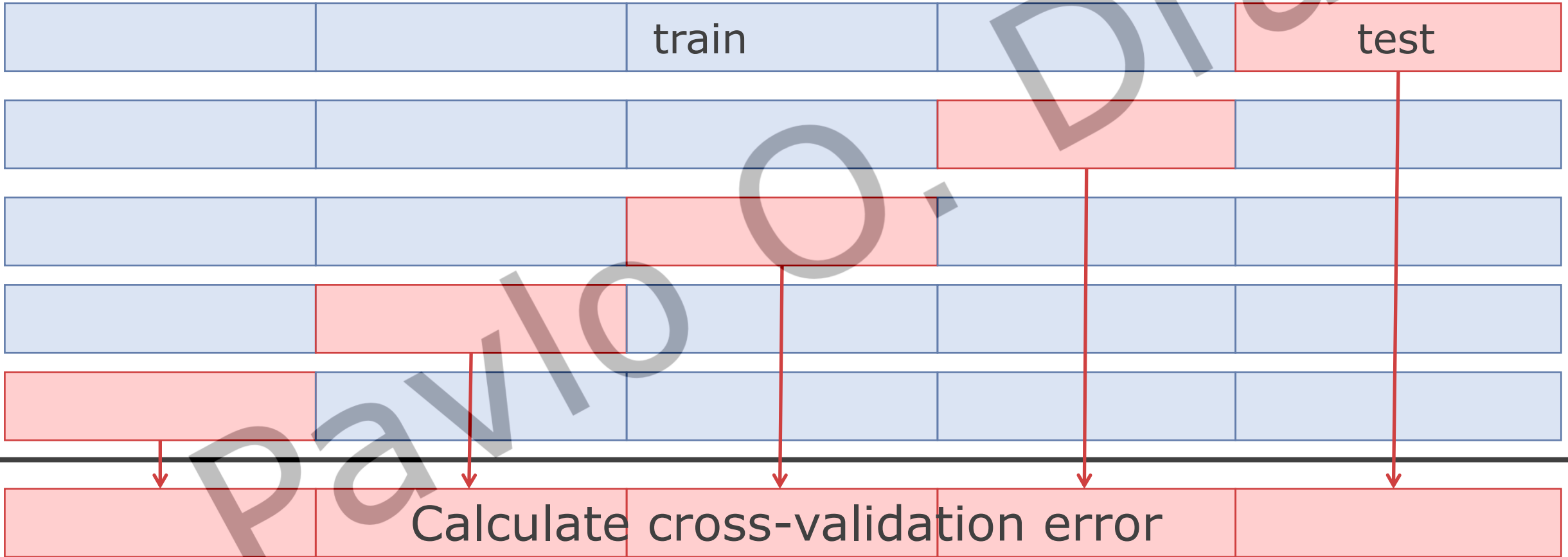


- Often ML error for its own training set is close to zero
- Using errors in the validation set would be also incorrect, because their minimization is a part of the training process
- We should estimate errors on a **completely independent test set**



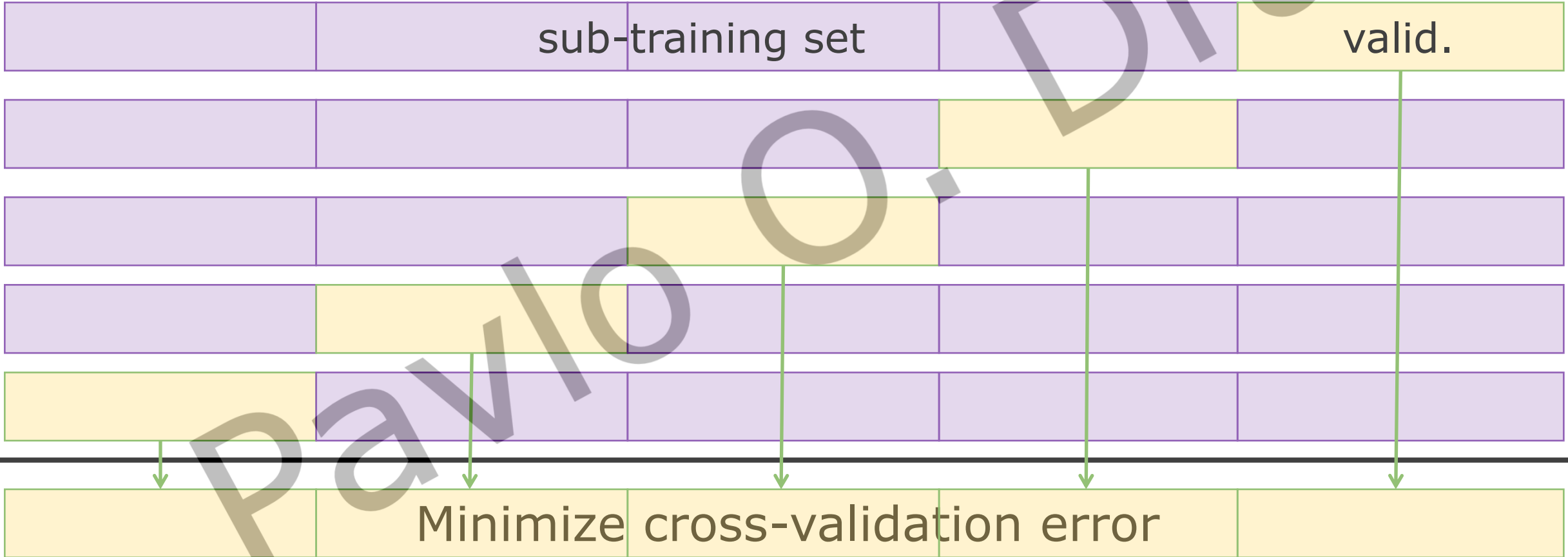
# 5-fold Cross-validation

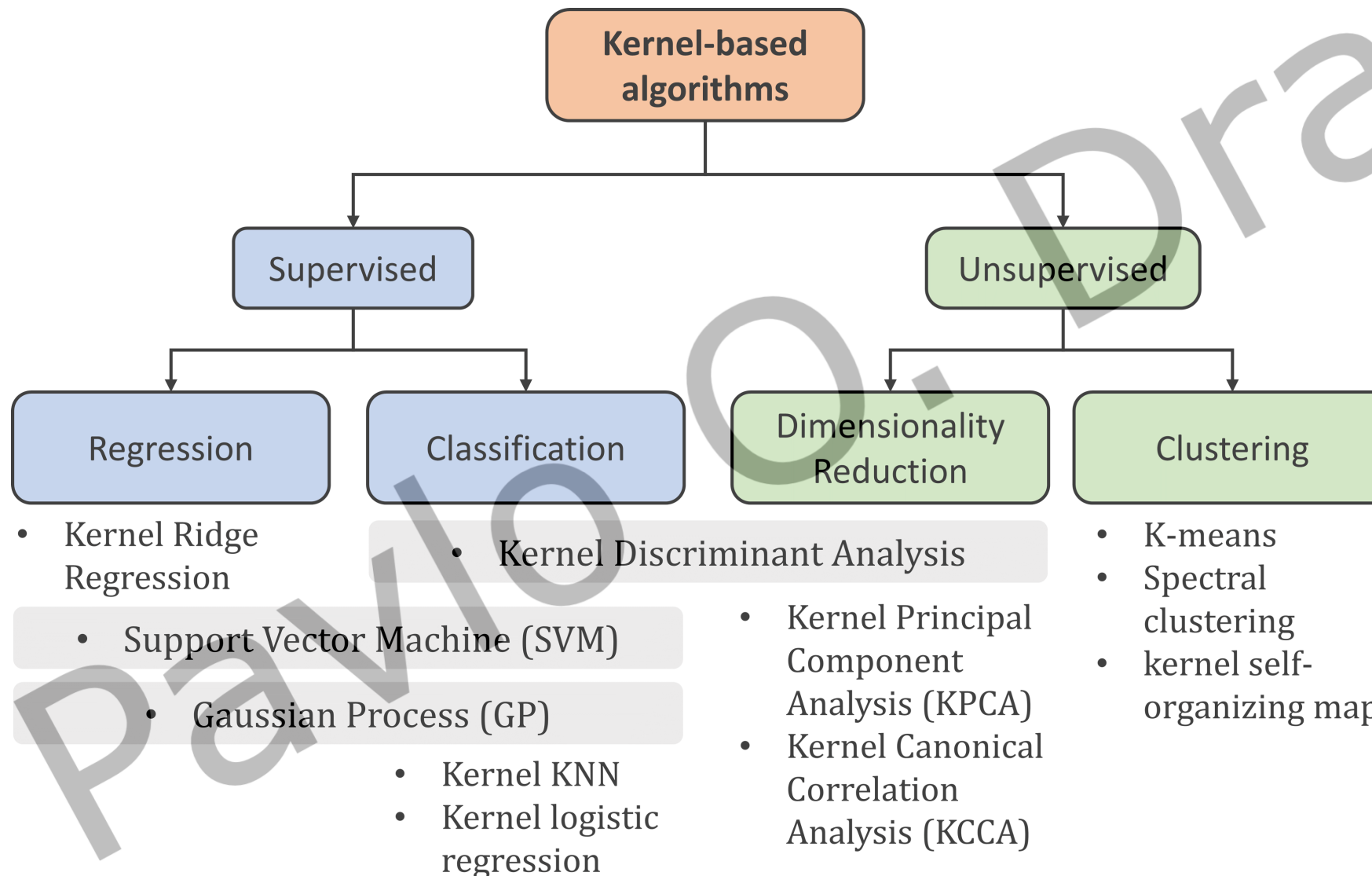
Entire set with randomly shuffled original data



# 5-fold Cross-validation

Training set with randomly shuffled items





- Kernel ridge regression (KRR)

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

- Gaussian processes (GP, kriging)

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

Prediction  
functions are  
the same for  
KRR and GP!

- Support vector machines (SVM)

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i y_i k(\mathbf{x}_i, \mathbf{x}'), \quad 0 < \alpha_i < C$$

Pavlio

Dral

- Kernel ridge regression gives the same prediction as Gaussian processes:

$$f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

- Gaussian processes also provide:
  - variance  $V$

$$V(\mathbf{x}') = k(\mathbf{x}', \mathbf{x}') - \mathbf{k}'^T (\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{k}'$$

$$\mathbf{k}' = \begin{pmatrix} k(\mathbf{x}_1, \mathbf{x}') \\ \vdots \\ k(\mathbf{x}_{N_{tr}}, \mathbf{x}') \end{pmatrix}$$

- Marginal likelihood:

$$\log p(\mathbf{y}|\mathbf{X}) = -\frac{1}{2} \mathbf{y}^T \boldsymbol{\alpha} - \frac{1}{2} \log |\mathbf{K} + \lambda \mathbf{I}| - \frac{N_{tr}}{2} \log 2\pi$$

Hyperparameters in kernel function can be found by optimizing log marginal likelihood, for which derivatives are taken, e.g.  $\frac{\partial \log p(\mathbf{y}|\mathbf{X}, \sigma)}{\partial \sigma}$

Advantages of kernel methods:

- Nonparametric models, i.e., do not assume a specific behavior of data (compare to parametric model such as linear regression)
- Explicitly incorporate training data, thus very flexible and accurate
- Closed (analytical) solution, i.e. fast training

$$\alpha = (\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{y} \quad f(\mathbf{x}') = \sum_{i=1}^{N_{tr}} \alpha_i k(\mathbf{x}_i, \mathbf{x}')$$

Disadvantages:

- Slow training for lots of training data (scales as  $O(N_{tr}^3)$ )
- Requires lots of RAM to store the kernel matrix (scales as  $O(N_{tr}^2)$ )
- Prediction time slows down with more training data (scales as  $O(N_{tr}^1)$ )

**Table 1** Required memory (RAM) for storing kernel matrix built with increasing number of training points.

Training set size	RAM size
$100 = 10^2$	78 kB
$1000 = 10^3$	7.6 MB
$10,000 = 10^4$	0.75 GB
$50,000 = 5 \times 10^4$	19 GB
$100,000 = 10^5$	75 GB
$500,000 = 5 \times 10^5$	1.8 TB
$1000,000 = 10^6$	7.3 TB

**Table 2** CPU time needed for calculating regression coefficients for increasing number of training points, assuming that it takes 10 s for 10,000 training points.

Training set size	Time
$100 = 10^2$	0.01 milliseconds
$1000 = 10^3$	0.01 s
$10,000 = 10^4$	10 s
$50,000 = 5 \times 10^4$	21 min
$100,000 = 10^5$	2.8 h
$500,000 = 5 \times 10^5$	15 days
$1000,000 = 10^6$	3.9 months



## Solutions:

- Reduce the training set by selecting the most relevant points[1,2]
- Sparsification techniques[3]
- Construct high-dimensional kernels as products of one-dimensional kernels[4]

See, for example:

[1] Dral, Owens, Yurchenko, Thiel, *J. Chem. Phys.* **2017**, 146, 244108

[2] Hu, Xie, Li, Li, Lan, *J. Phys. Chem. Lett.* **2018**, 9, 2725

[3] Bartók, Csányi, *Int. J. Quantum Chem.* **2015**, 115, 1051

[4] Unke, Meuwly, *J. Chem. Inf. Model.* **2017**, 57, 1923

# ***Neural networks***

Pavlo O. Dral

Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Kernel ridge regression (KRR)

Neural networks (NN)

Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Kernel ridge regression (KRR)

Neural networks (NN)

## Linear regression

$$\hat{y} = f(\mathbf{x}; \mathbf{w}, b) = b + w_1x_1 + w_2x_2 + \cdots + w_px_p = \mathbf{x}^T \mathbf{w} + b$$



Neural networks (NNs): the single hidden layer, feed-forward network

$$\hat{y} = f(\mathbf{x}; \boldsymbol{\alpha}, \mathbf{a}, \mathbf{w}, b) = b + w_1h_1(\mathbf{x}; \boldsymbol{\alpha}_1, a_1) + \cdots + w_Mh_M(\mathbf{x}; \boldsymbol{\alpha}_M, a_M) = \mathbf{h}^T \mathbf{w} + b$$

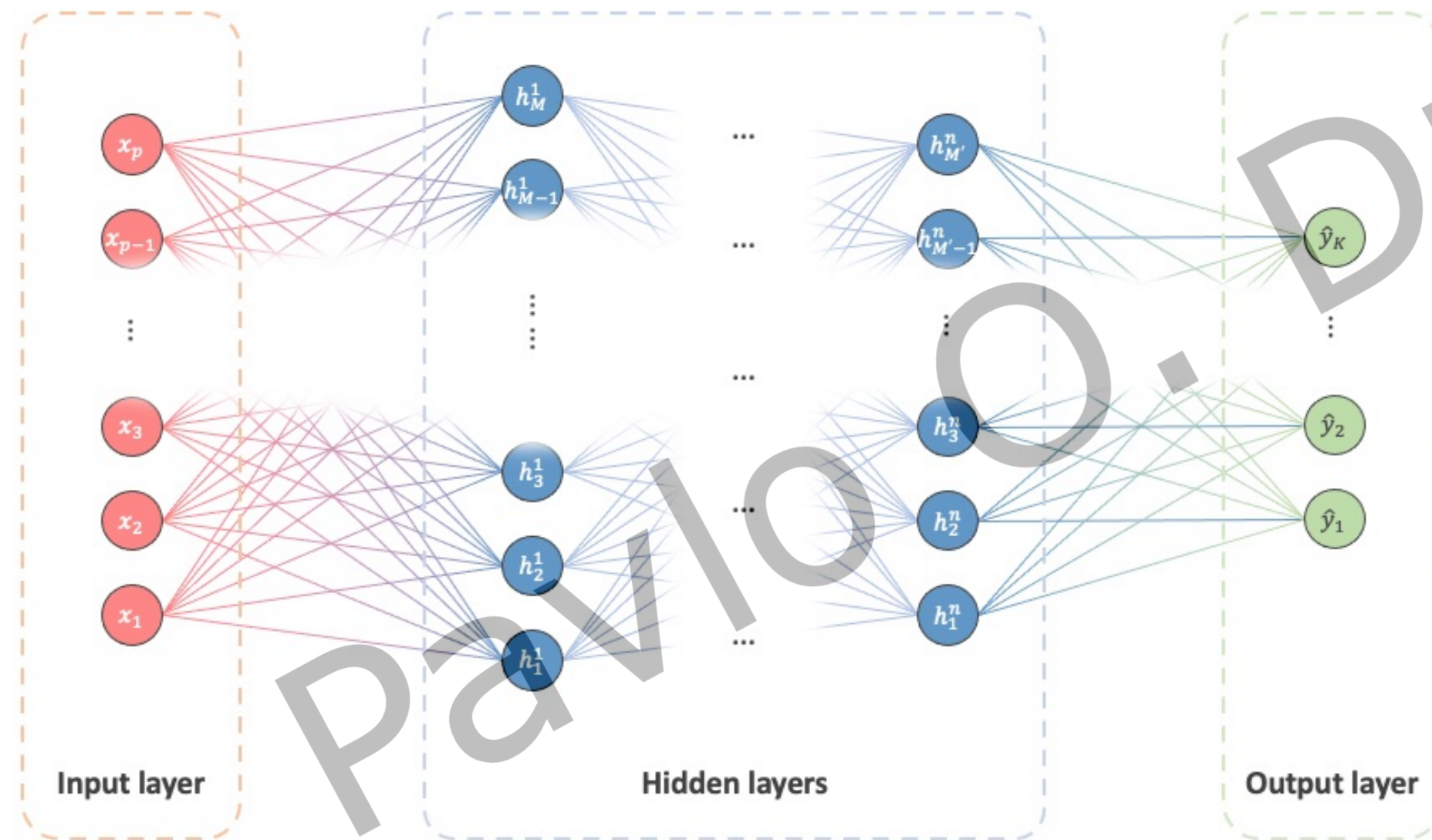
$$h_m(\mathbf{x}; \boldsymbol{\alpha}_m, a_m) = g(\mathbf{a}_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \cdots + \alpha_{mp}x_p) = g(\mathbf{x}^T \boldsymbol{\alpha}_m + a_m)$$

$$\hat{y} = f(\mathbf{x}; \boldsymbol{\alpha}, \mathbf{a}, \mathbf{w}, b) = f^{(2)}(\mathbf{h}; \mathbf{w}, b) = f^{(2)}(f^{(1)}(\mathbf{x}))$$

Activation functions:

$$g(v) = \exp(-a(v - c)^2) \quad \text{radial basis function (RBF)}$$

# Neural network (NN)



NN consists of

- layers
- nodes=units=neurons

The single hidden layer, feed-forward NN

$w, \alpha \dots$  weights

$a, b \dots$  biases

$$f(\mathbf{x}; \boldsymbol{\alpha}, \mathbf{a}, \mathbf{w}, b) = f^{(2)}(\mathbf{h}; \mathbf{w}, b)$$

## Linear regression

$$\hat{y} = f(\mathbf{x}; \mathbf{w}, b) = b + w_1x_1 + w_2x_2 + \dots + w_px_p = \mathbf{x}^T \mathbf{w} + b$$



Neural networks (NNs): the single hidden layer, feed-forward network

$$\hat{y} = f(\mathbf{x}; \boldsymbol{\alpha}, \mathbf{a}, \mathbf{w}, b) = b + w_1h_1(\mathbf{x}; \boldsymbol{\alpha}_1, a_1) + \dots + w_Mh_M(\mathbf{x}; \boldsymbol{\alpha}_M, a_M) = \mathbf{h}^T \mathbf{w} + b$$

$$h_m(\mathbf{x}; \boldsymbol{\alpha}_m, a_m) = g(a_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \dots + \alpha_{mp}x_p) = g(\mathbf{x}^T \boldsymbol{\alpha}_m + a_m)$$

$g$  is the activation function. If:

- $g$  is the identity function, NN is equivalent to linear regression  $g(v) = v$

$$g(a_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \dots + \alpha_{mp}x_p) = a_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \dots + \alpha_{mp}x_p$$

- Typically,  $g$  is used for the nonlinear transformation making the NN flexible

$$g(v) = \exp(-a(v - c)^2) \quad \text{radial basis function (RBF)}$$

**Table 1.** Overview of a selection of popular activation functions.

Names	Equation
linear function	$g(v) = v$
identity function[2]	
rectified linear unit (ReLU) [2]	$g(v) = \max(0, v)$
exponential linear unit (ELU)[5]	$g(v) = \begin{cases} v & \text{if } v \geq 0 \\ a(\exp(v) - 1) & \text{otherwise} \end{cases}$ <p>where <math>a</math> is a parameter</p>
continuously differentiable exponential linear unit (CELU)[6]	$g(v) = \begin{cases} v & \text{if } v \geq 0 \\ a \left( \exp\left(\frac{v}{a}\right) - 1 \right) & \text{otherwise} \end{cases}$ <p>where <math>a</math> is a parameter</p>
Gaussian error linear unit (GELU)[7]	$g(v) = v \cdot \frac{1}{2} \left[ 1 + \operatorname{erf}\left(\frac{v}{\sqrt{2}}\right) \right]$ <p>faster approximated versions:</p> $g(v) = 0.5v \left( 1 + \tanh \left[ \sqrt{\frac{2}{\pi}} (v + 0.044715v^3) \right] \right)$ $g(v) = v \cdot \sigma(1.702v) = v \cdot \frac{1}{1 + \exp(-1.702v)}$

P. O. Dral, A. Kananenka, F. Ge, B.-X. Xue, Neural Networks. In *Quantum Chemistry in the Age of Machine Learning*, 1st ed.; P. O. Dral, Ed. Elsevier: 2023.

Pavlo Dral



---

radial basis function

(RBF)[1-2]

$$g(v) = \exp(-a(v - c)^2)$$

where  $a$  and  $c$  are parameters

---

logistic sigmoid

function[1-2]

$$g(v) = \sigma(v) = \frac{1}{1 + \exp(-v)}$$

---

softplus function[2]

$$g(v) = \log(1 + \exp(v))$$

---

hyperbolic tangent

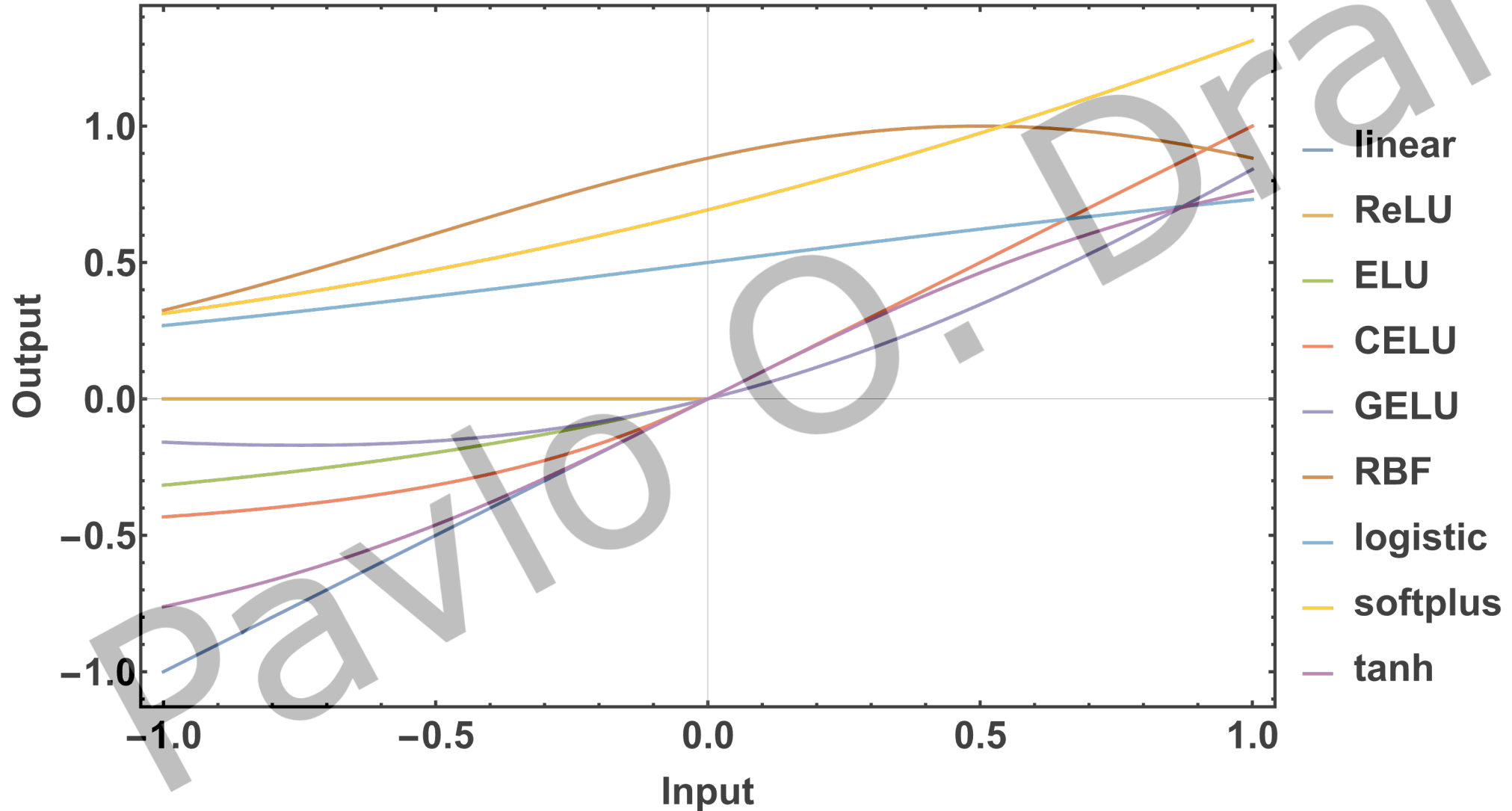
function[2]

$$g(v) = T(v) = \tanh(v)$$

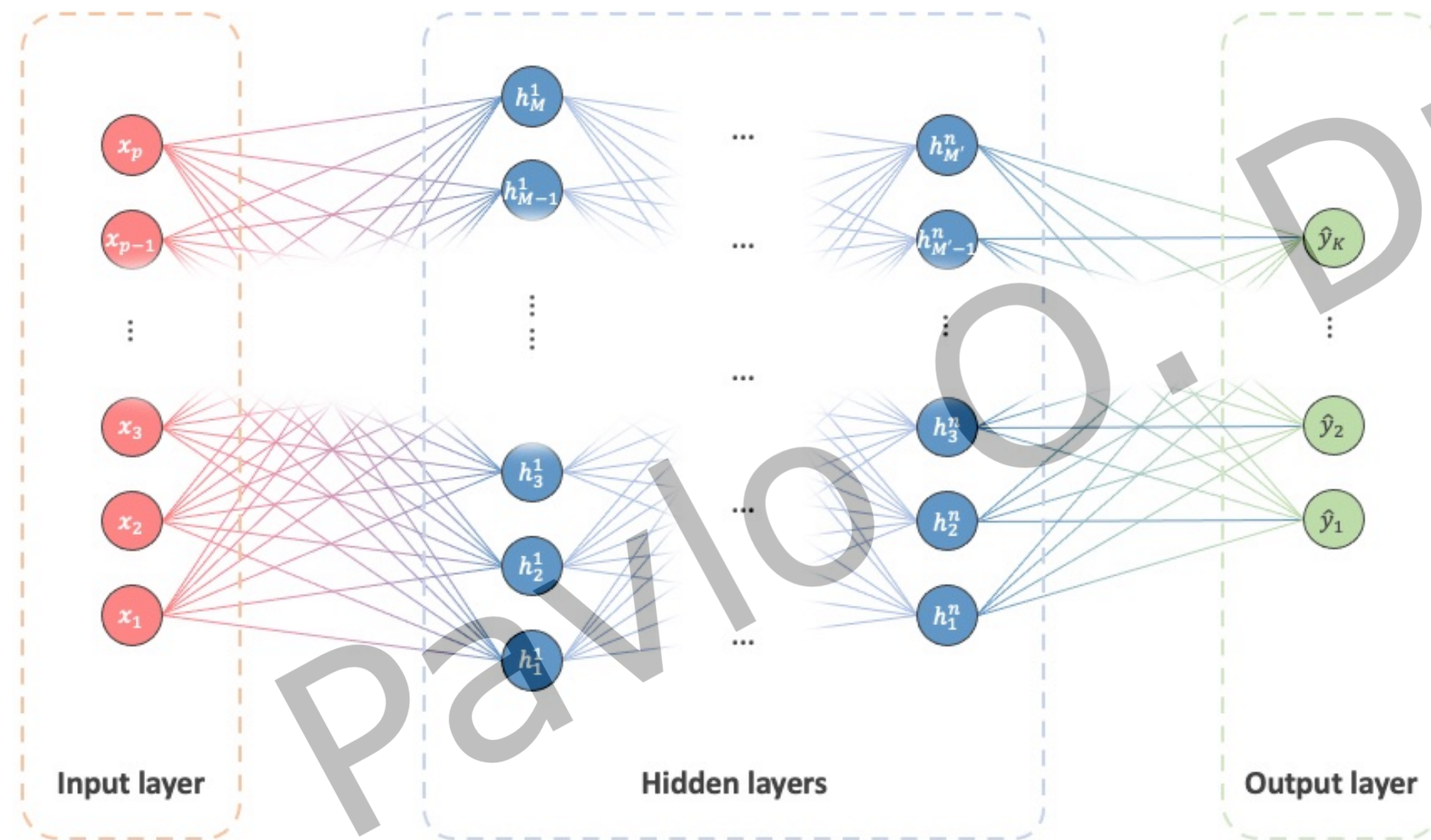
---

---

# Activation functions



# Neural network (NN)



NN consists of

- layers
- nodes=units=neurons

The single hidden layer,  
feed-forward NN

$w, \alpha \dots$  weights

$a, b \dots$  biases

$$f(\mathbf{x}; \alpha, \mathbf{a}, \mathbf{w}, b) = f^{(2)}(\mathbf{h}; \mathbf{w}, b)$$

To train NN means to find its weights  $\theta$  usually by solving this minimization task:

$$\arg \min_{\theta} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \theta) - y_i)^2$$

To avoid overfitting this solution can be regularized using weight decay approach (recall ridge regression and KRR):

$$\arg \min_{\theta} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \theta) - y_i)^2 + \lambda \sum_{j=1}^{N_p} \theta_j^2$$

$$\theta = \mathbf{w}, \alpha, \mathbf{a}, b$$

parameters

$$\hat{y} = f(\mathbf{x}; \alpha, \mathbf{a}, \mathbf{w}, b) = b + w_1 h_1(\mathbf{x}; \alpha_1, a_1) + \dots + w_M h_M(\mathbf{x}; \alpha_M, a_M) = \mathbf{h}^T \mathbf{w} + b$$

$$h_m(\mathbf{x}; \alpha_m, a_m) = g(a_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \dots + \alpha_{mp}x_p) = g(\mathbf{x}^T \alpha_m + a_m)$$

To train NN means to find its weights  $\theta$  usually by solving this minimization task:

$$\arg \min_{\theta} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \theta) - y_i)^2$$

To avoid overfitting this solution can be regularized using weight decay approach (recall ridge regression and KRR):

$$\arg \min_{\theta} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \theta) - y_i)^2 + \lambda \sum_{j=1}^{N_p} \theta_j^2$$

$$\theta = \mathbf{w}, \alpha, \mathbf{a}, b$$

parameters

$$\hat{y} = f(\mathbf{x}; \alpha, \mathbf{a}, \mathbf{w}, b) = b + w_1 h_1(\mathbf{x}; \alpha_1, a_1) + \dots + w_M h_M(\mathbf{x}; \alpha_M, a_M) = \mathbf{h}^T \mathbf{w} + b$$

$$h_m(\mathbf{x}; \alpha_m, a_m) = g(a_m + \alpha_{m1}x_1 + \alpha_{m2}x_2 + \dots + \alpha_{mp}x_p) = g(\mathbf{x}^T \alpha_m + a_m)$$

**In contrast to linear regression and kernel methods, closed solution is unknown**

Issues with NNs:

In contrast to linear regression and kernel methods, no closed solution exists

Solutions are unstable and difficult to find.

Computationally expensive optimization problem should be solved and it therefore often can be speed up by using GPUs instead of CPUs.

GPUs are however much more expensive and difficult to get and optimization is still quite slow.

One of the popular approaches for fitting is **back-propagation**.

Back-propagation:

$$L(\boldsymbol{\theta}) = \sum_{i=1}^N (f(\mathbf{x}_i; \boldsymbol{\theta}) - y_i)^2$$

gradient descent update with learning rate  $\gamma$

$$\theta_k^{(r+1)} = \theta_k^{(r)} - \gamma \frac{\partial L(\boldsymbol{\theta})}{\partial \theta_k}$$

Well parallelized:

$$L(\boldsymbol{\theta}) = \sum_{i=1}^N L_i = \sum_{i=1}^N (f(\mathbf{x}_i; \boldsymbol{\theta}) - y_i)^2$$

The training set is often split into the minibatches (**batches**)

Update of parameters after the sweep over the entire training set is called an *epoch*.

Issues with NNs:

Overfitting, regularization methods to deal with it:

- weight decay

$$\arg \min_{\theta} \sum_{i=1}^{N_{tr}} (f(\mathbf{x}_i; \theta) - y_i)^2 + \lambda \sum_{j=1}^{N_p} \theta_j^2$$

- early stopping
- data augmentation





## Issues with NNs:

- Input values should be scaled, usually standardized to center the inputs and scale them so that their standard deviation is 1 (Z-score normalization)
- It is also important to center reference data
- Number of hidden layers and units should be adjusted often by manual experimentation

Pavlo O. Dral

## Issues with NNs:

- Initial guess of weights strongly influences the final parameter values
- Starting with zero values prevents back-propagation algorithm to find better solutions
- Starting with too large values often leads to large generalization errors

Pavlo O. Dral

Issues with NNs:

- Initial guess of weights strongly influences the final parameter values
- Starting with zero values prevents back-propagation algorithm to find better solutions
- Starting with too large values often leads to large generalization errors

Thus, one can get lot of different NNs fitted on the same data!

One can exploit this:

- Take average of multiple NNs to get more stable prediction
- Use deviation between NN predictions to estimate prediction uncertainty (e.g. useful in active learning)

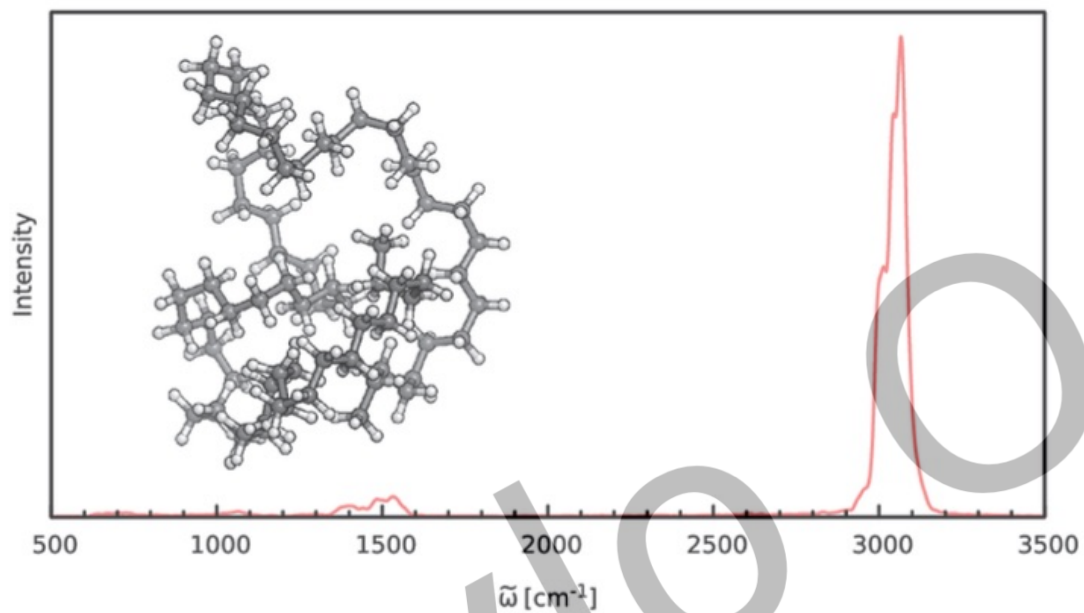
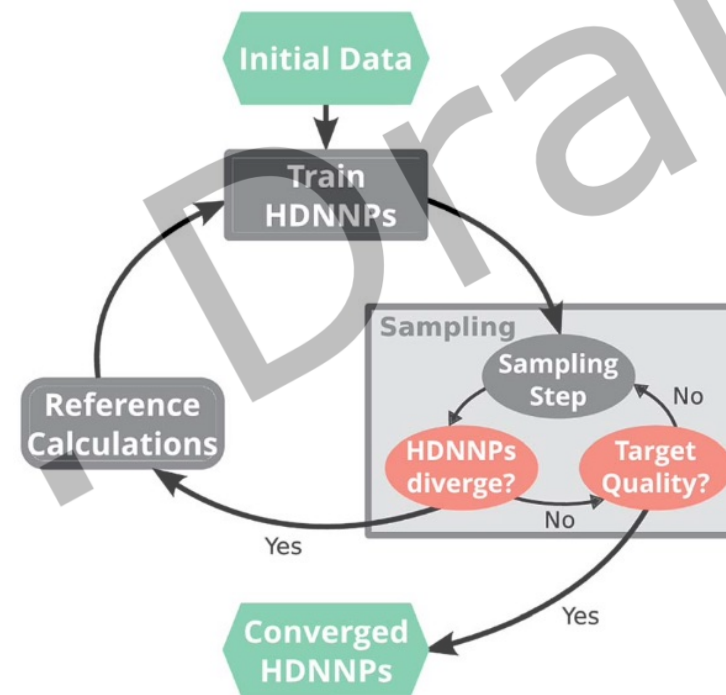


Fig. 6 IR spectrum of the  $C_{69}H_{140}$  alkane as predicted by the ML model based on the B2PLYP method.

















Deep learning is based on neural networks (NN) with large depth (for feed-forward neural network – more than one hidden unit) in contrast to shallow neural network

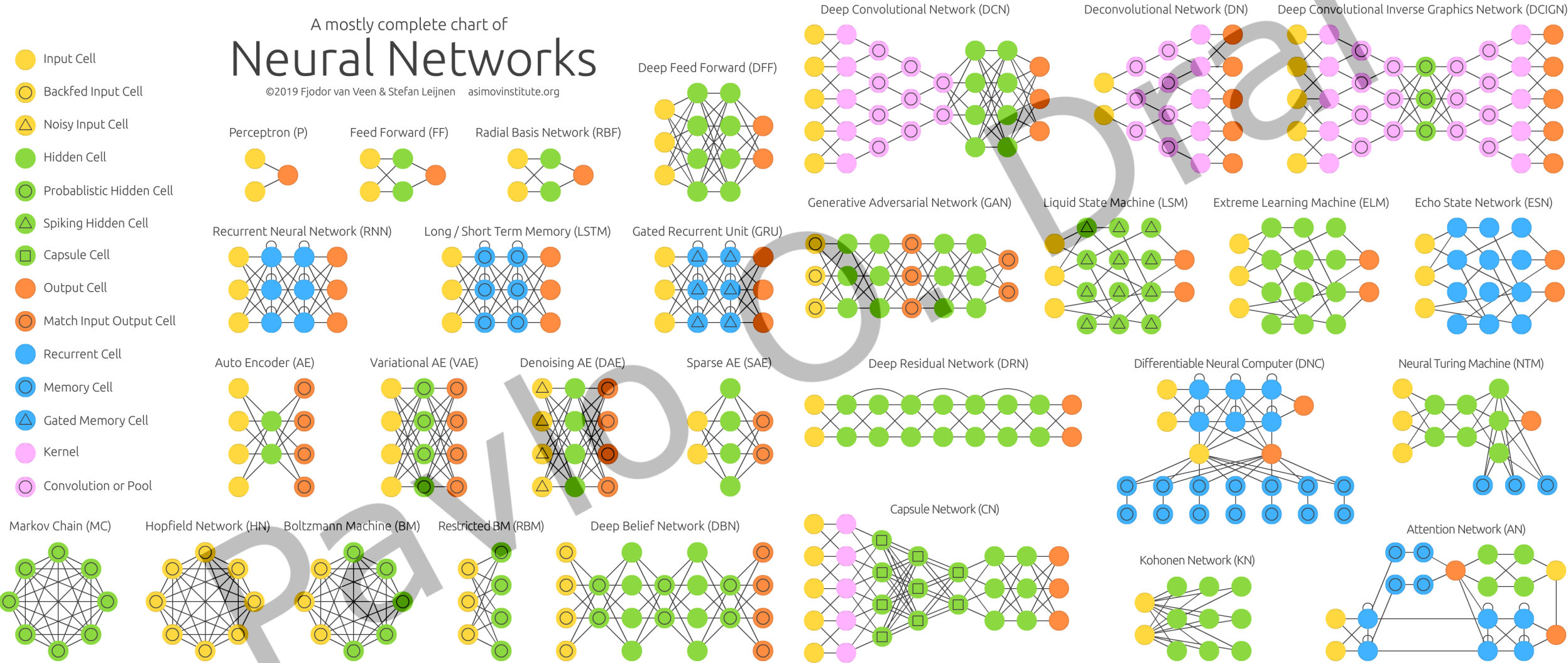
Some of other types of neural networks:

- Convolutional networks
- Recurrent neural networks
- Autoencoders

## A mostly complete chart of Neural Networks

©2019 Fjodor van Veen & Stefan Leijnen asimovinstitute.org

-  Input Cell
-  Backfed Input Cell
-  Noisy Input Cell
-  Hidden Cell
-  Probabilistic Hidden Cell
-  Spiking Hidden Cell
-  Capsule Cell
-  Output Cell
-  Match Input Output Cell
-  Recurrent Cell
-  Memory Cell
-  Gated Memory Cell
-  Kernel
-  Convolution or Pool



***Parametric vs nonparametric  
algorithms***

Pavlo O. Dral

**What method to choose?  
Kernel methods or neural networks?**

Pavlo O. Dral



$f(x; \text{parameters})$

Linear regression

$$f(\mathbf{x}_i; \boldsymbol{\beta}) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots$$

Number of parameters is fixed: parametric model

Neural networks are also parametric models

Kernel ridge regression (KRR)

$$f(\mathbf{x}_i; \mathbf{p}) = \sum_{j=1}^{N_{\text{tr}}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j; \mathbf{b})$$

Number of parameters depends on number of training points:  
nonparametric model, e.g. KRR

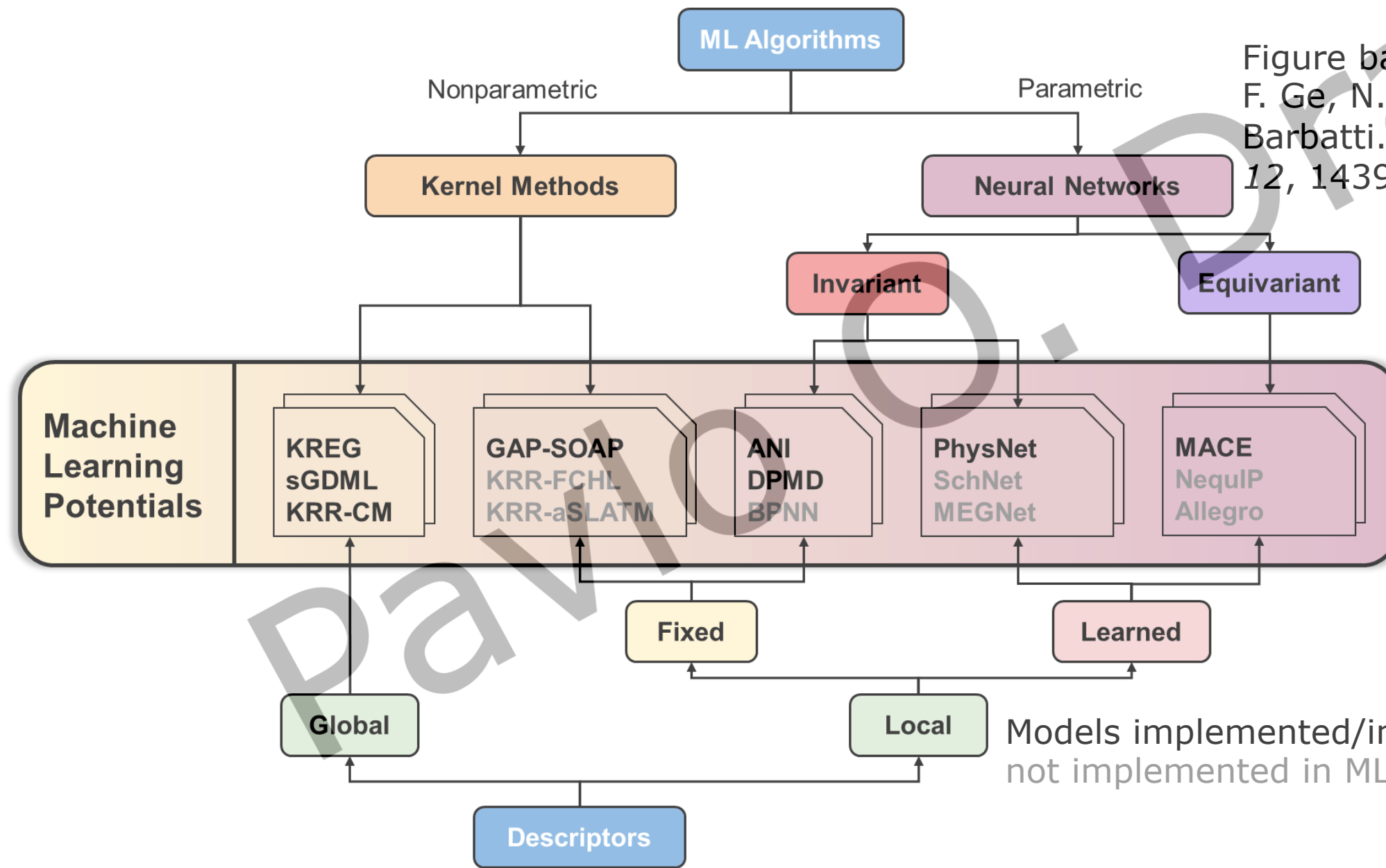
All have some advantages and disadvantages, but often provide results with similar accuracy.

In many cases **it is not possible to claim that one fitting method is better than another.** The **choice will depend on experience and taste.**[1]

However: You should be aware of the **law of the hammer** and do not try to use a hammer for every problem only because you already have a hammer.



Figure based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, *12*, 14396–14413



Models implemented/interfaced in MLatom  
not implemented in MLatom

# Supervised Machine Learning

$$E = f(\mathbf{R})$$

Input (x)  $\rightarrow$   $f(x)$   $\rightarrow$  Output (y)

Given collection of known  $\{x,y\}$  find a function  $f(x)$

training set

train

ML model

Use this function for making new predictions given just  $\{x'\}$

- Data
- **Choice of x (descriptor)**
- Choice of y (labels)
- Fitting function (ML algorithm, ML model)
- Optimization of ML model parameters

$$f(\mathbf{x}_i) = \sum_{j=1}^{N_{\text{tr}}} \alpha_j k(\mathbf{x}_i, \mathbf{x}_j) \quad k(\mathbf{x}_i, \mathbf{x}_j) = \exp\left(-\frac{1}{2\sigma^2} \sum_s^{N_x} (x_{i,s} - x_{j,s})^2\right) \quad x = \left(\dots \frac{Req}{R} \dots\right)^T$$

the Gaussian kernel function  
( $\sigma$  is the kernel width)

RE descriptor

Analytical solution for the regression coefficients  $\alpha$  given  $N_{\text{tr}}$  training points

$$\begin{pmatrix} k(\mathbf{x}_1, \mathbf{x}_1) + \lambda & \dots & k(\mathbf{x}_1, \mathbf{x}_{N_{\text{tr}}}) \\ \vdots & \ddots & \vdots \\ k(\mathbf{x}_{N_{\text{tr}}}, \mathbf{x}_1) & \dots & k(\mathbf{x}_{N_{\text{tr}}}, \mathbf{x}_{N_{\text{tr}}}) + \lambda \end{pmatrix} \begin{pmatrix} \alpha_1 \\ \vdots \\ \alpha_{N_{\text{tr}}} \end{pmatrix} = \begin{pmatrix} y_1 \\ \vdots \\ y_{N_{\text{tr}}} \end{pmatrix}$$

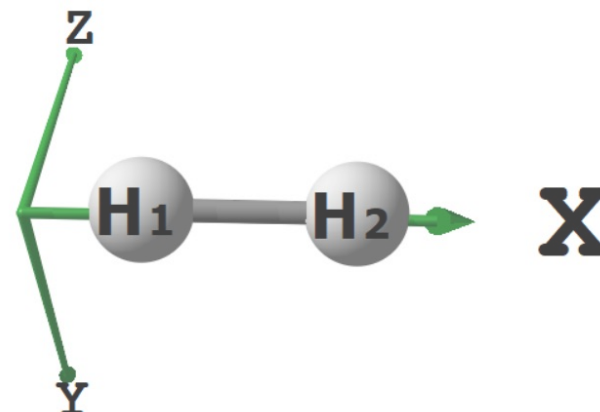
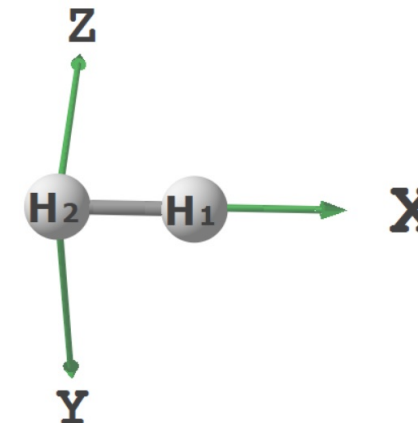
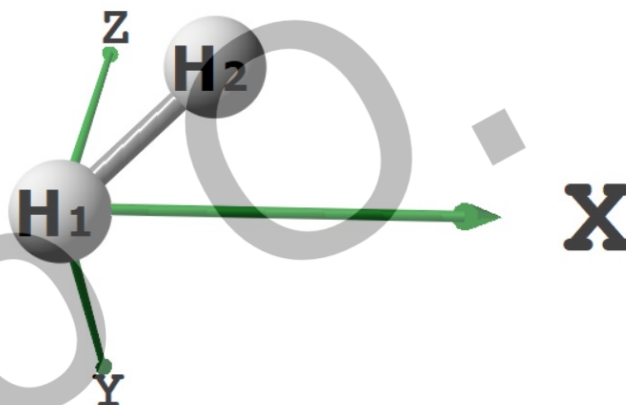
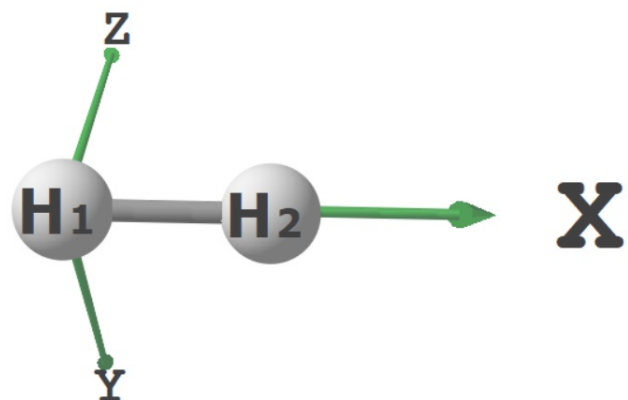
$\lambda$  is the regularization parameter ensuring transferability

$$(\mathbf{K} + \lambda \mathbf{I})\alpha = \mathbf{y}$$

# Requirements to MLPs

- rotationally and translationally invariant (or **equivariant!**)

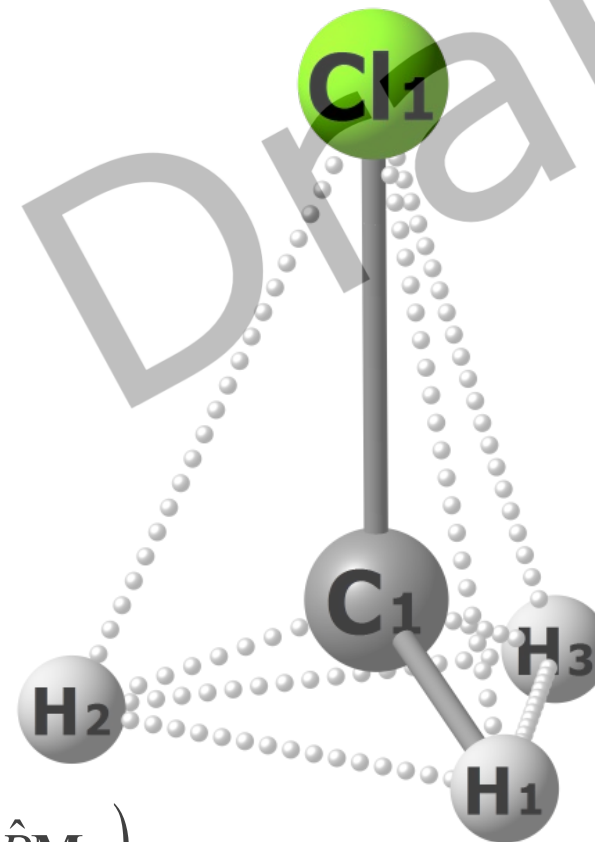
- same-element atom index invariant



# Molecular Descriptor for $\text{CH}_3\text{Cl}$

$$\mathbf{M} = \begin{pmatrix} r_{C-Cl}^{eq} \\ r_{C-Cl} \\ r_{C-H1}^{eq} \\ r_{C-H1} \\ r_{Cl-H1}^{eq} \\ r_{Cl-H1} \\ r_{H1-H2}^{eq} \\ r_{H1-H2} \\ \dots \end{pmatrix}$$

- Sort atoms, e.g. hydrogens can be sorted by their nuclear repulsion
- Normalized permutationally invariant kernel



$$\overline{K}(\mathbf{M}_i, \mathbf{M}_j) = \frac{\sum_{\hat{P}}^{N_{perm}} K(\mathbf{M}_i, \hat{P}\mathbf{M}_j)}{\sqrt{\sum_{\hat{P}}^{N_{perm}} K(\mathbf{M}_i, \hat{P}\mathbf{M}_i)} \sqrt{\sum_{\hat{P}}^{N_{perm}} K(\mathbf{M}_j, \hat{P}\mathbf{M}_j)}} \quad [2]$$

[1] Dral, Owens, Yurchenko, Thiel, *J. Chem. Phys.* **2017**, *146*, 244108

[2] Bartók, Csányi, *Int. J. Quantum Chem.* **2015**, *115*, 1051



Behler–Parrinello  
GAP-SOAP

**ANI**

PhysNet

SchNet

FCHL

aSLATM

...

Overview & benchmark in:

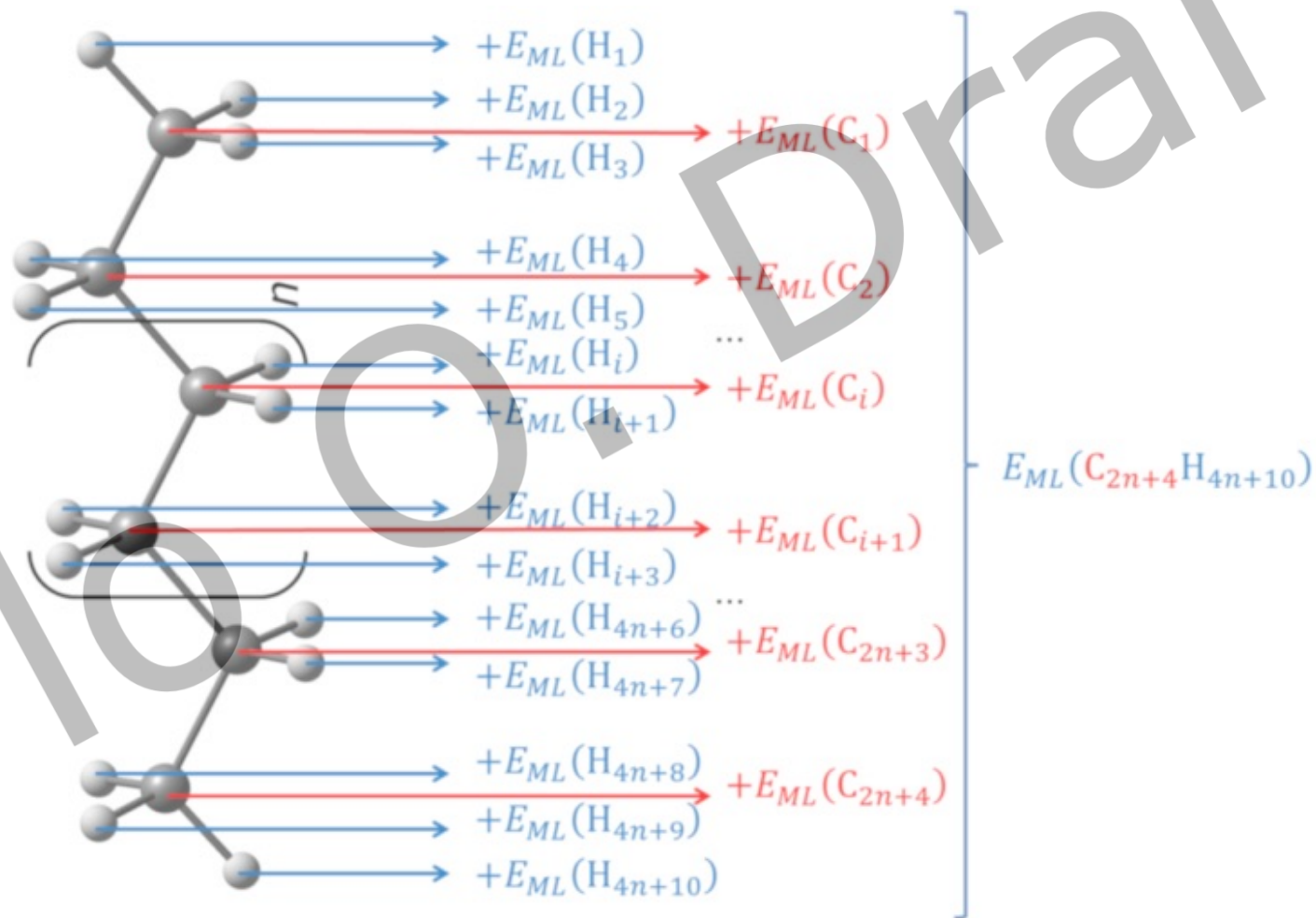
M. Pinheiro Jr, F. Ge,

N. Ferré, P. O. Dral,

M. Barbatti.

*Chem. Sci.* **2021**, *12*,

14396–14413



Approach: Behler, Parrinello, *Phys. Rev. Lett.* **2007**, *98*, 146401

ANI environment vectors (AEVs) consist of radial and angular atomic terms. Each element has its own subAEV:

$$G_k^R = \sum_{j \neq i} e^{-\eta(R_{ij} - R_s^{(k)})^2} f_c(R_{ij})$$

$$G_{p,q}^A = 2^{1-\zeta} \sum_{j,k \neq i} \left(1 + \cos(\theta_{ijk} - \theta_s^{(q)})\right)^\zeta e^{-\eta\left(\frac{R_{ij} + R_{ik}}{2} - R_s^{(p)}\right)^2} f_c(R_{ij}) f_c(R_{ik}).$$

the cutoff function

$$f_{\text{cut}}(r) = \begin{cases} 1, & r \leq r_{\text{cut}} - r_{\Delta}, \\ \frac{1}{2} \left( \cos\left(\pi \frac{r - r_{\text{cut}} + r_{\Delta}}{r_{\Delta}}\right) + 1 \right), & r_{\text{cut}} - r_{\Delta} < r \leq r_{\text{cut}}, \\ 0, & r > r_{\text{cut}}, \end{cases}$$

Gaussian approximation potential (GAP)[1] with Smooth Overlap of Atomic Positions (SOAP)[2] descriptor

the atomic neighborhood density

$$\rho_i(\mathbf{r}) = \sum_j \exp\left(-\frac{|\mathbf{r} - \mathbf{r}_{ij}|^2}{2\sigma_{atom}^2}\right) f_{cut}(|\mathbf{r}_{ij}|),$$

the cutoff function

$$f_{cut}(r) = \begin{cases} 1, & r \leq r_{cut} - r_{\Delta}, \\ \frac{1}{2} \left( \cos\left(\pi \frac{r - r_{cut} + r_{\Delta}}{r_{\Delta}}\right) + 1 \right), & r_{cut} - r_{\Delta} < r \leq r_{cut}, \\ 0, & r > r_{cut}, \end{cases}$$

[1] A. P. Bartók, M. C. Payne, R. Kondor, G. Csányi, Phys. Rev. Lett. **2010**, 104, 136403

[2] A. P. Bartók, R. Kondor, G. Csányi, Phys. Rev. B **2013**, 87, 187115

PhysNet is using message-passing NN and so called 'learned' local descriptors

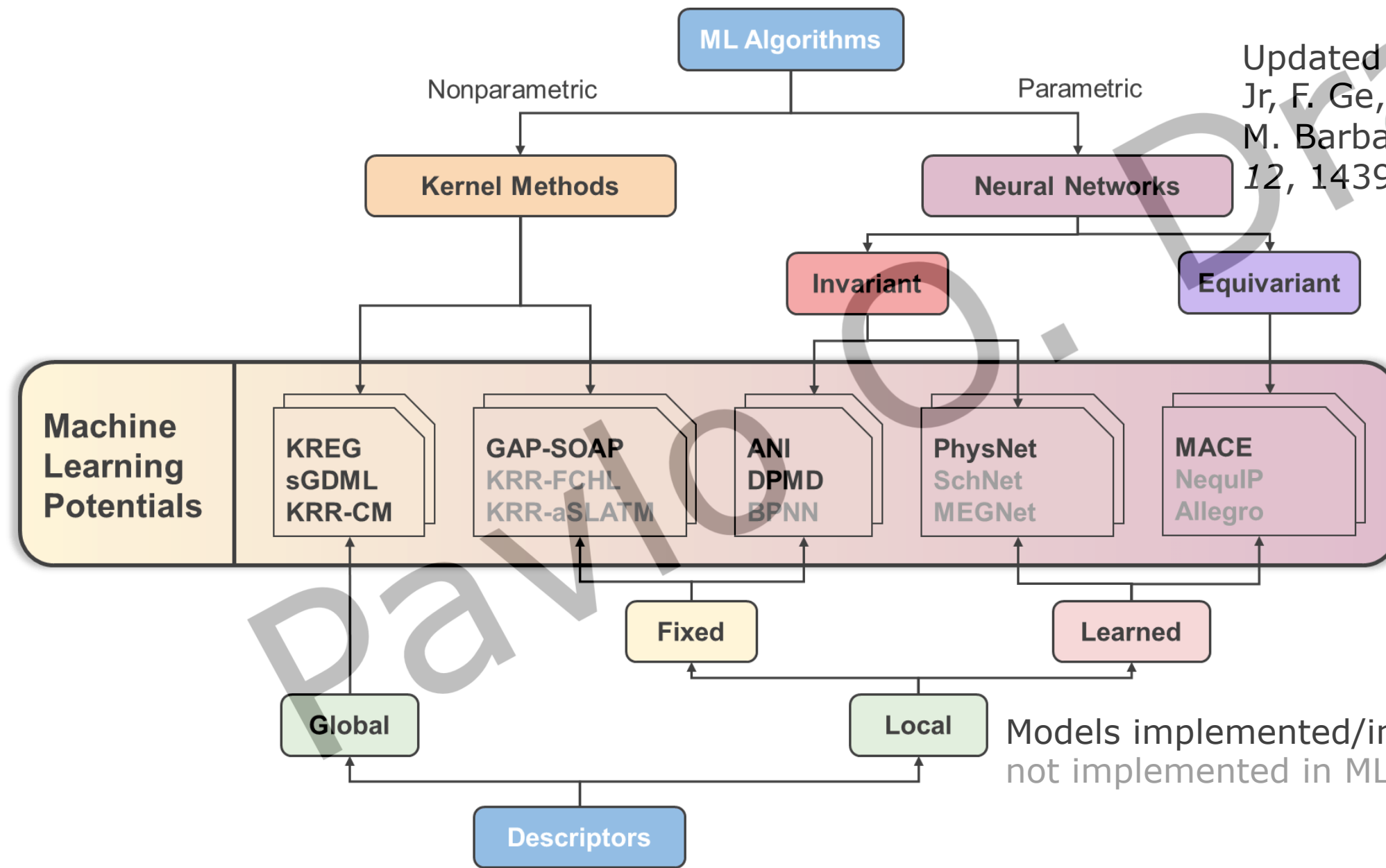
the embedding vector

$$\mathbf{x}_i^0 = \mathbf{e}_{z_i}$$

the coordinates are transformed to

$$g_k(r_{ij}) = f_c(r_{ij}) \cdot e^{-\beta_k(e^{-r_{ij}} - \mu_k)^2}$$

Updated based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, 12, 14396–14413





## Certificate of appreciation Highly cited article

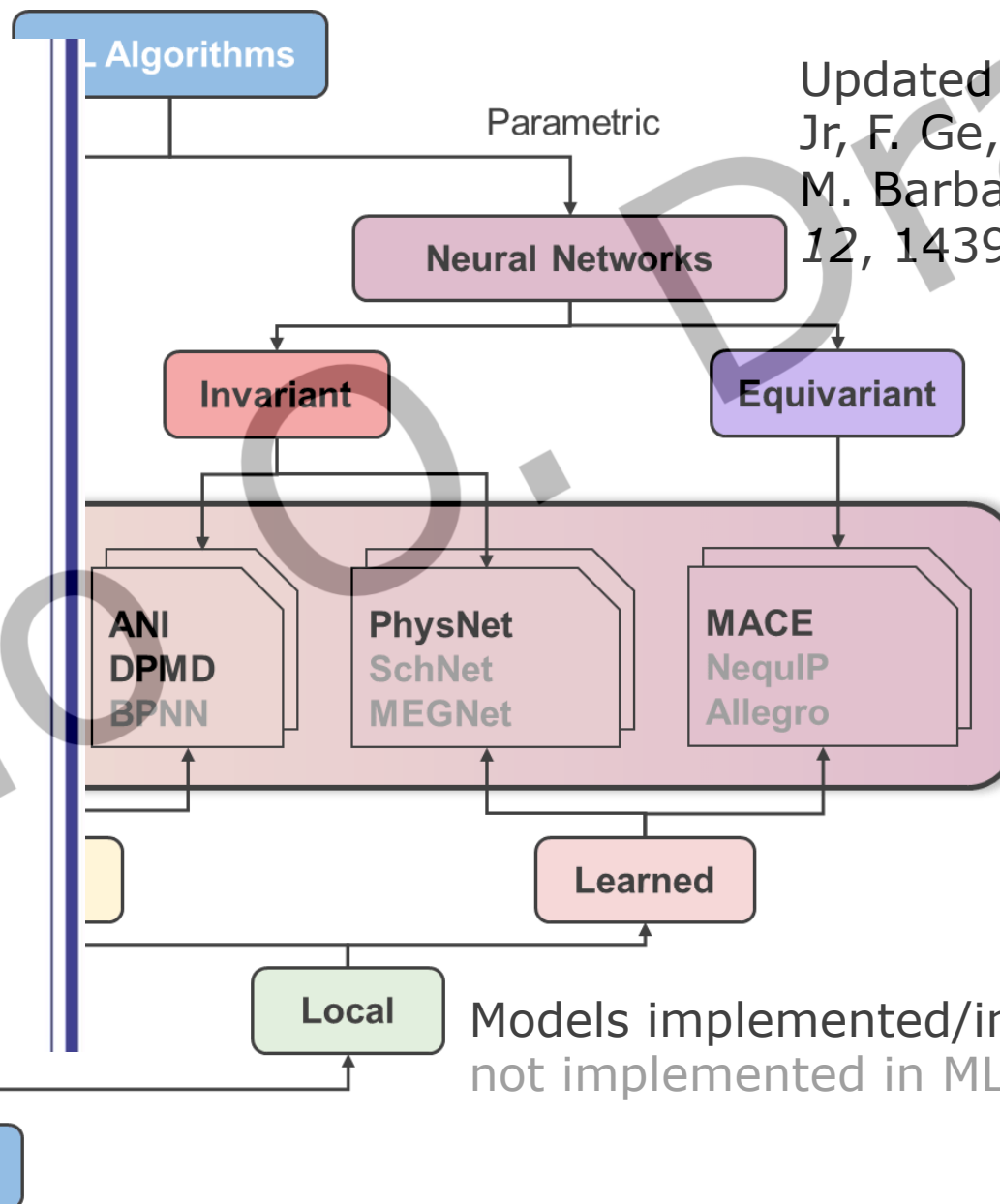
This certificate is awarded to

*Choosing the right molecular machine learning potential*

Max Pinheiro, Fuchun Ge, Nicolas Ferré, Pavlo O. Dral and Mario Barbatti

*Chemical Science*, 2021, 12, 43, 14396

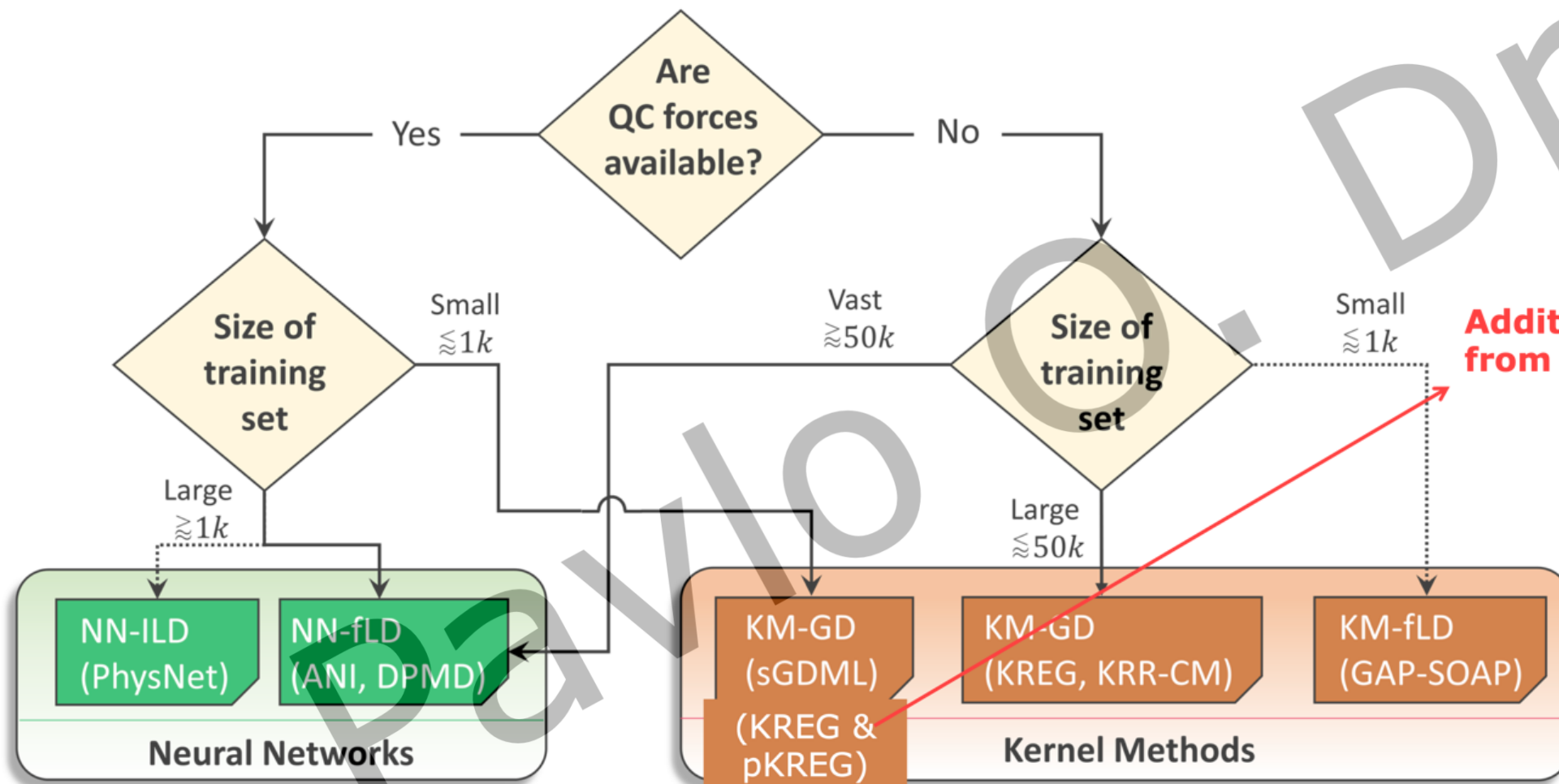
For publishing research in the top 5% of highly cited works  
from the Royal Society of Chemistry Journals



Updated based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, 12, 14396–14413

Models implemented/interfaced in **MLatom**  
not implemented in MLatom

## Recommendations from 2021

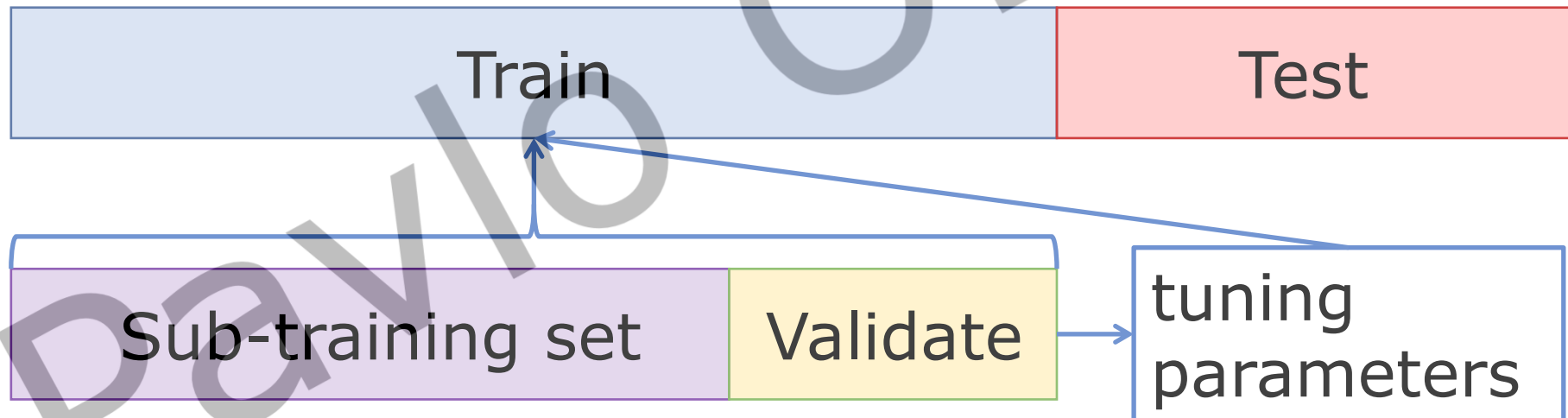


**Addition from 2023**

KREG & pKREG:  
Y.-F. Hou, F. Ge,  
P. O. Dral. *J. Chem. Theory Comput.* **2023**, *19*, 2369–2379

**After 2021, open questions such as: where is the place for equivariant NNs?**

- Often ML error for its own training set is close to zero
- Using errors in the validation set would be also incorrect, because their minimization is a part of the training process
- We should estimate errors on a **completely independent test set**



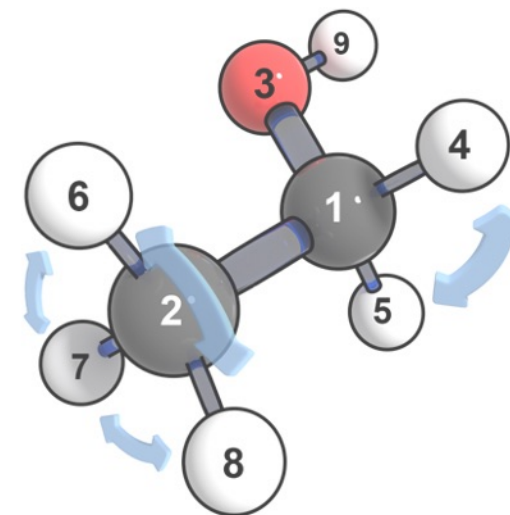
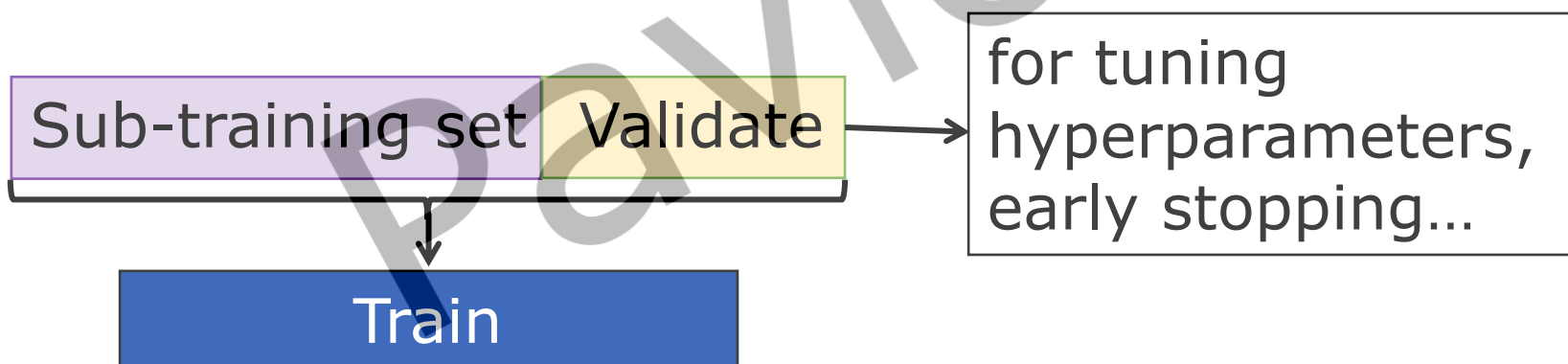


**KRR-CM** – kernel ridge regression with Gaussian kernel and Coulomb matrix descriptor

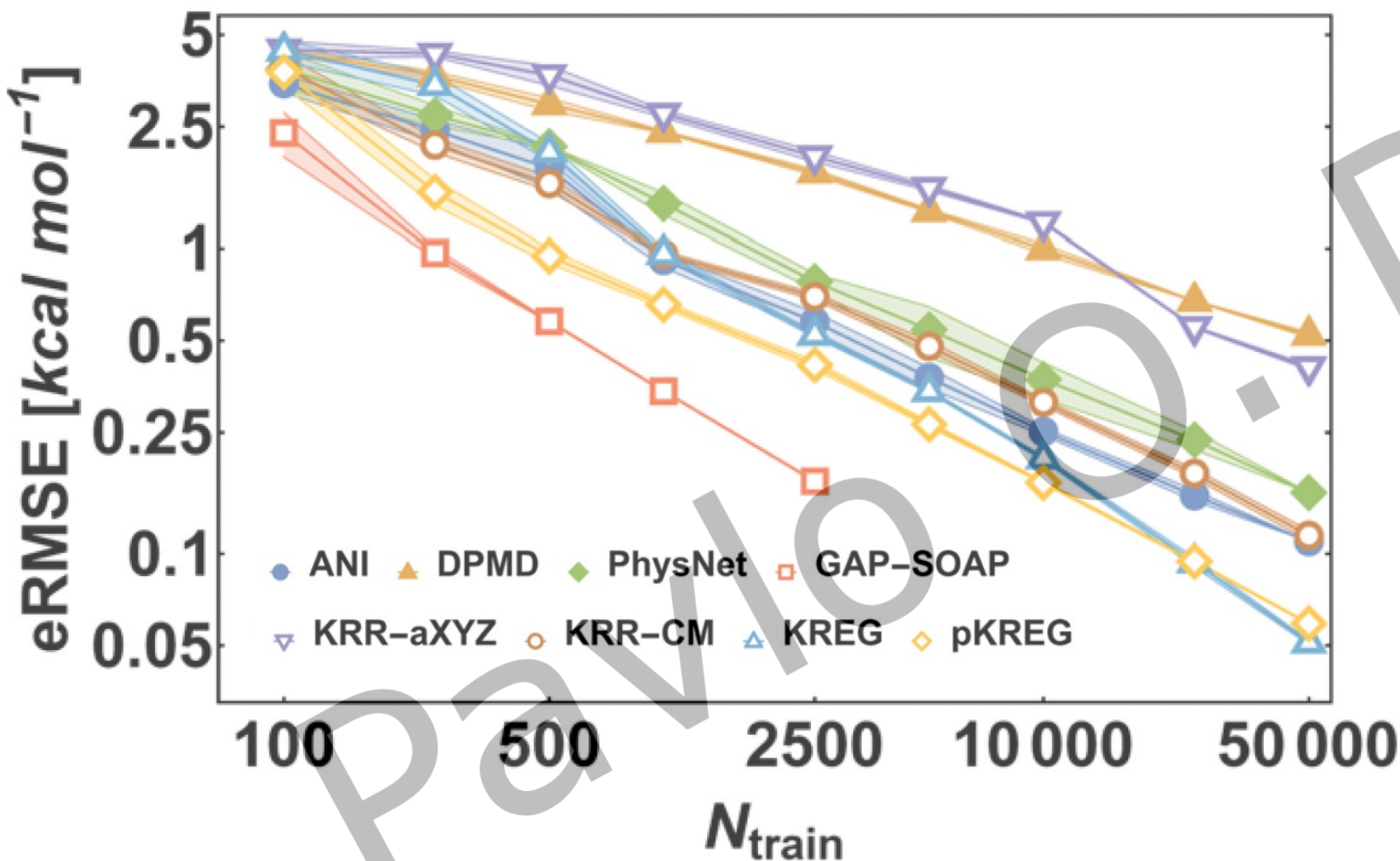
Test set RMSEs in kcal/mol

Number of training points	KRR-CM	KREG
<b>100</b>	<b>3.90±0.41</b>	<b>4.45±0.36</b>
<b>2500</b>	<b>0.70±0.02</b>	<b>0.52±0.01</b>

What do we mean by '100 training points?' Is the validation set included?  
Let's use the term '**sub-training set**'!



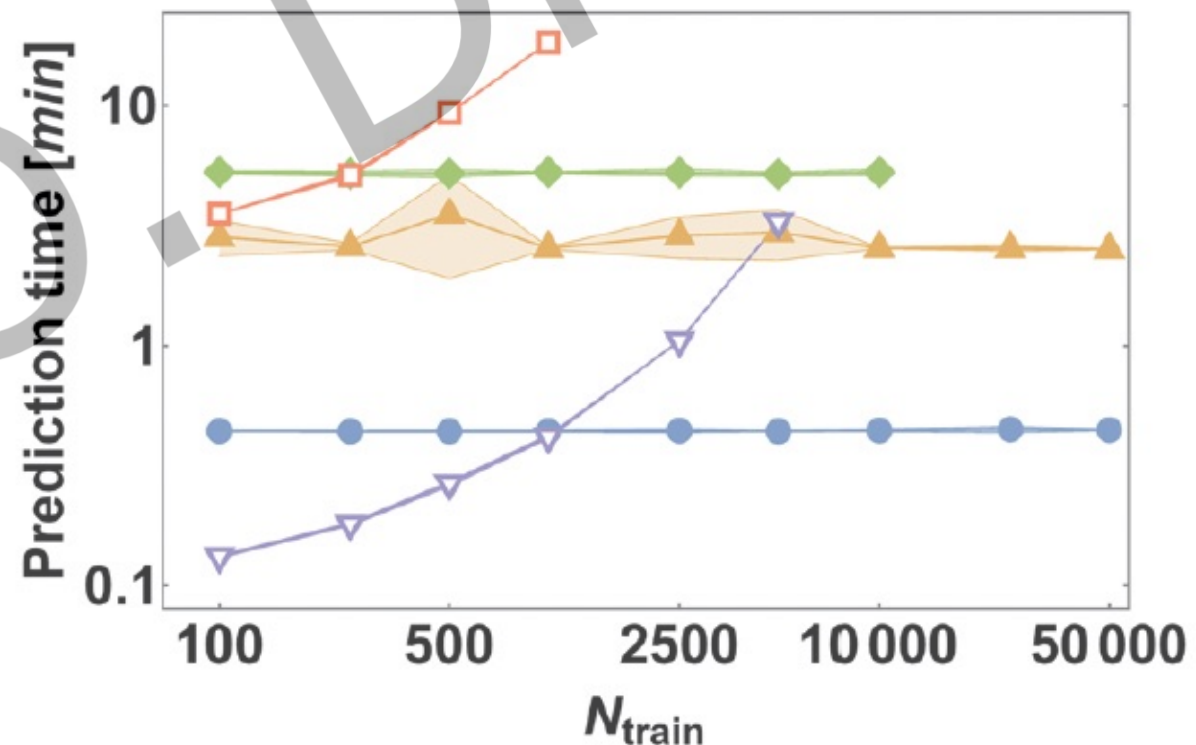
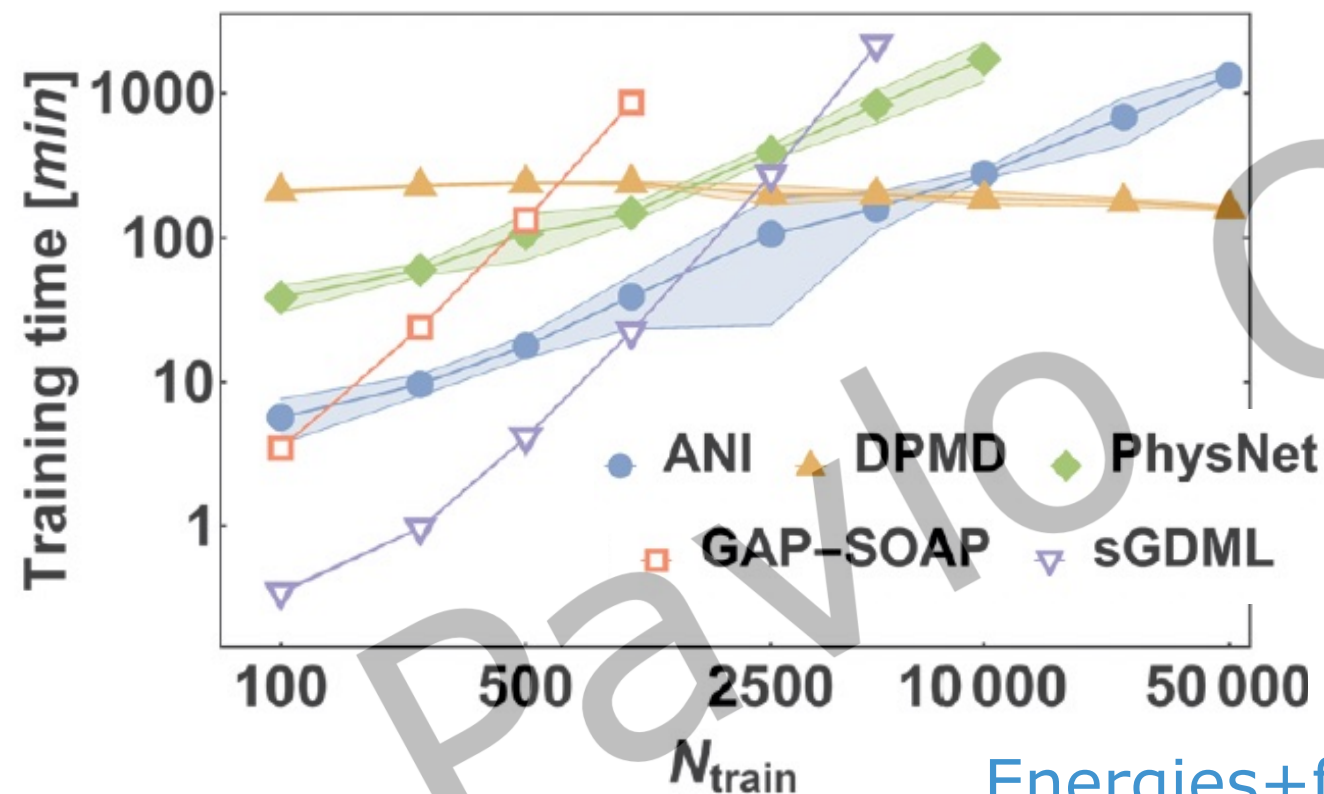
# Learning curves (energies-only)



$$\varepsilon = \varepsilon_a + \frac{a}{N_{tr}^b}$$

$$\log(\varepsilon) \approx \log(a) - b \log N_{tr}$$

For small training sets kernel methods (open markers) are often both more accurate and faster for training and prediction than neural networks (filled markers)



Energies+forces

The same hardware!

# Supervised Machine Learning

Input (x) → f(x) → Output (y)

$$E = f(\mathbf{R})$$

$$F_{A,d} = -\frac{\partial E}{\partial x_{A,d}}$$

Given collection of **known {x,y}** find a **function f(x)**  
 training set                      train                      ML model

Use this function for making new predictions given just {x'}

- Data
- Choice of x (descriptor)
- **Choice of y (labels)**
- Fitting function (ML algorithm, ML model)
- Optimization of ML model parameters

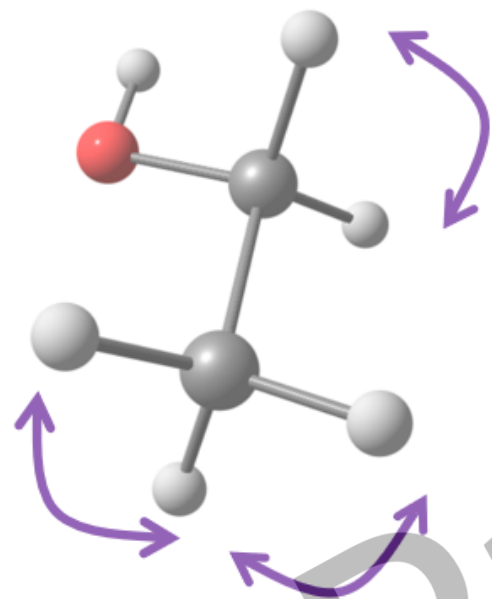
$$L = \omega_E L_E + \omega_F L_F$$

$$L = \omega_E \sum_{j=1}^{N_{\text{tr}}} (f(\mathbf{x}_j; \boldsymbol{\alpha}) - y_j)^2 + \omega_F \sum_{j=1}^{N_{\text{tr}}} \sum_{d=1}^p \left( \frac{\partial f(\mathbf{x}_j; \boldsymbol{\alpha})}{\partial x_{j,d}} - \frac{\partial y_j}{\partial x_{j,d}} \right)^2$$

$$f(\mathbf{x}') = \sum_{i=1}^{N_{\text{tr}}} \alpha_j k(\mathbf{x}_i, \mathbf{x}') + \sum_{i=1}^{N_{\text{tr}}} \sum_{d=1}^p \alpha_{id}^{\text{drv}} \frac{\partial k(\mathbf{x}_i, \mathbf{x}')}{\partial x_{i,d}}$$

$$\frac{\partial f(\mathbf{x}')}{\partial x'_t} = \sum_{i=1}^{N_{\text{tr}}} \alpha_i \frac{\partial k(\mathbf{x}_i, \mathbf{x}')}{\partial x'_t} + \sum_{i=1}^{N_{\text{tr}}} \sum_{d=1}^p \alpha_{id}^{\text{drv}} \frac{\partial^2 k(\mathbf{x}_i, \mathbf{x}')}{\partial x_{i,d} \partial x'_t}$$

**(p)KREG**



**Permutations**

$$\begin{pmatrix} \mathbf{K} + \lambda_{\mathbf{v}} & \frac{\partial \mathbf{K}}{\partial \mathbf{M}} \\ \frac{\partial \mathbf{K}}{\partial \mathbf{M}} & \frac{\partial^2 \mathbf{K}}{\partial \mathbf{M}^2} + \lambda_{\text{gxyz}} \end{pmatrix} \boldsymbol{\alpha} = \begin{pmatrix} \mathbf{y} - \mathbf{y}_{\text{prior}} \\ \frac{\partial \mathbf{y}}{\partial \mathbf{M}} \end{pmatrix}$$

**RE descriptor**

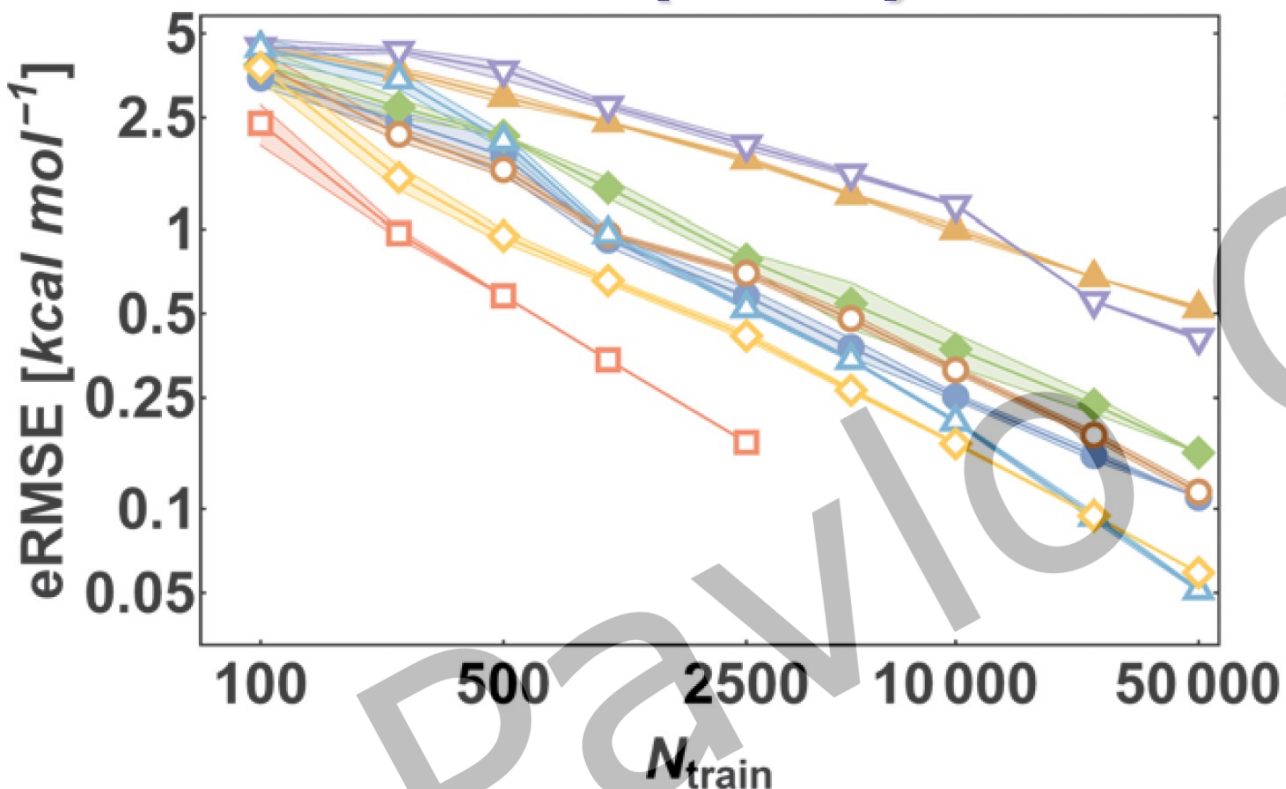
$$\mathbf{x}^T = [\dots r_{a,b \neq a}^{\text{ref}} / r_{a,b \neq a} \dots]$$

**Gaussian kernel**

$$k(\mathbf{x}_i, \mathbf{x}_j) = \exp\left(-\|\mathbf{x}_i - \mathbf{x}_j\|^2 / 2\sigma^2\right)$$

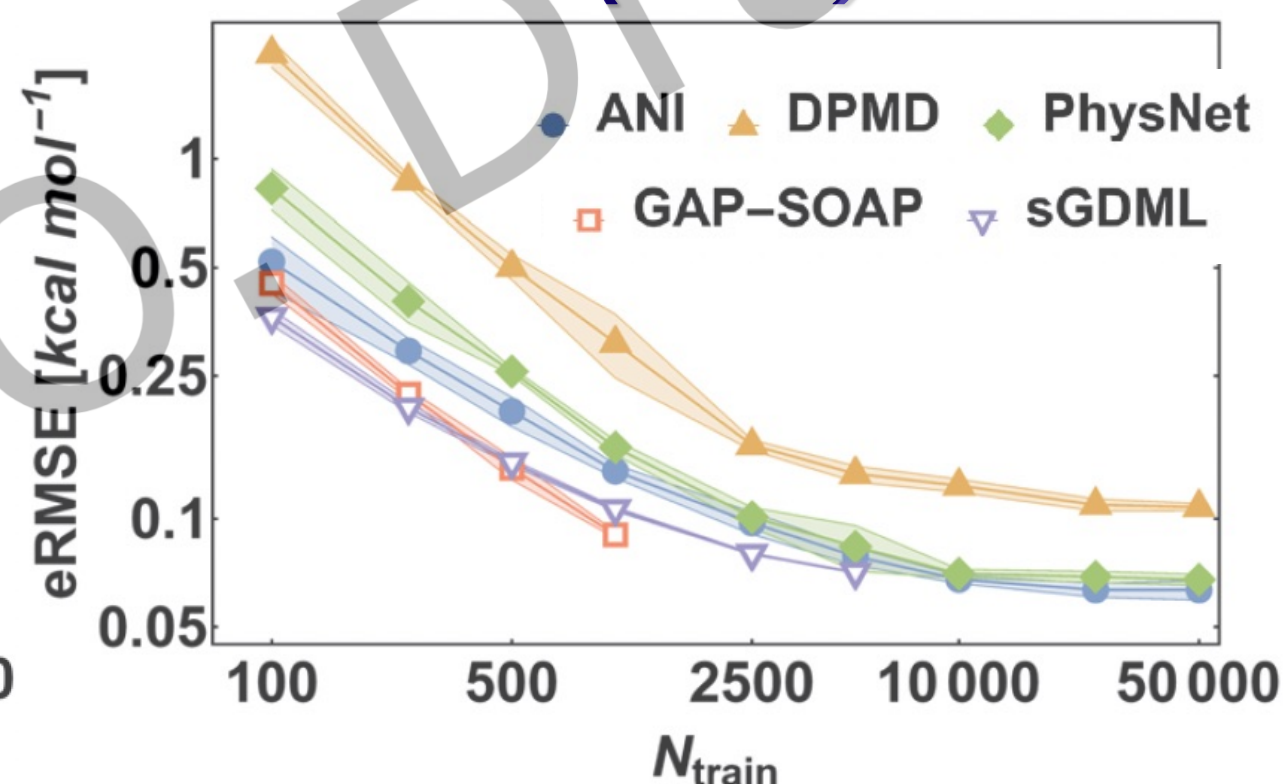
## Energies-only

### Ethanol (MD17)



## Energies+gradients

### Ethanol (MD17)



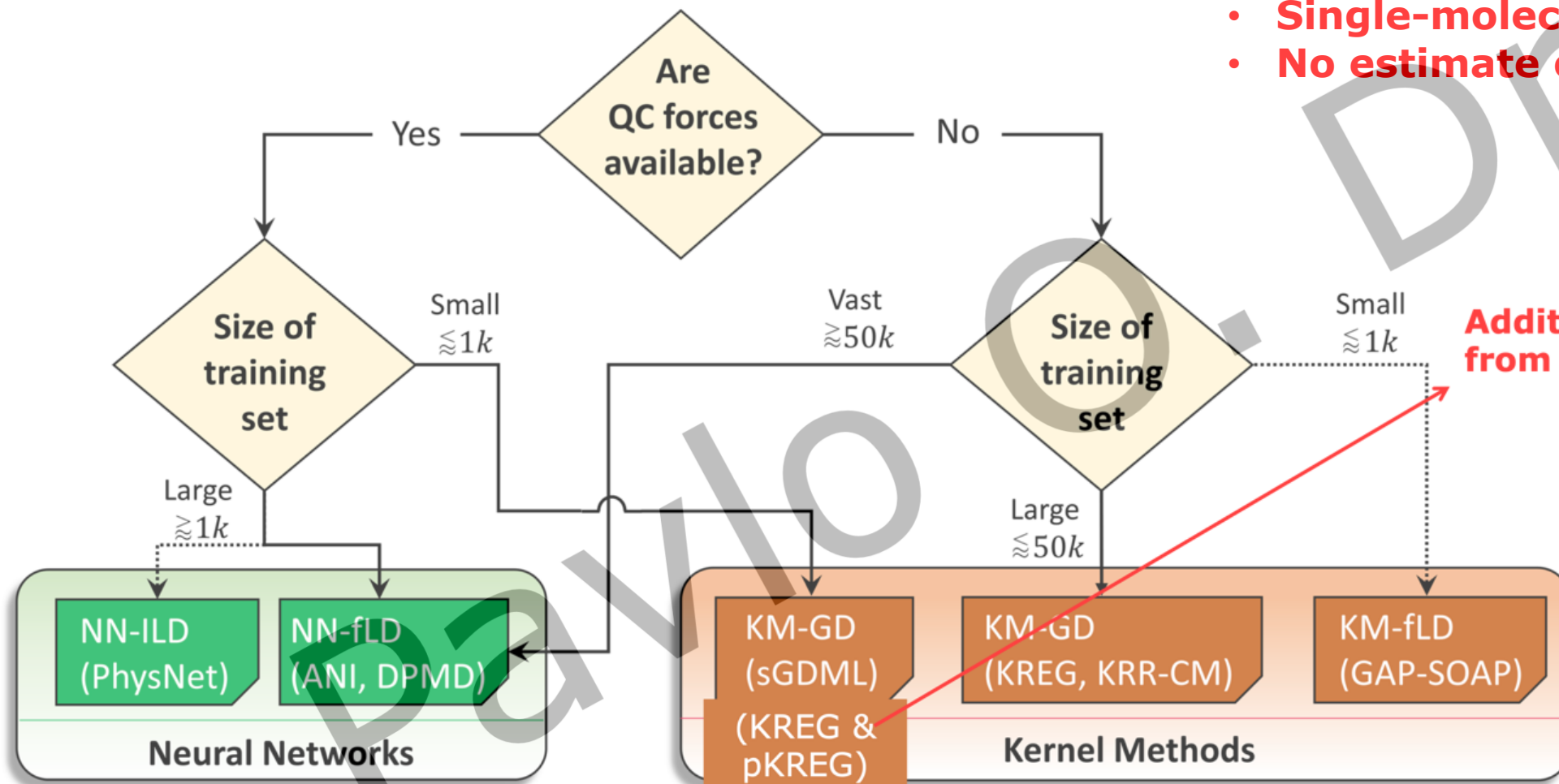
● ANI ▲ DPMD ◆ PhysNet □ GAP-SOAP  
▼ KRR-aXYZ ○ KRR-CM ▲ KREG ◇ pKREG



## Recommendations from 2021

## Limitations:

- **Single-molecule PES**
- **No estimate of equivariant MLPs**

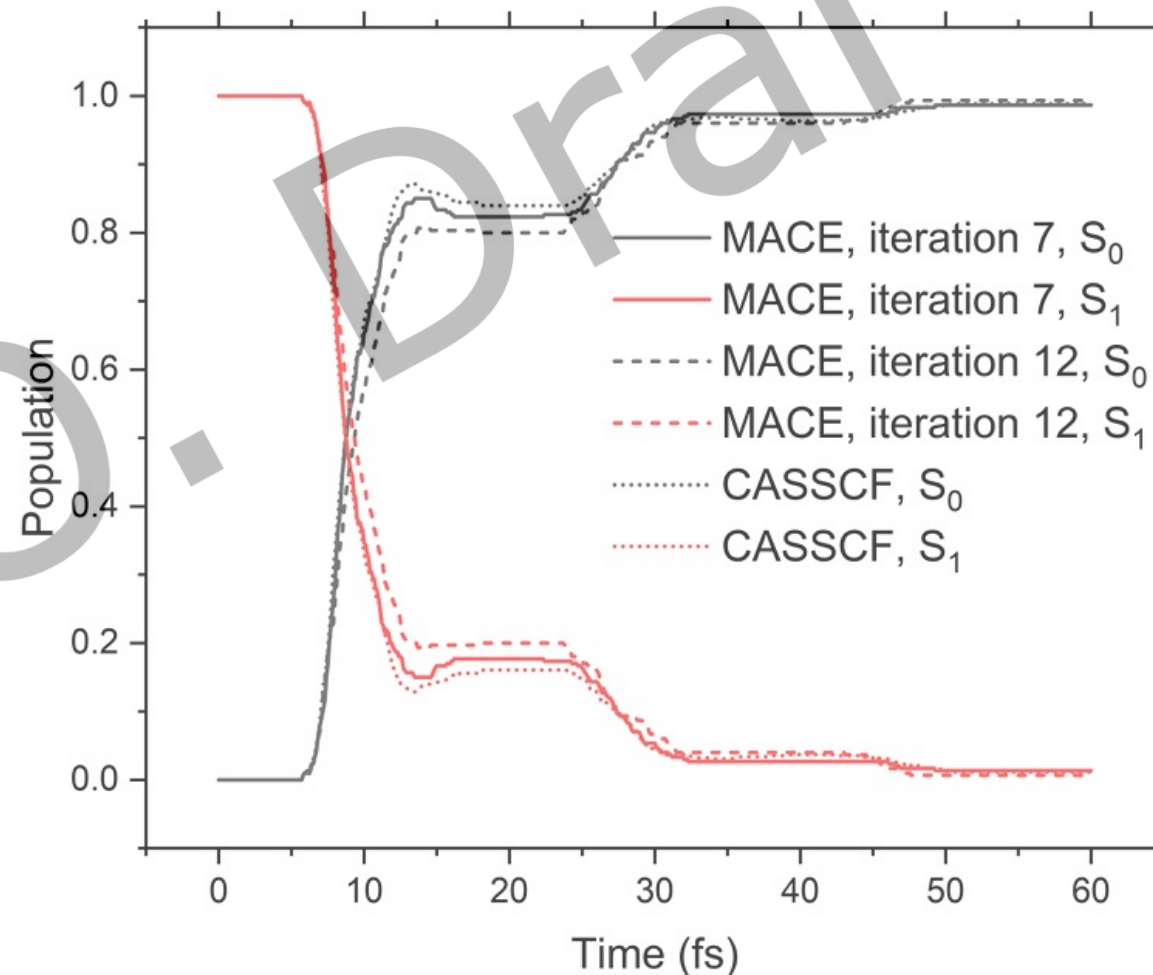


KREG & pKREG:  
Y.-F. Hou, F. Ge,  
P. O. Dral. *J. Chem. Theory Comput.* **2023**, *19*, 2369–2379

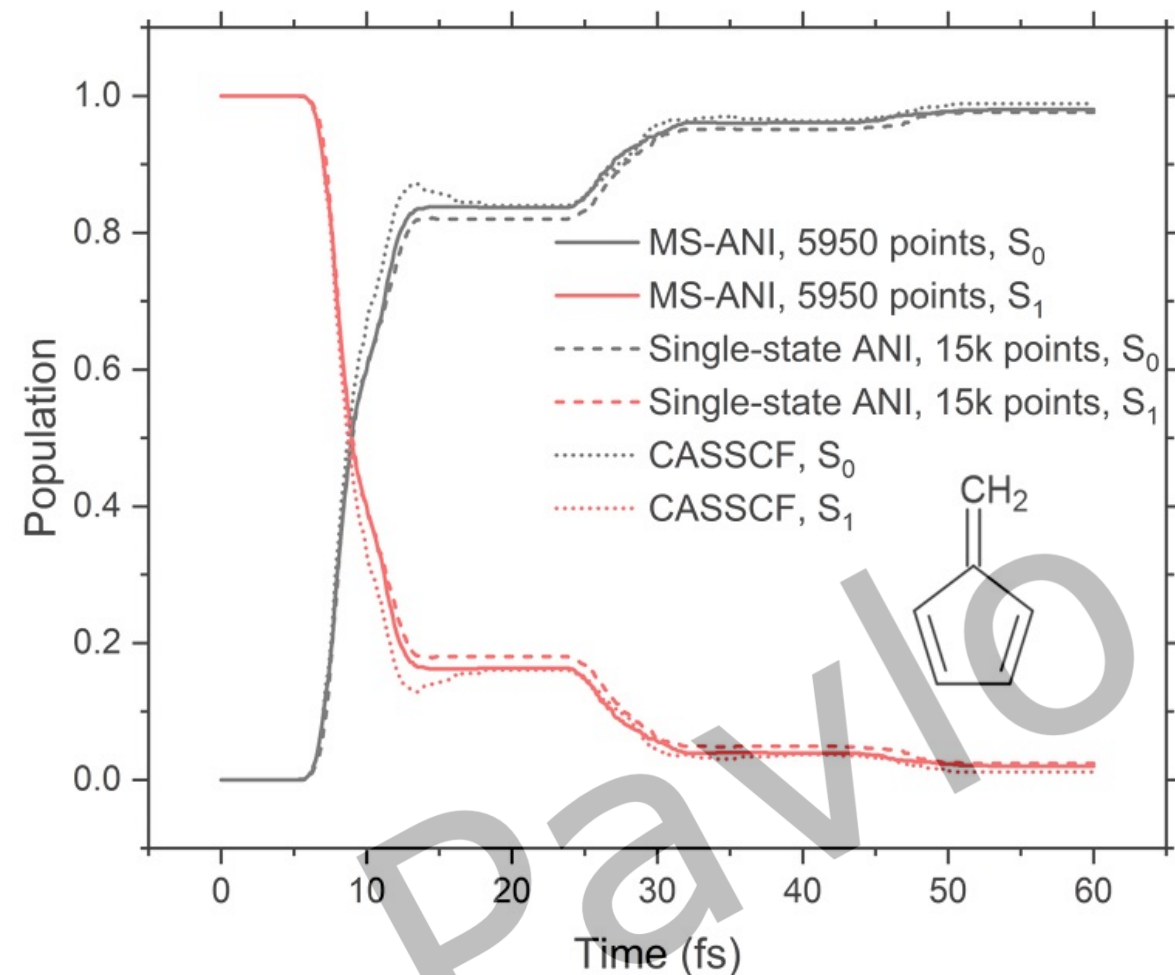
**After 2021, open questions such as: where is the place for equivariant NNs?**

Crashed after 28 days of active learning after 13 iterations and producing 3850 training points

**Use equivariant MLPs if you can  
like use CCSD(T) if you can  
(mostly we use ANI-type like mostly  
people use DFT)**



M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>



19 iterations of active learning

**Three days** on

RTX 4090 GPU and 16 Intel Xeon Gold 6226R CPUs for the entire procedure with trainings and labeling

M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

pubs.acs.org/JPCA

Perspective

# Machine Learning Interatomic Potentials and Long-Range Physics

Dylan M. Anstine and Olexandr Isayev\*



Cite This: *J. Phys. Chem. A* 2023, 127, 2417–2431



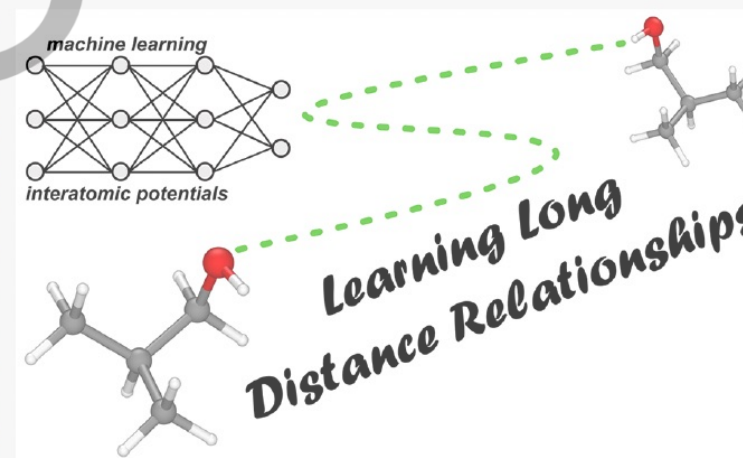
Read Online

ACCESS |

 Metrics & More

 Article Recommendations

**ABSTRACT:** Advances in machine learned interatomic potentials (MLIPs), such as those using neural networks, have resulted in short-range models that can infer interaction energies with near ab initio accuracy and orders of magnitude reduced computational cost. For many atom systems, including macromolecules, biomolecules, and condensed matter, model accuracy can become reliant on the description of short- and long-range physical interactions. The latter terms can be difficult to incorporate into an MLIP framework. Recent research has produced numerous models with considerations for nonlocal electrostatic and dispersion interactions, leading to a large range of applications that can be addressed using MLIPs. In light of this, we present a Perspective focused on key methodologies and models being used where the presence of nonlocal physics and chemistry are crucial for describing system properties. The strategies covered include MLIPs augmented with dispersion corrections, electrostatics calculated with charges predicted from atomic environment descriptors, the use of self-consistency and message passing iterations to propagated nonlocal system information, and charges obtained via equilibration schemes. We aim to provide a pointed discussion to support the development of machine learning-based interatomic potentials for systems where contributions from only nearsighted terms are deficient.



pubs.acs.org/CR

Review

## Four Generations of High-Dimensional Neural Network Potentials

Jörg Behler\*



Cite This: *Chem. Rev.* 2021, 121, 10037–10072



Read Online

ACCESS |

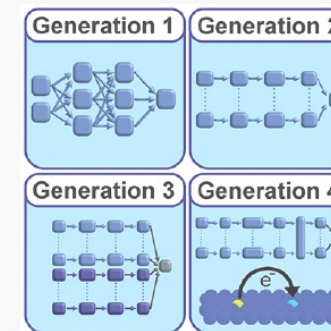


Metrics & More



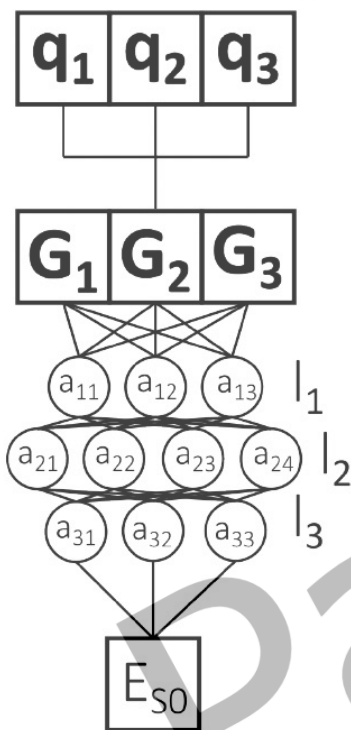
Article Recommendations

**ABSTRACT:** Since their introduction about 25 years ago, machine learning (ML) potentials have become an important tool in the field of atomistic simulations. After the initial decade, in which neural networks were successfully used to construct potentials for rather small molecular systems, the development of high-dimensional neural network potentials (HDNNPs) in 2007 opened the way for the application of ML potentials in simulations of large systems containing thousands of atoms. To date, many other types of ML potentials have been proposed continuously increasing the range of problems that can be studied. In this review, the methodology of the family of HDNNPs including new recent developments will be discussed using a classification scheme into four generations of potentials, which is also applicable to many other types of ML potentials. The first generation is formed by early neural network potentials designed for low-dimensional systems. High-dimensional neural network potentials established the second generation and are based on three key steps: first, the expression of the total energy as a sum of environment-dependent atomic energy contributions; second, the description of the atomic environments by atom-centered symmetry functions as descriptors fulfilling the requirements of rotational, translational, and permutation invariance; and third, the iterative construction of the reference electronic structure data sets by active learning. In third-generation HDNNPs, in addition, long-range interactions are included employing environment-dependent partial charges expressed by atomic neural networks. In fourth-generation HDNNPs, which are just emerging, in addition, nonlocal phenomena such as long-range charge transfer can be included. The applicability and remaining limitations of HDNNPs are discussed along with an outlook at possible future developments.

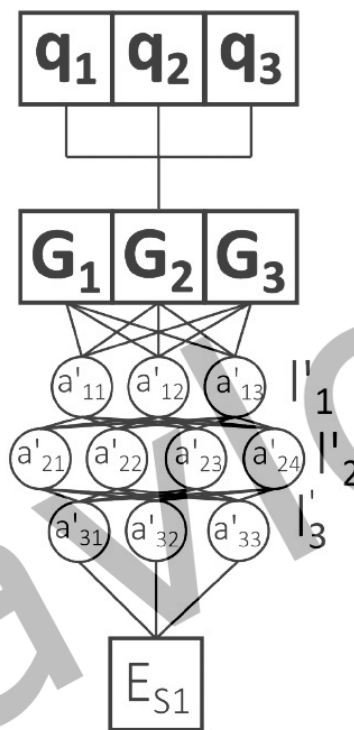


(a)  
Single-state model

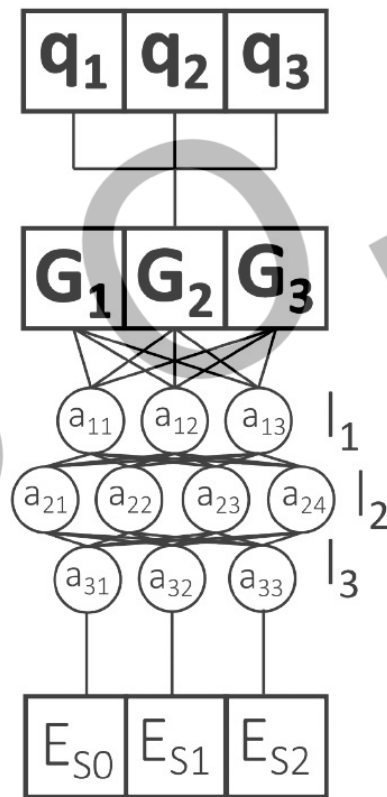
Network for  $S_0$

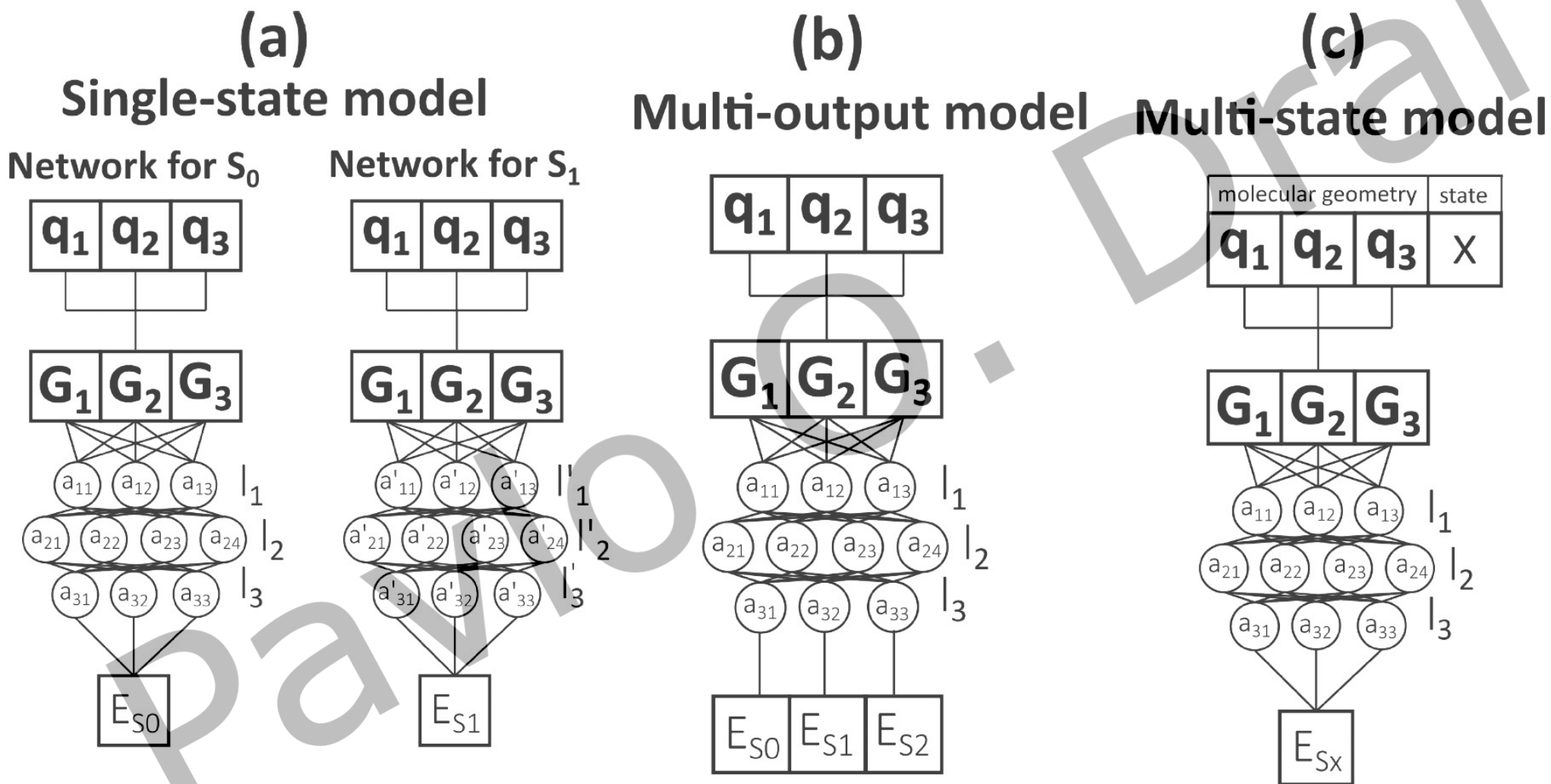


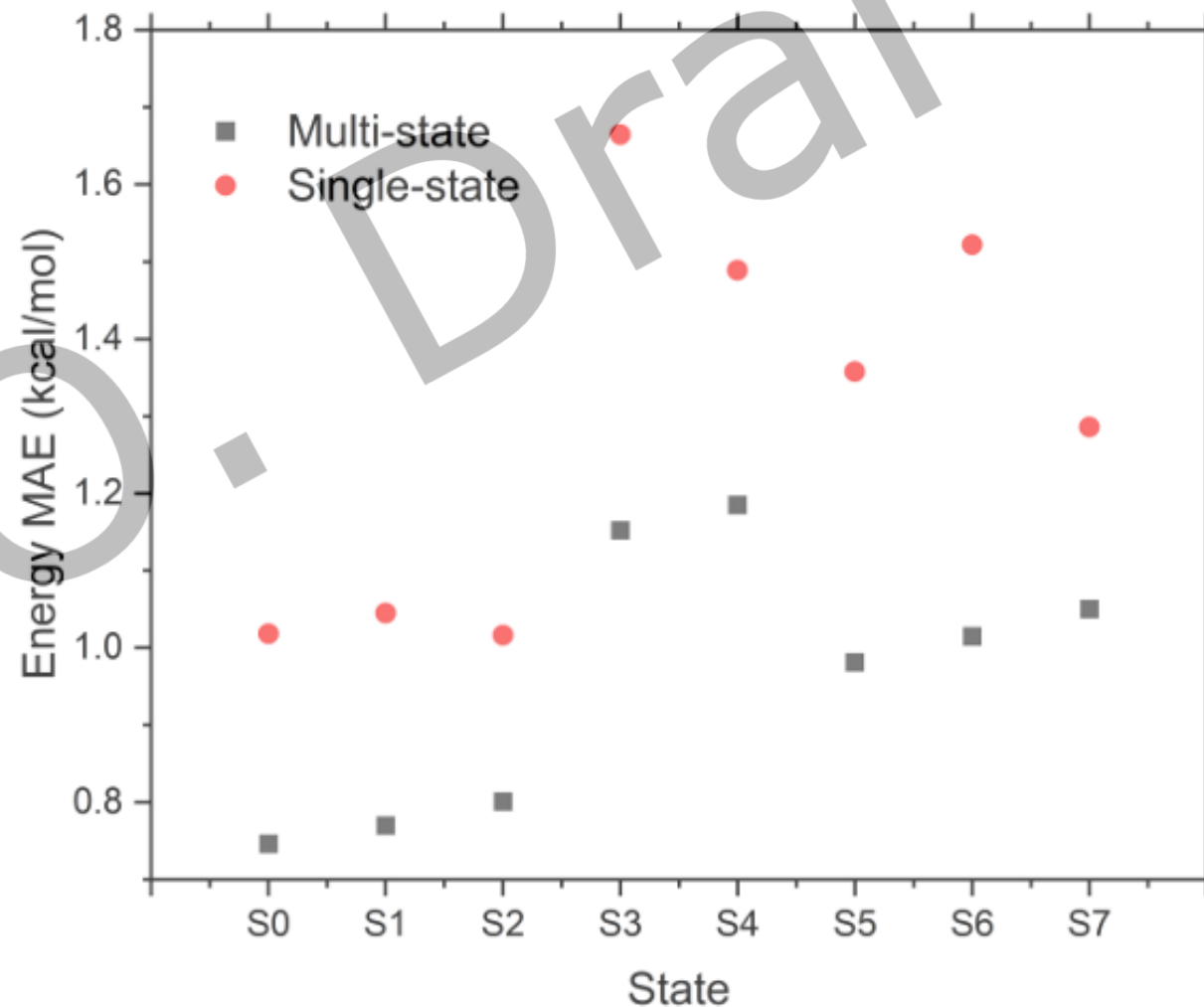
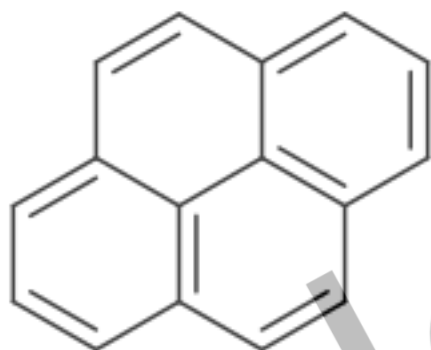
Network for  $S_1$



(b)  
Multi-output model



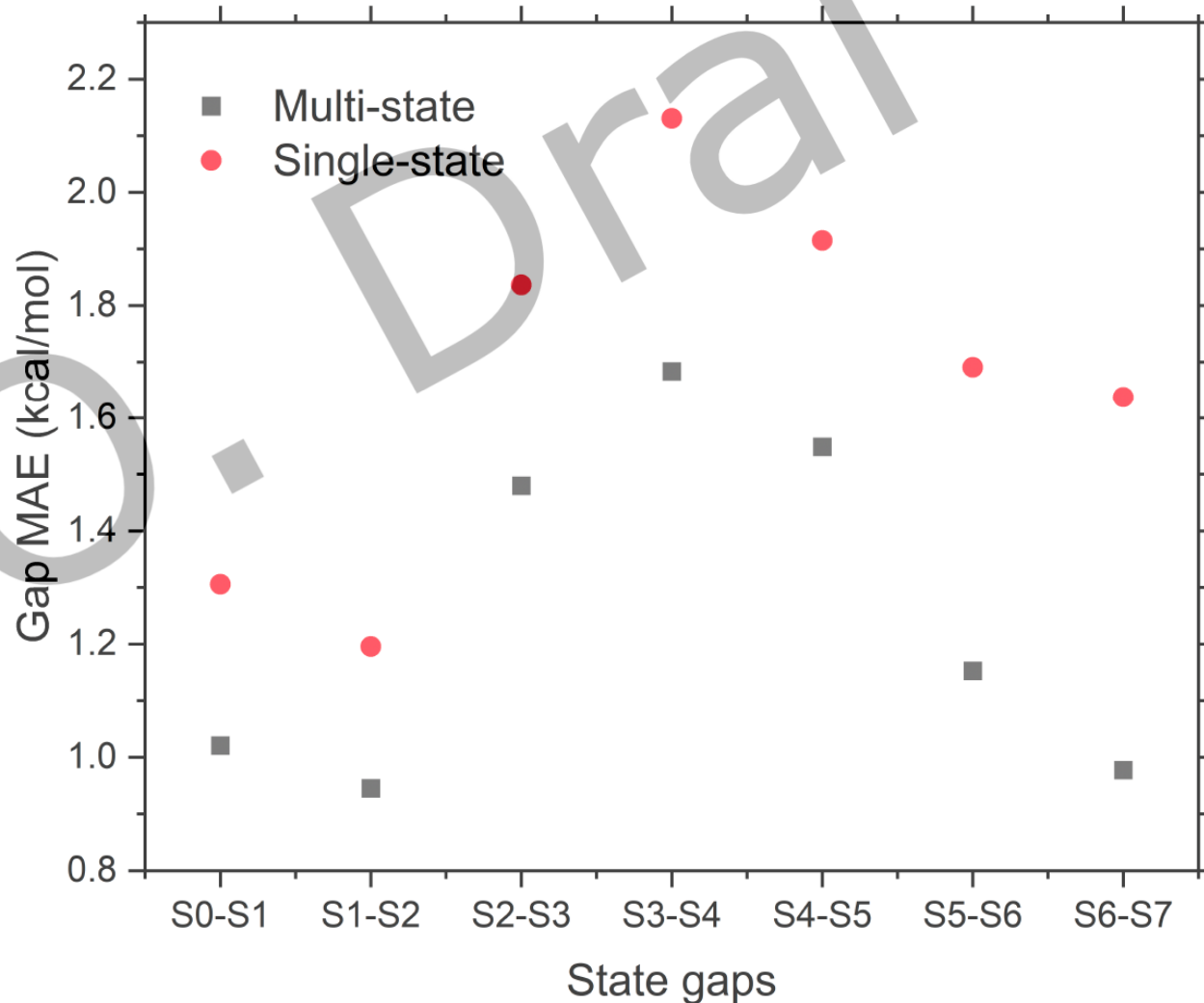




M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

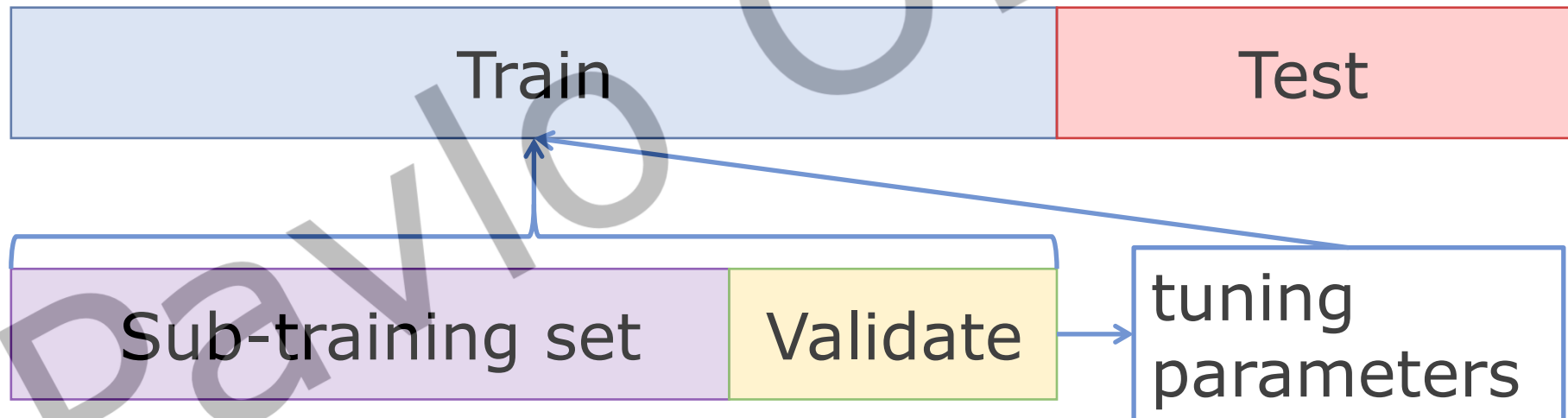


$$L = \omega_E L_E + \omega_F L_F + \omega_{\text{gap}} L_{\text{gap}}.$$



M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

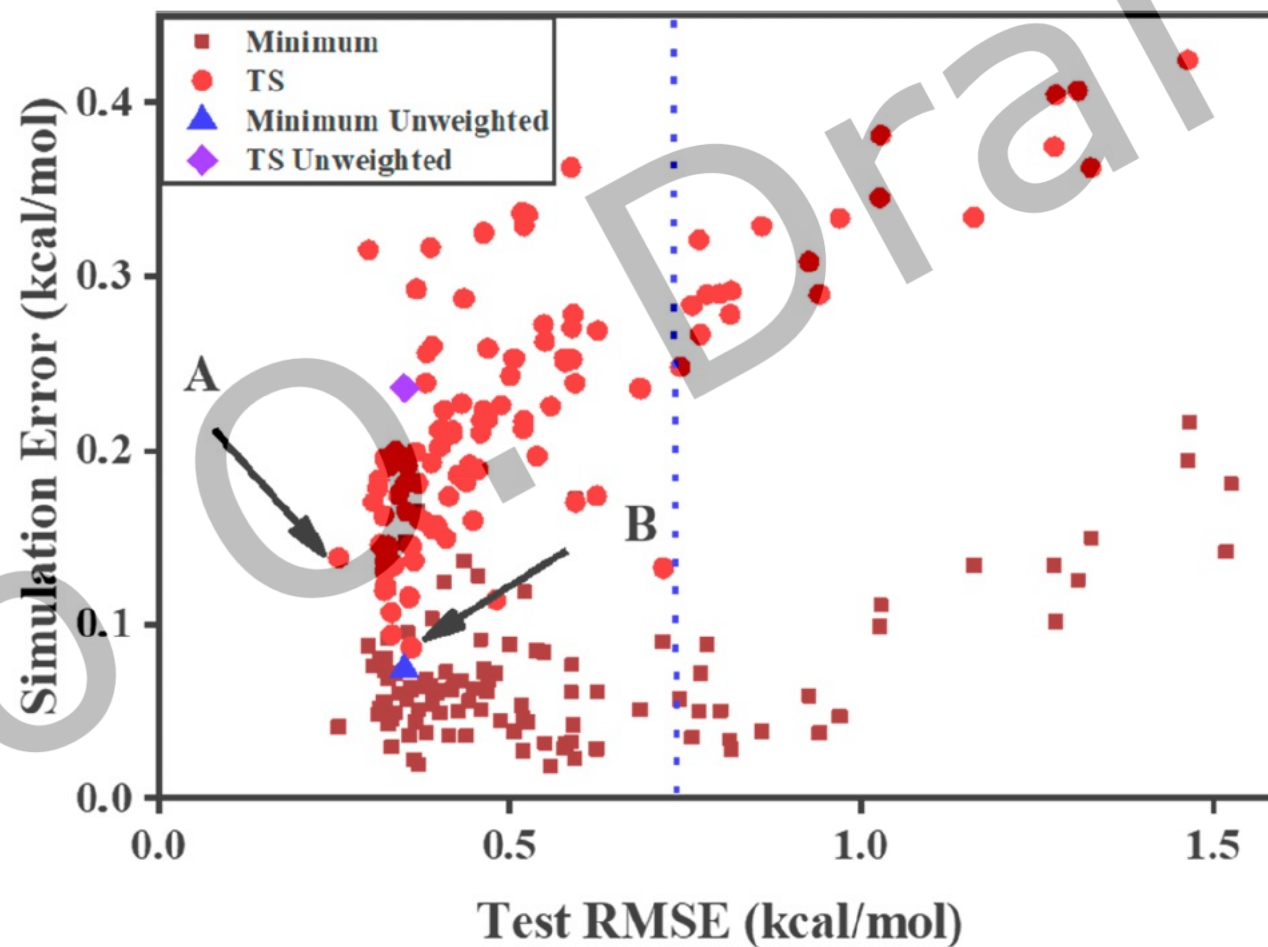
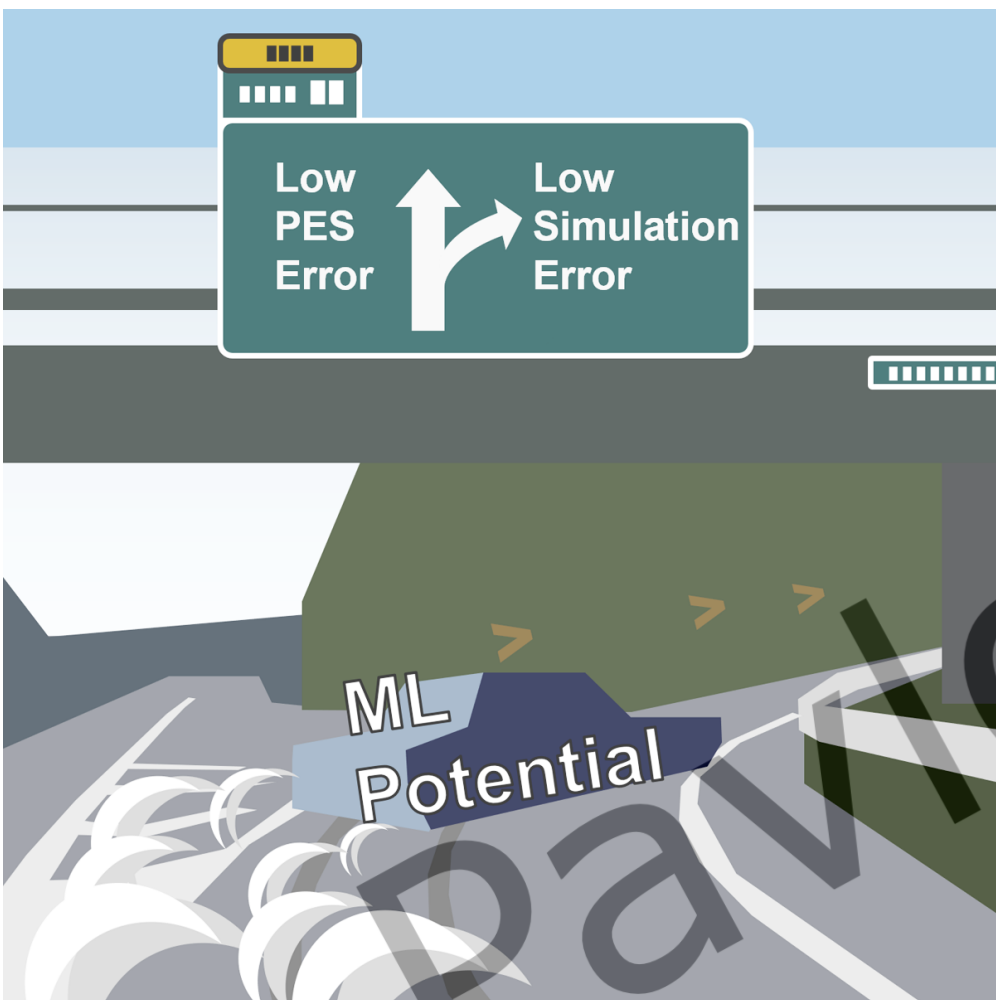
- Often ML error for its own training set is close to zero
- Using errors in the validation set would be also incorrect, because their minimization is a part of the training process
- We should estimate errors on a **completely independent test set**

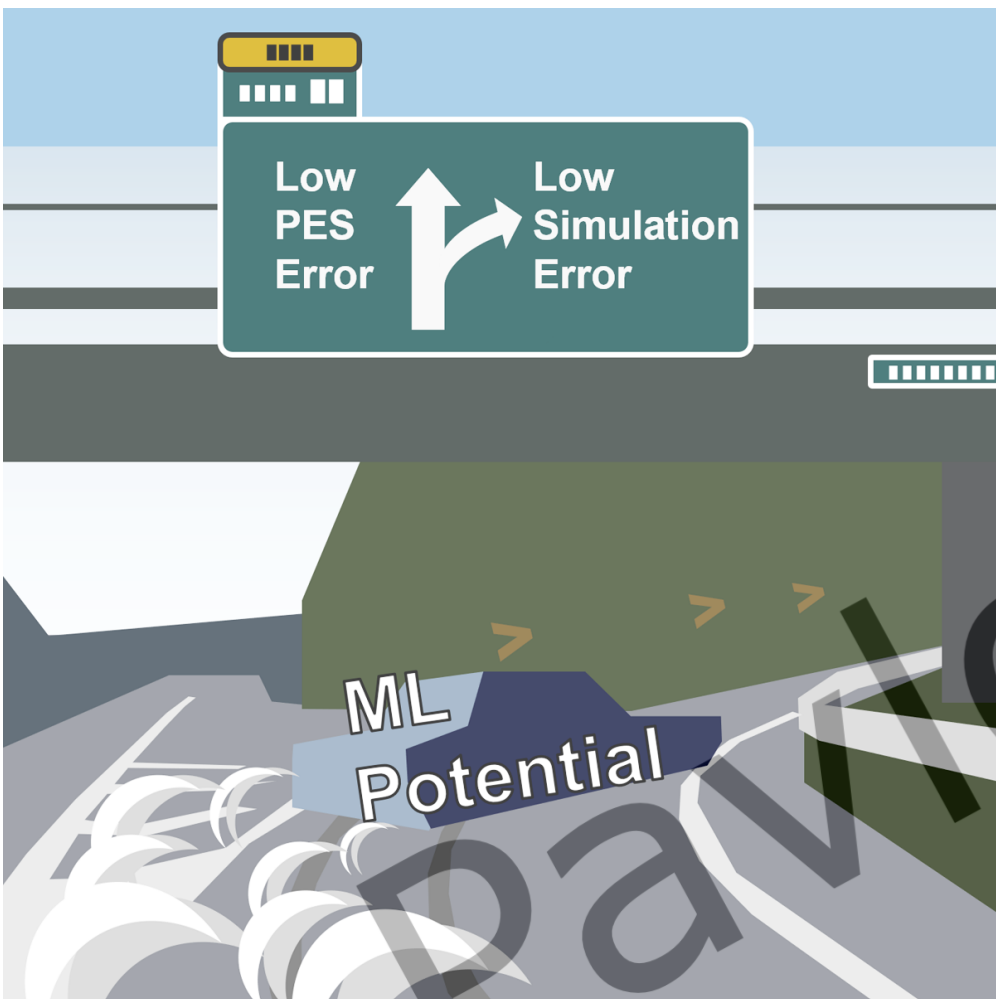


**The ultimate test is the performance in the required application!**

Pavlo O. Dral

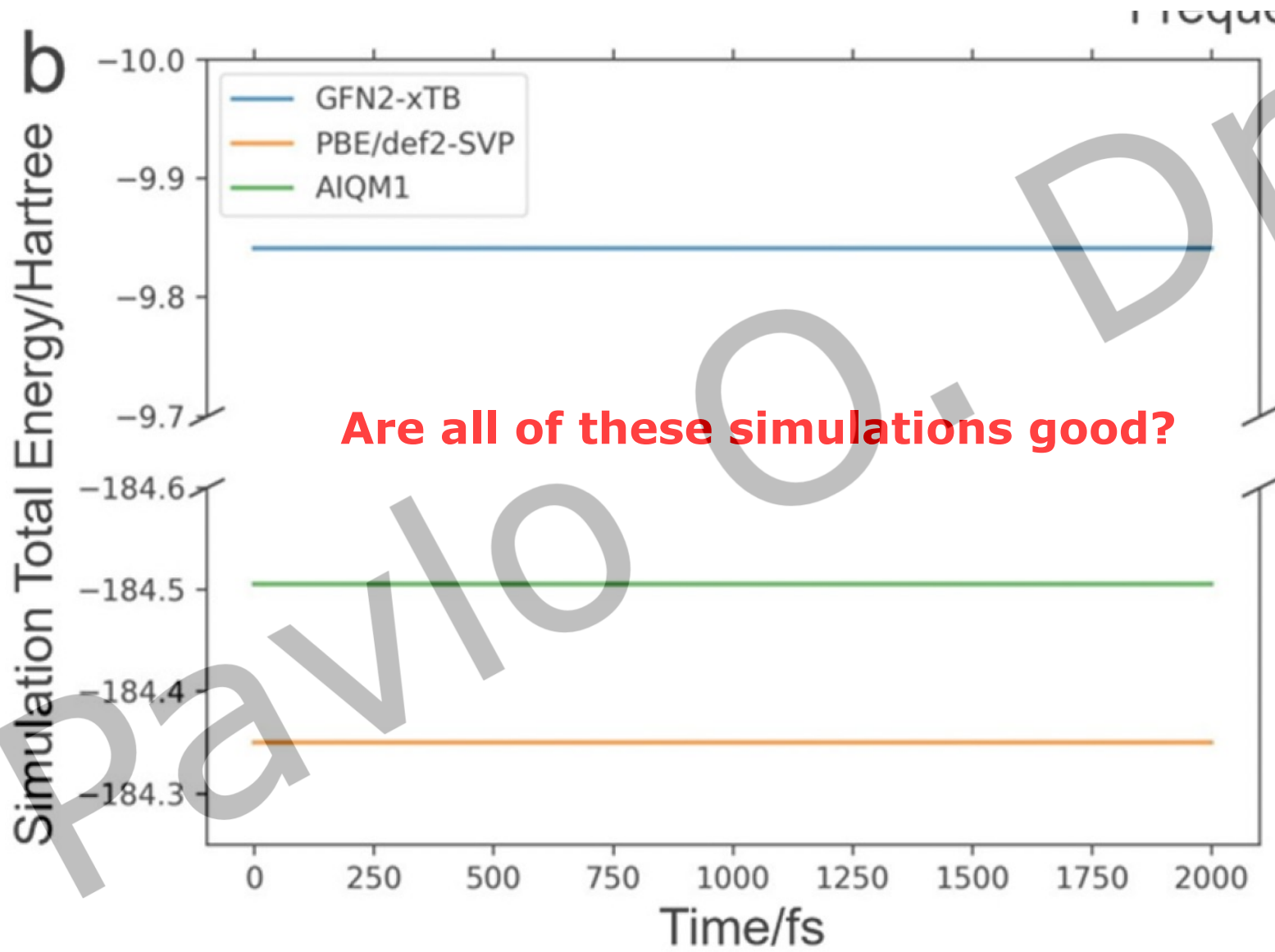
# Quality of simulation $\neq$ quality of fit

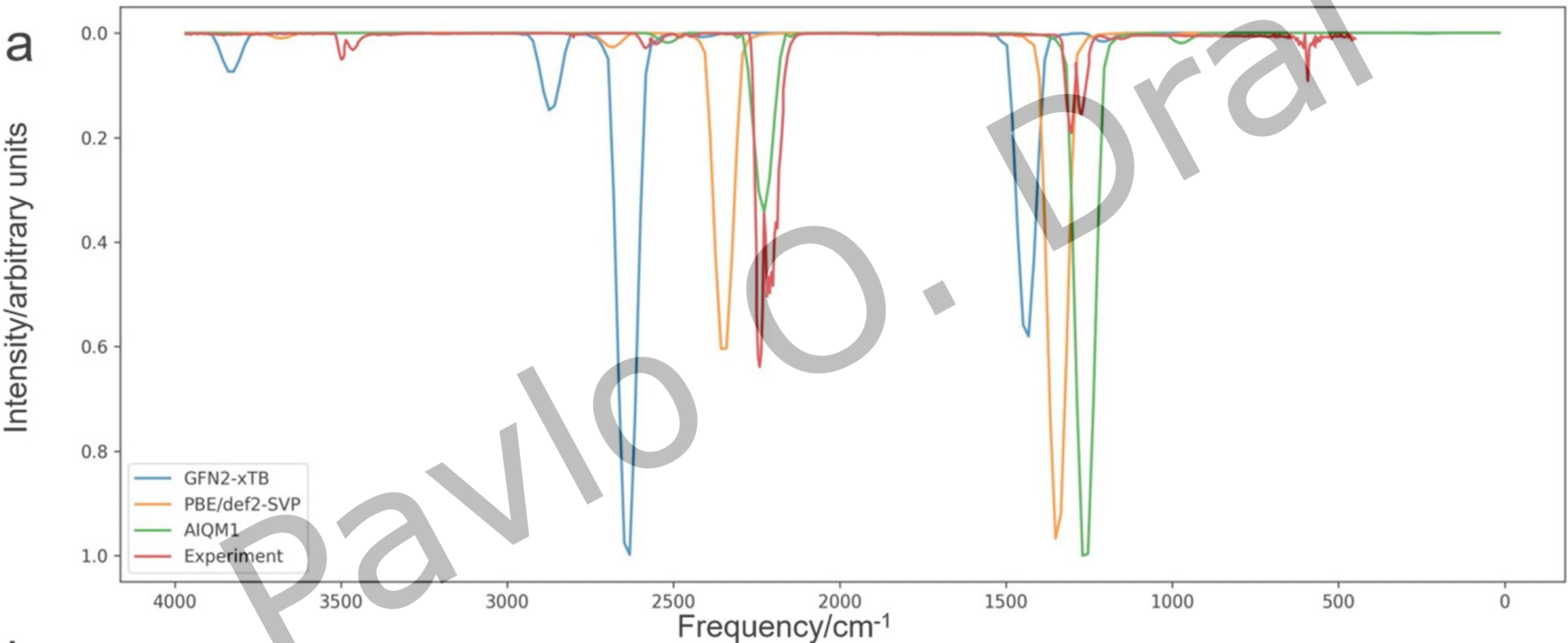




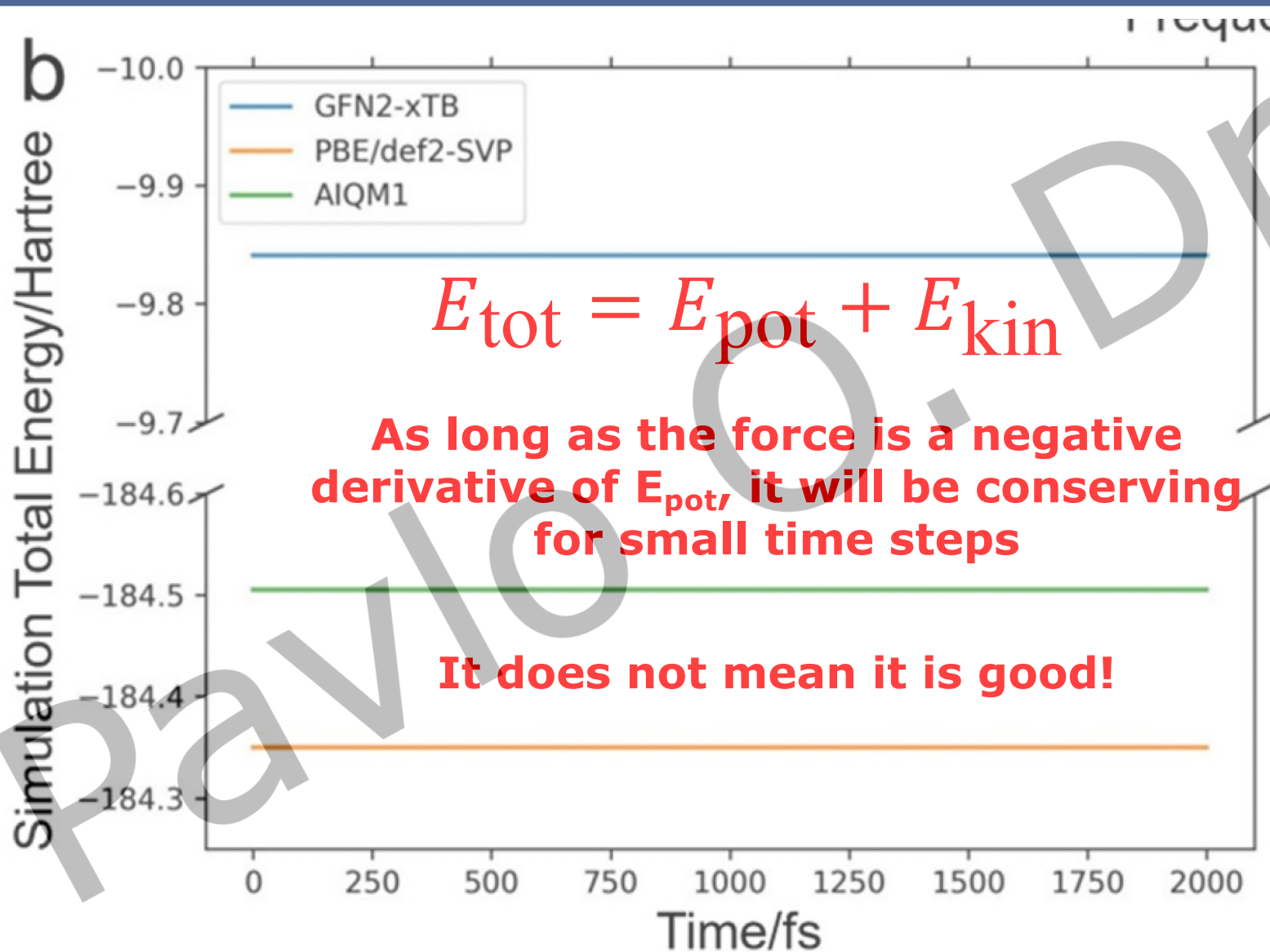
"One [diffusion Monte Carlo] simulation for a single conformer requires 30000 walkers and 55000 steps comprising roughly  **$1.6 \cdot 10^9$**  potential evaluations with B3LYP."

→ 60 hours on a single GPU with ANI...

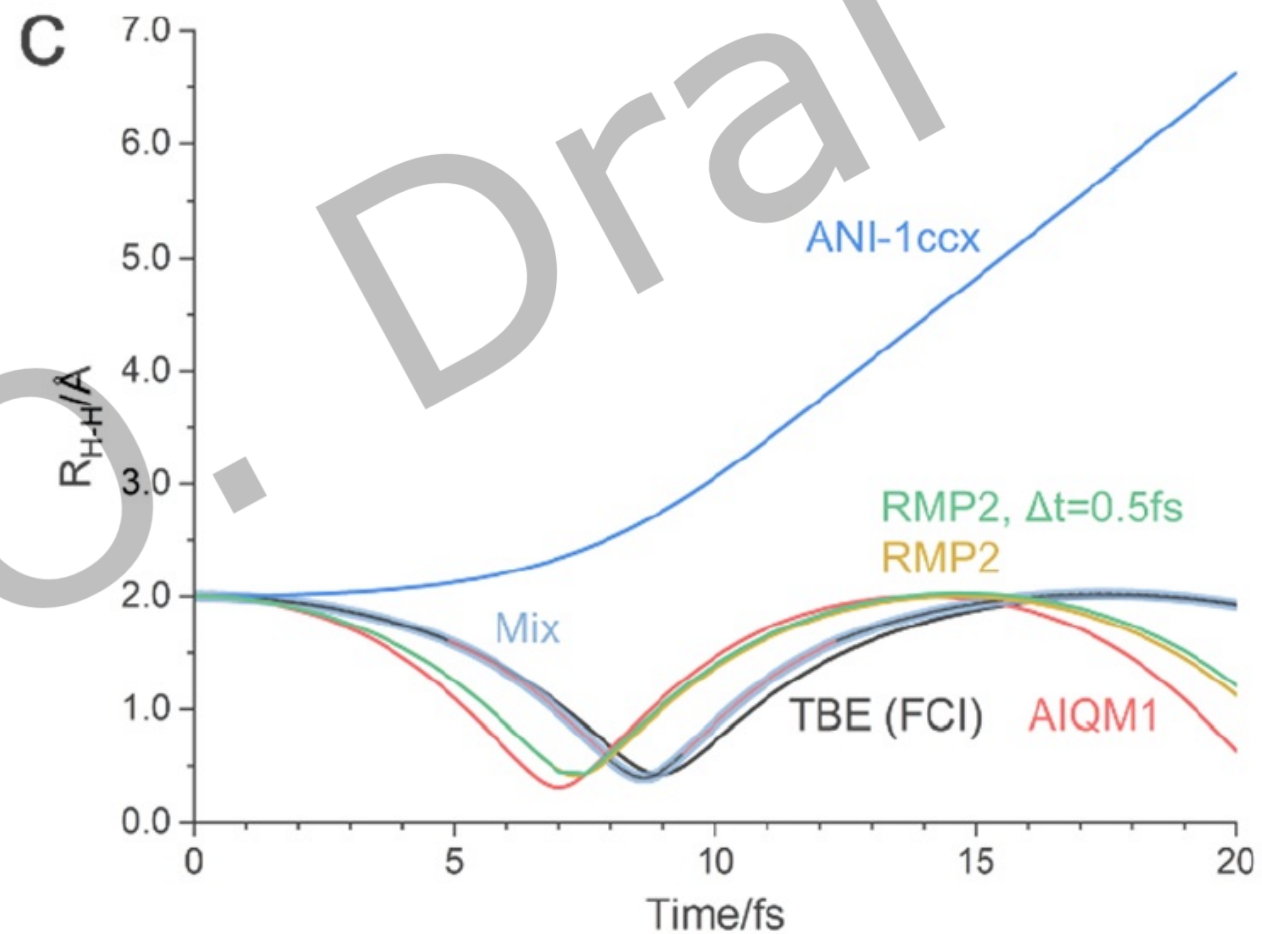
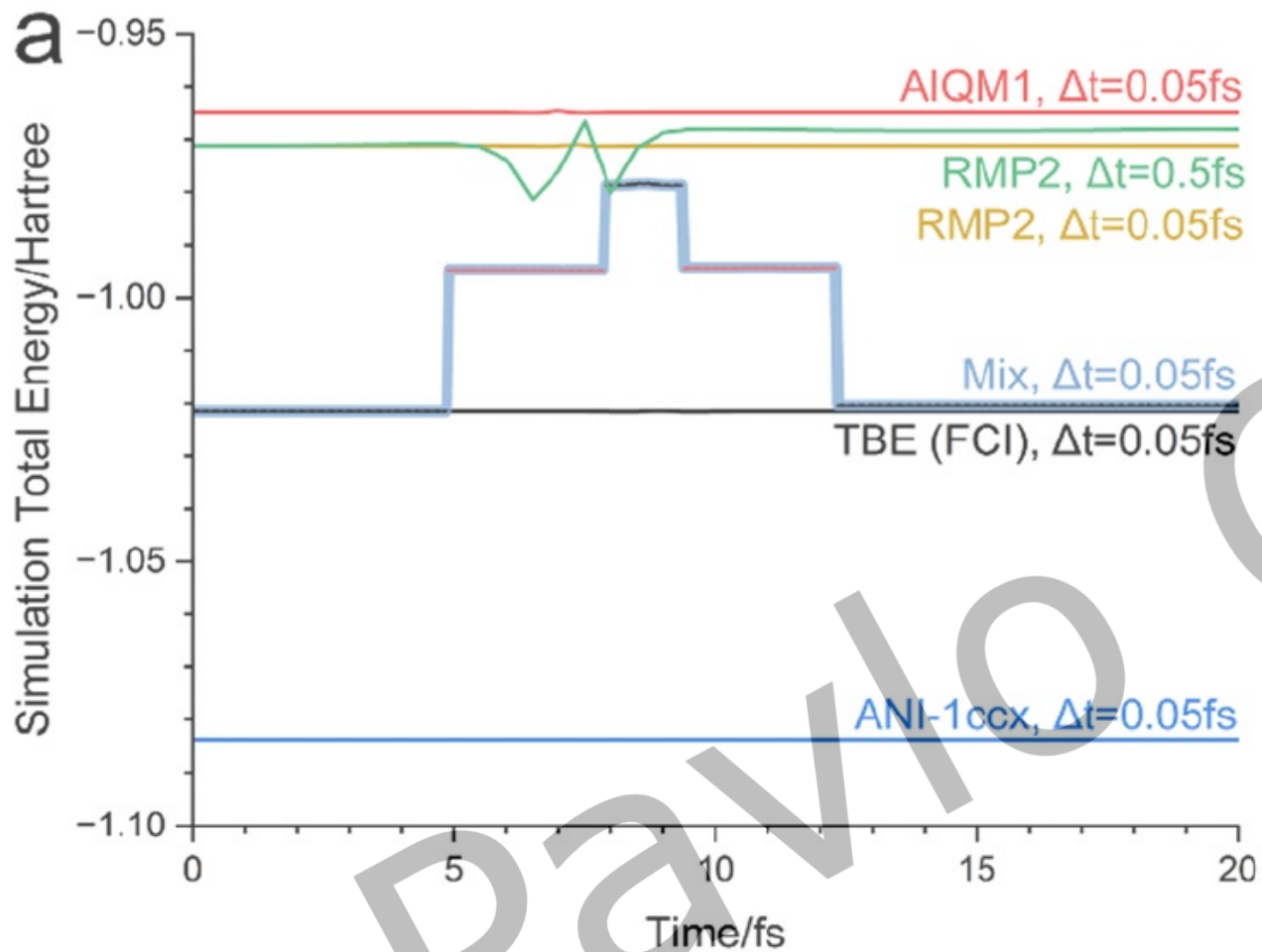


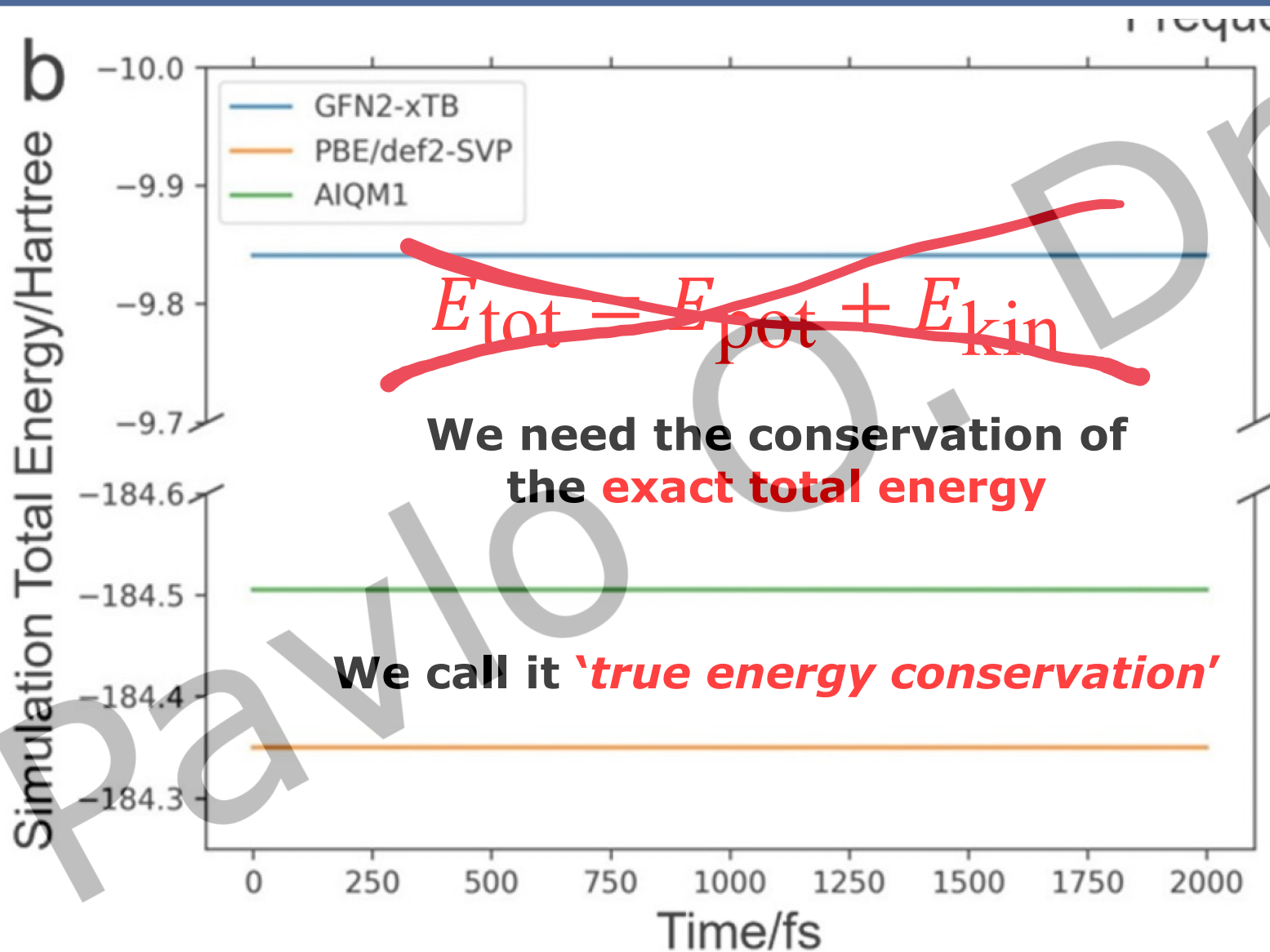


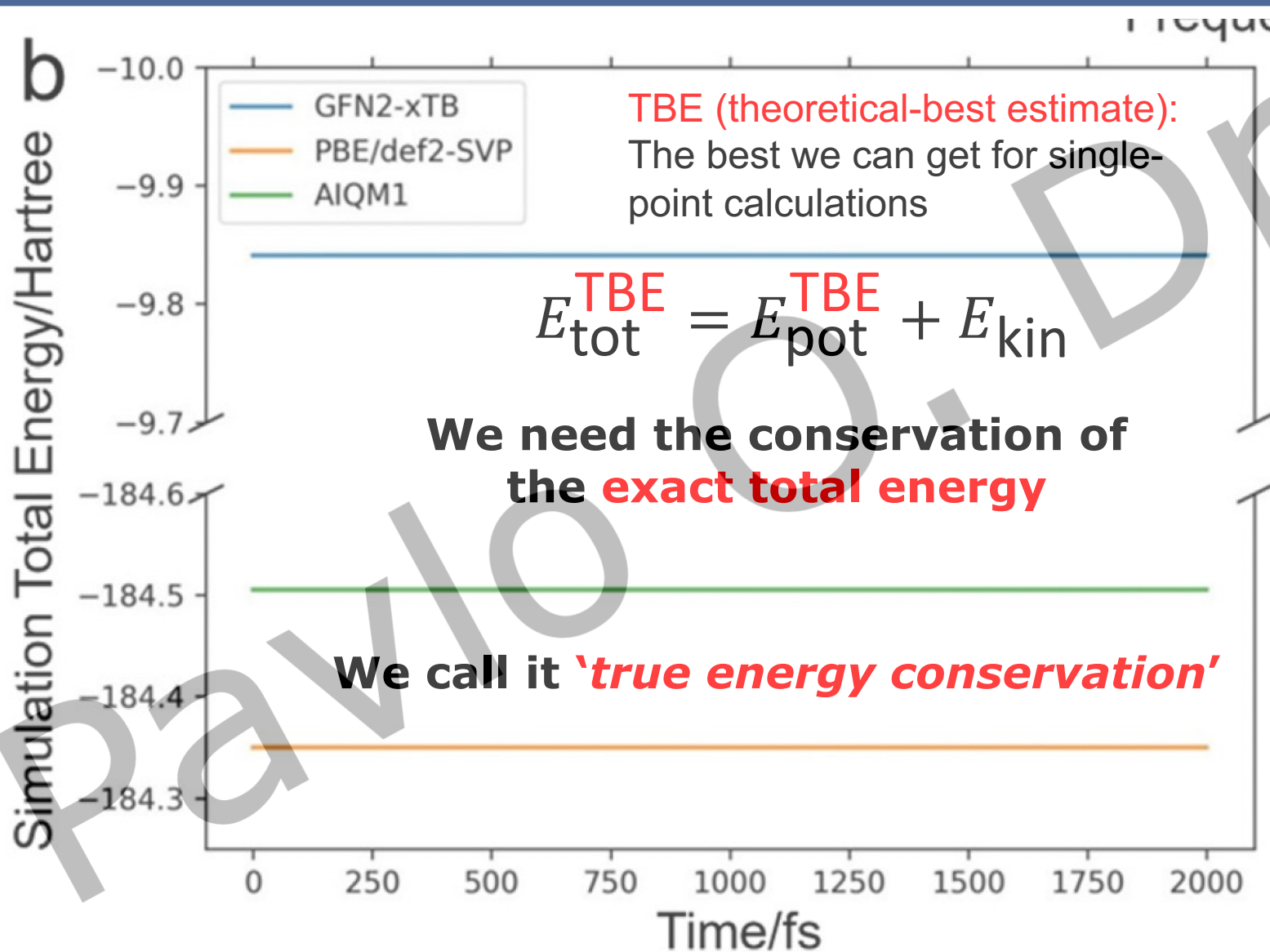
**Why are they so different in accuracy?**







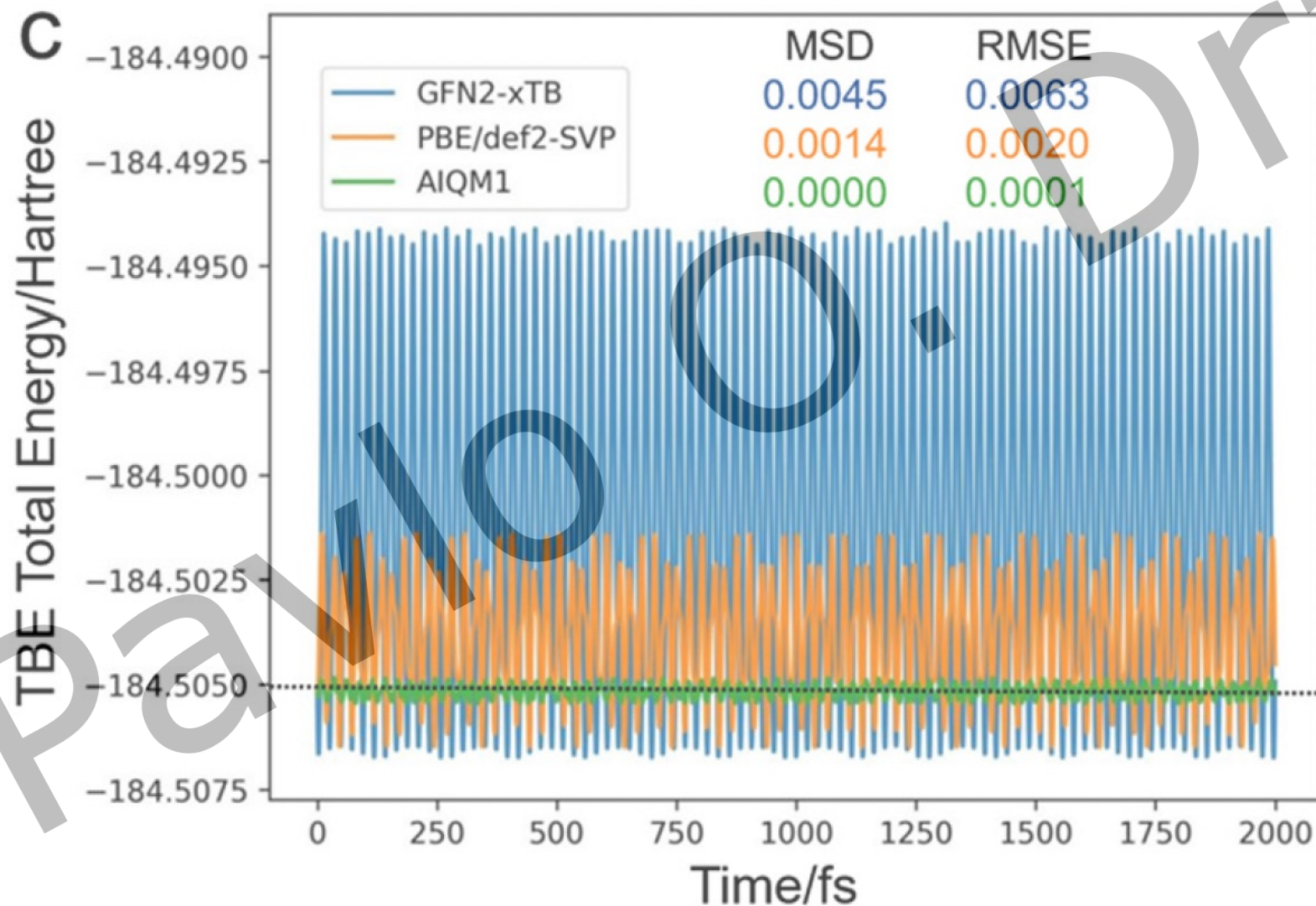


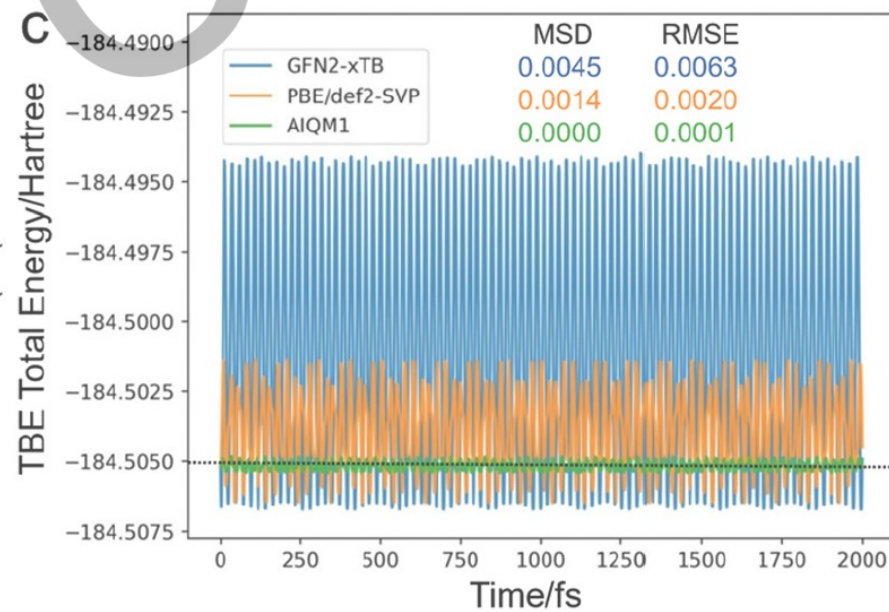
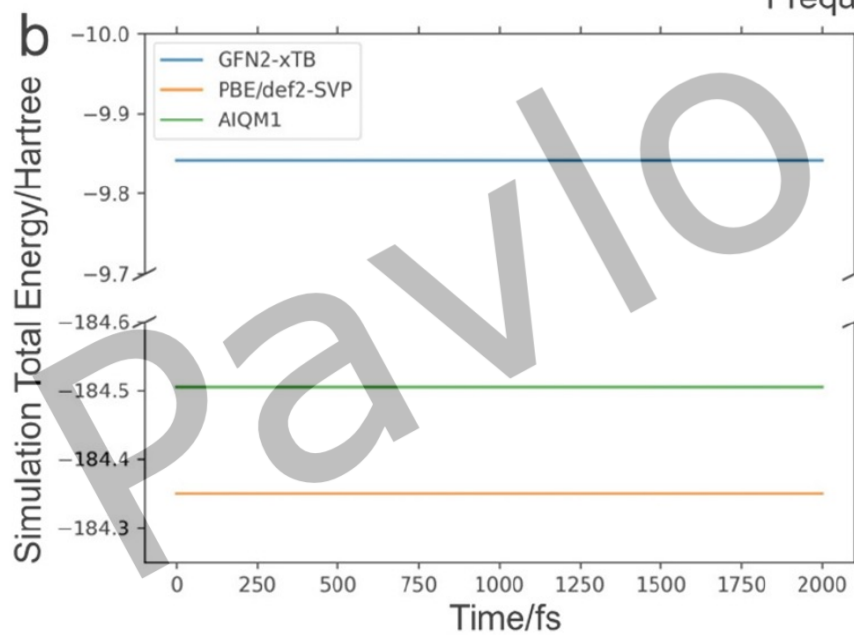
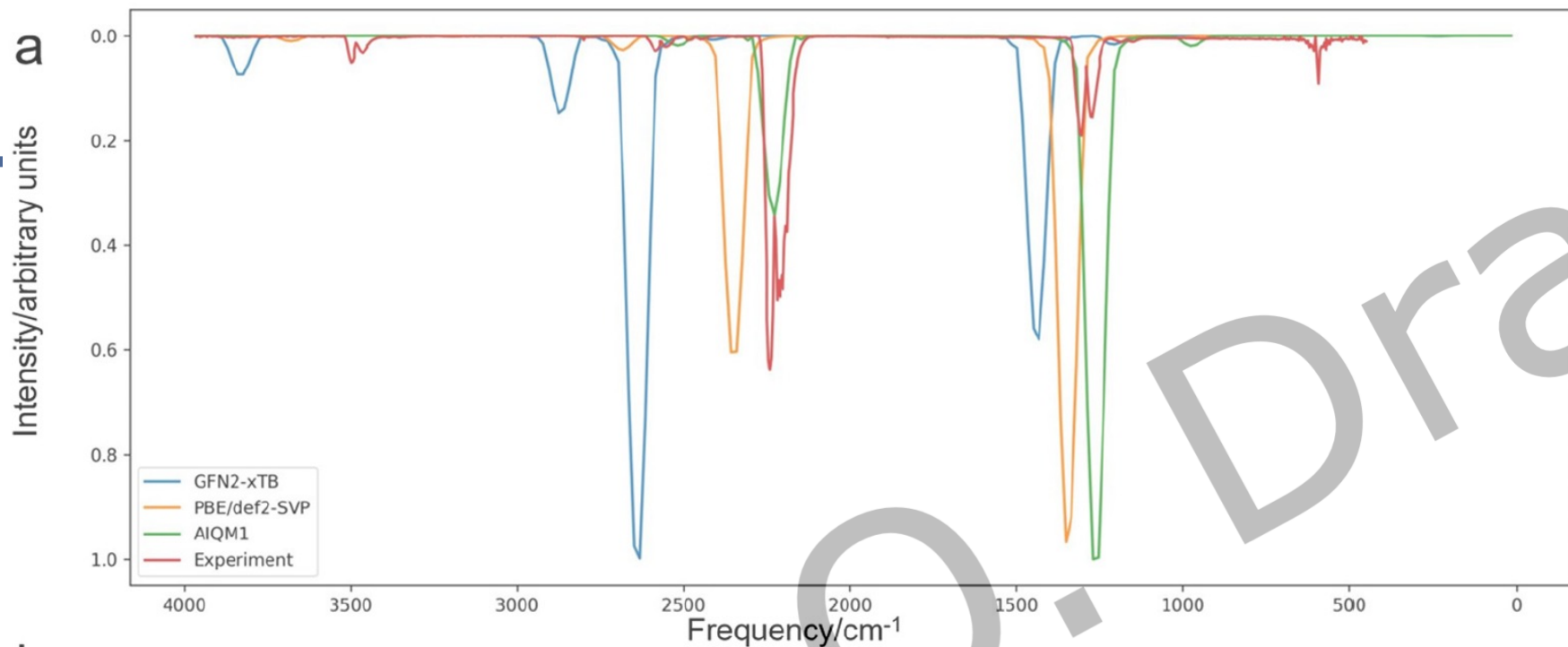


$$\text{RMSE} = \sqrt{1/\text{number of time steps} \sum \left( E_{\text{tot}}^{\text{TBE}}(\text{time step}) - E_{\text{tot}}^{\text{TBE}}(\text{time zero}) \right)^2}$$

$$\text{MSD} = 1/\text{number of time steps} \sum \left( E_{\text{tot}}^{\text{TBE}}(\text{time step}) - E_{\text{tot}}^{\text{TBE}}(\text{time zero}) \right)$$

**TBE: CCSD(T)\*/CBS**





# Supervised Machine Learning

$$E = f(\mathbf{R})$$

$$F_{A,d} = -\frac{\partial E}{\partial x_{A,d}}$$

Input (x) → f(x) → Output (y)

Given collection of **known {x,y}** find a **function f(x)**  
 training set                      train                      ML model

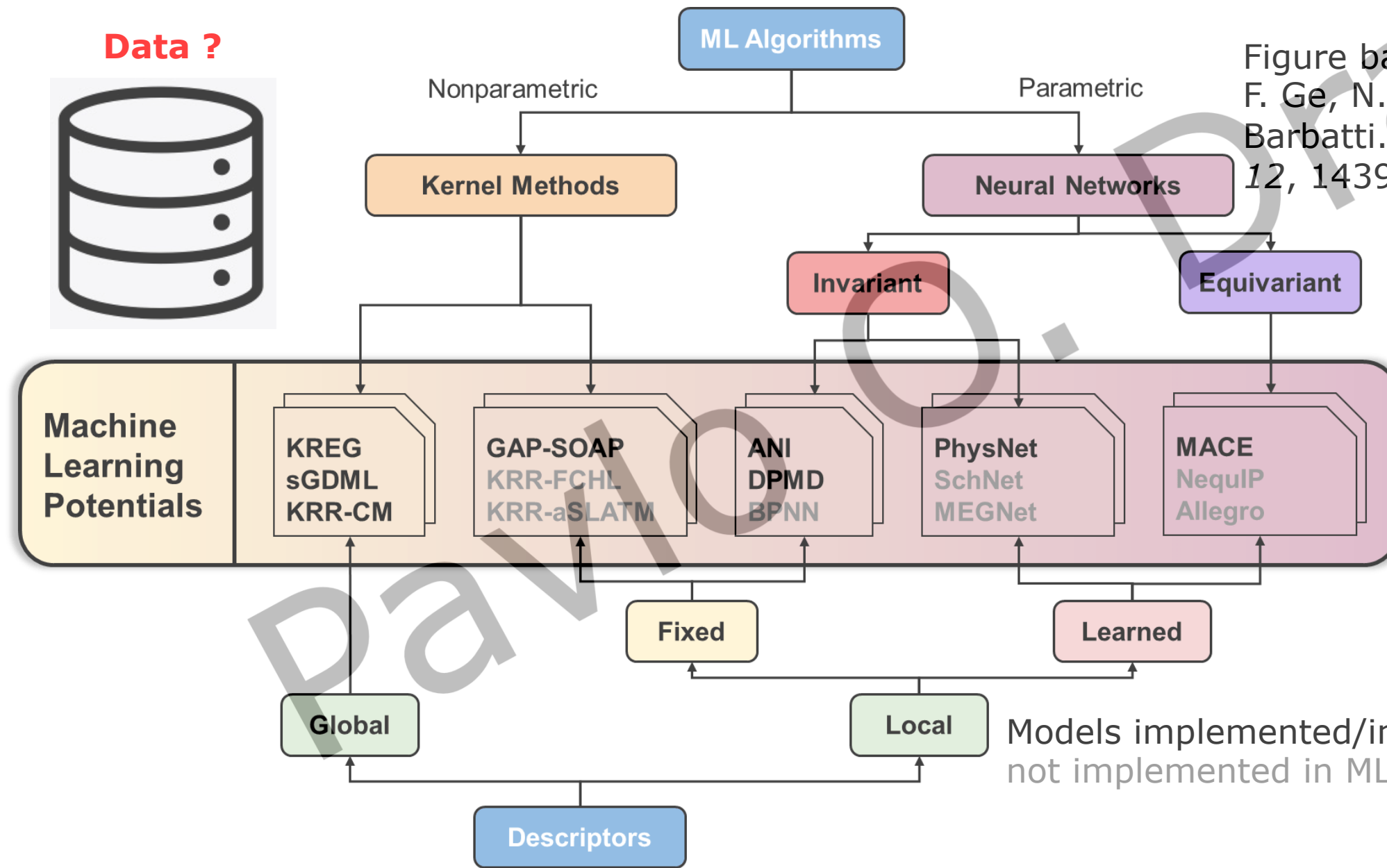
Use this function for making new predictions given just {x'}

- **Data**
- Choice of x (descriptor)
- Choice of y (labels)
- Fitting function (ML algorithm, ML model)
- Optimization of ML model parameters

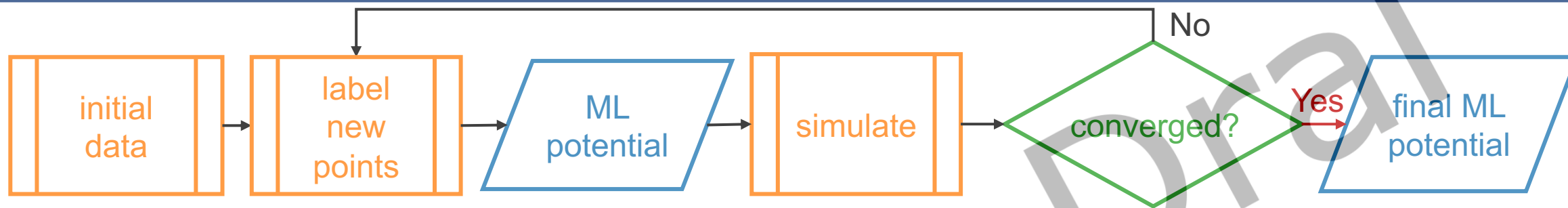
Data ?



Figure based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, *12*, 14396–14413

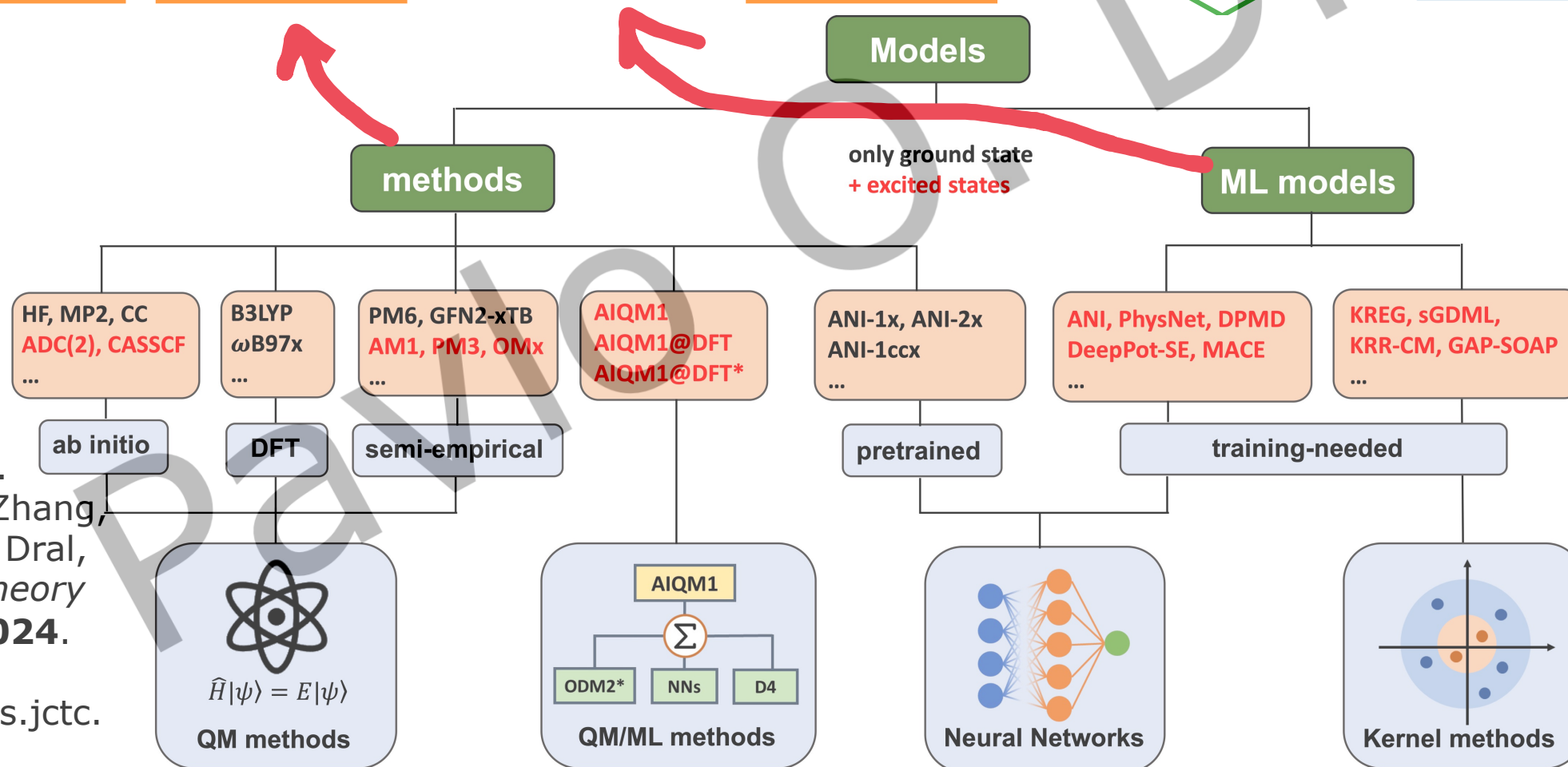
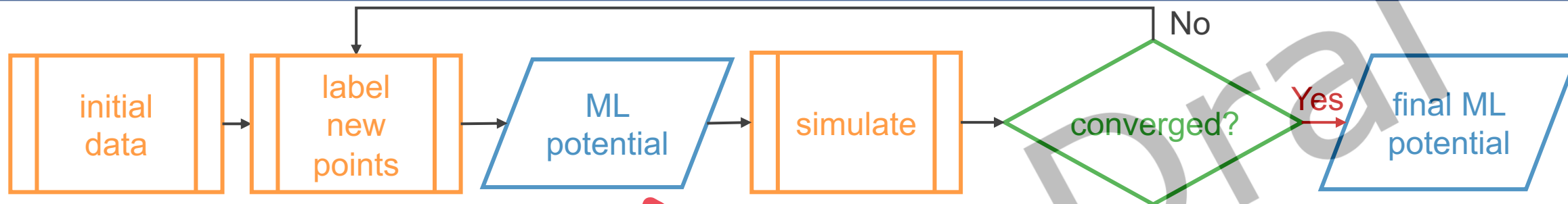


Models implemented/interfaced in **MLatom**  
not implemented in MLatom

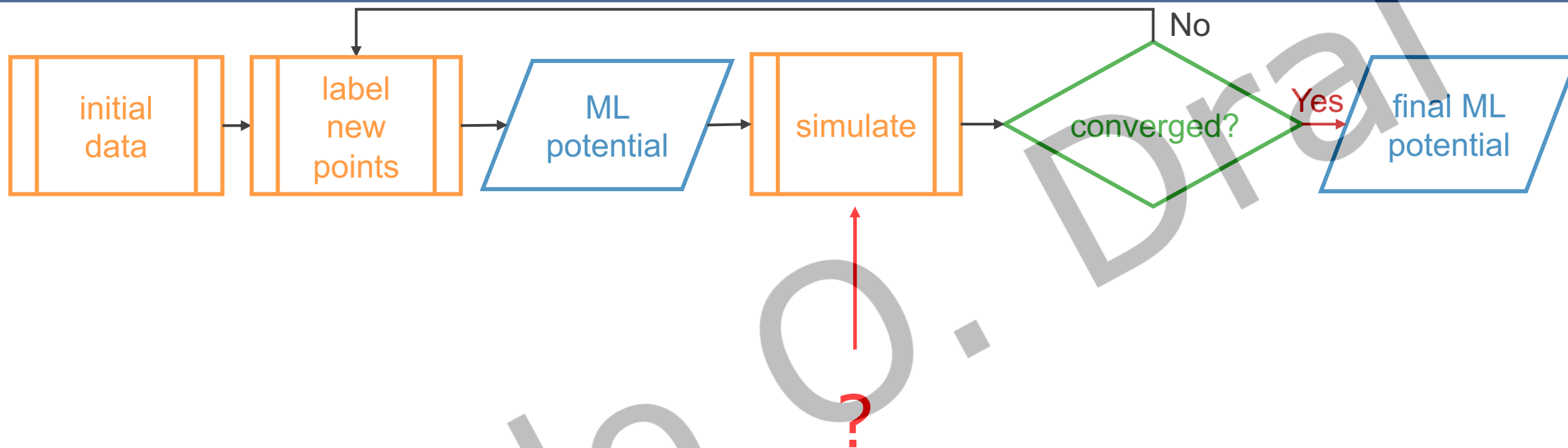


**Each step is highly non-trivial and you need methods & software!**





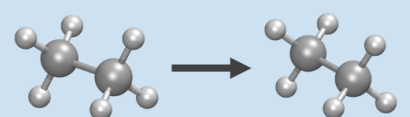
Y.-F. Hou, L. Zhang, Q. Zhang, F. Ge, P. O. Dral, *J. Chem. Theory Comput.* **2024**. DOI: 10.1021/acs.jctc.4c00821



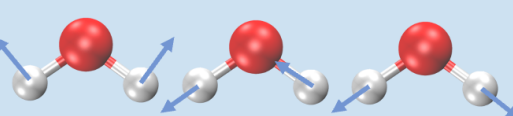
Pavlo

Single point calculations Energies, forces, Hessian matrix...

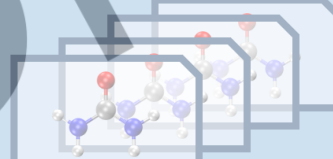
Geometry optimizations



Frequency calculations

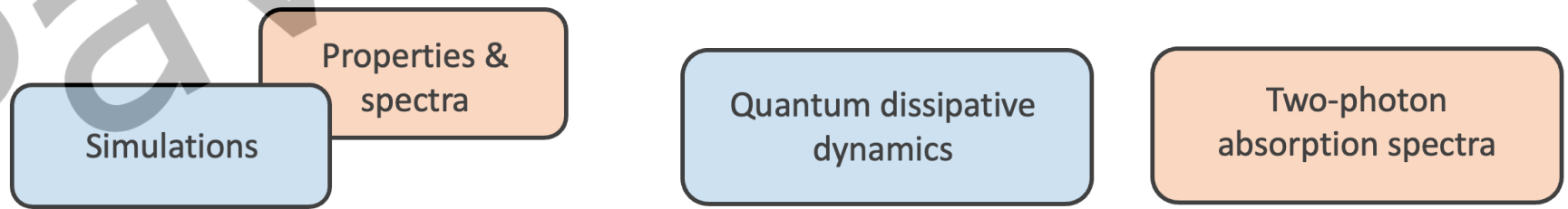
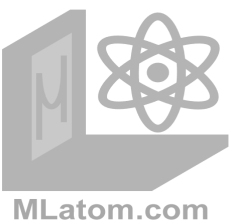
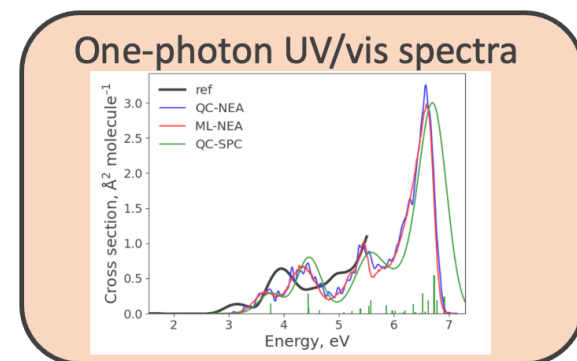
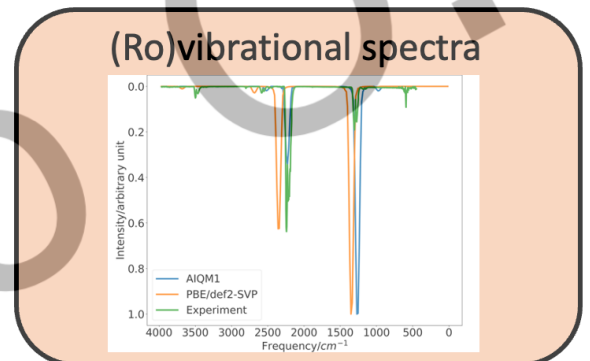


Molecular dynamics

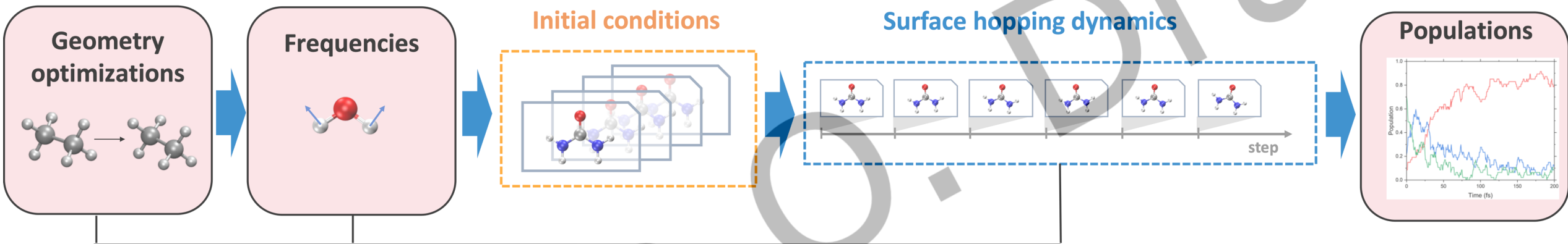


Thermochemistry calculations

Heat of formation:

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


# Surface-hopping dynamics



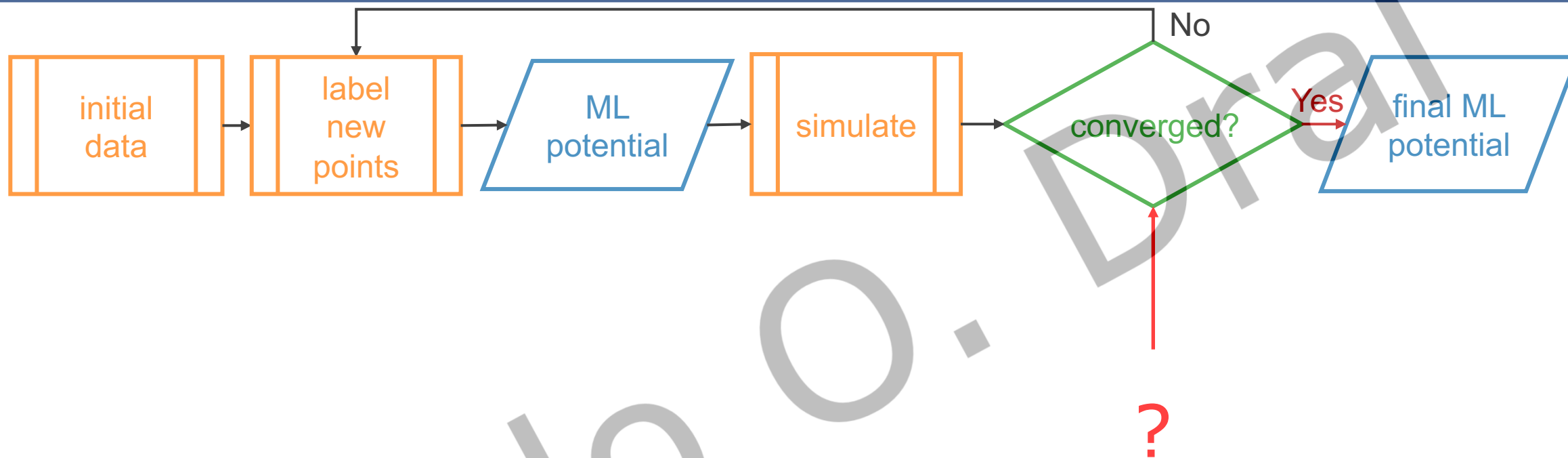
**Single point calculations**

$$E, \frac{\partial E}{\partial \mathbf{x}}, \frac{\partial^2 E}{\partial \mathbf{x}^2}$$

```
aiqm1=mlatom.models.methods(method='AIQM1')
geomopt=mlatom.optimize_geometry(model=aiqm1, ...)
...
mlatom.namd.surface_hopping_md(model=model, ...)
```

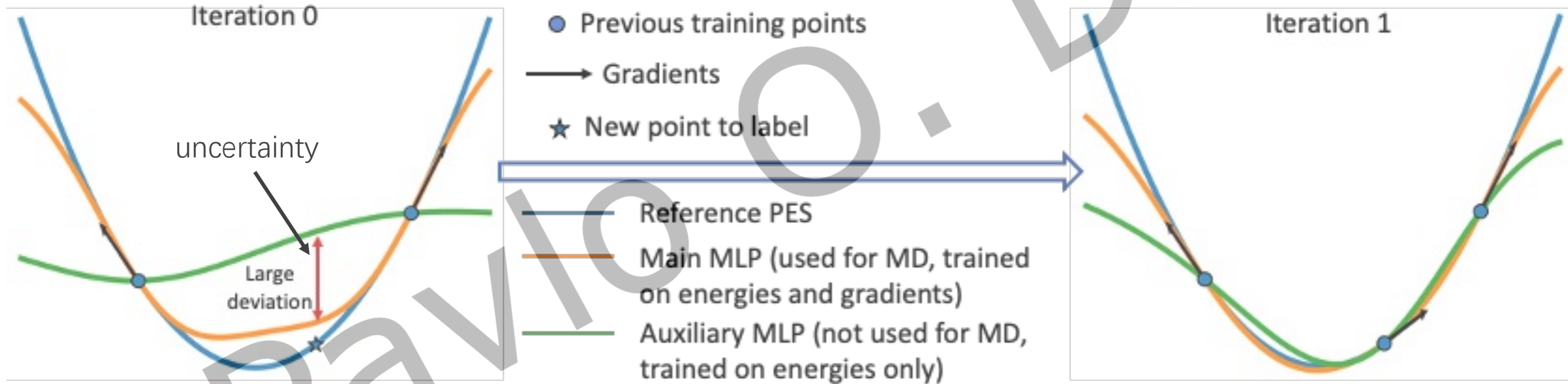
**Models**

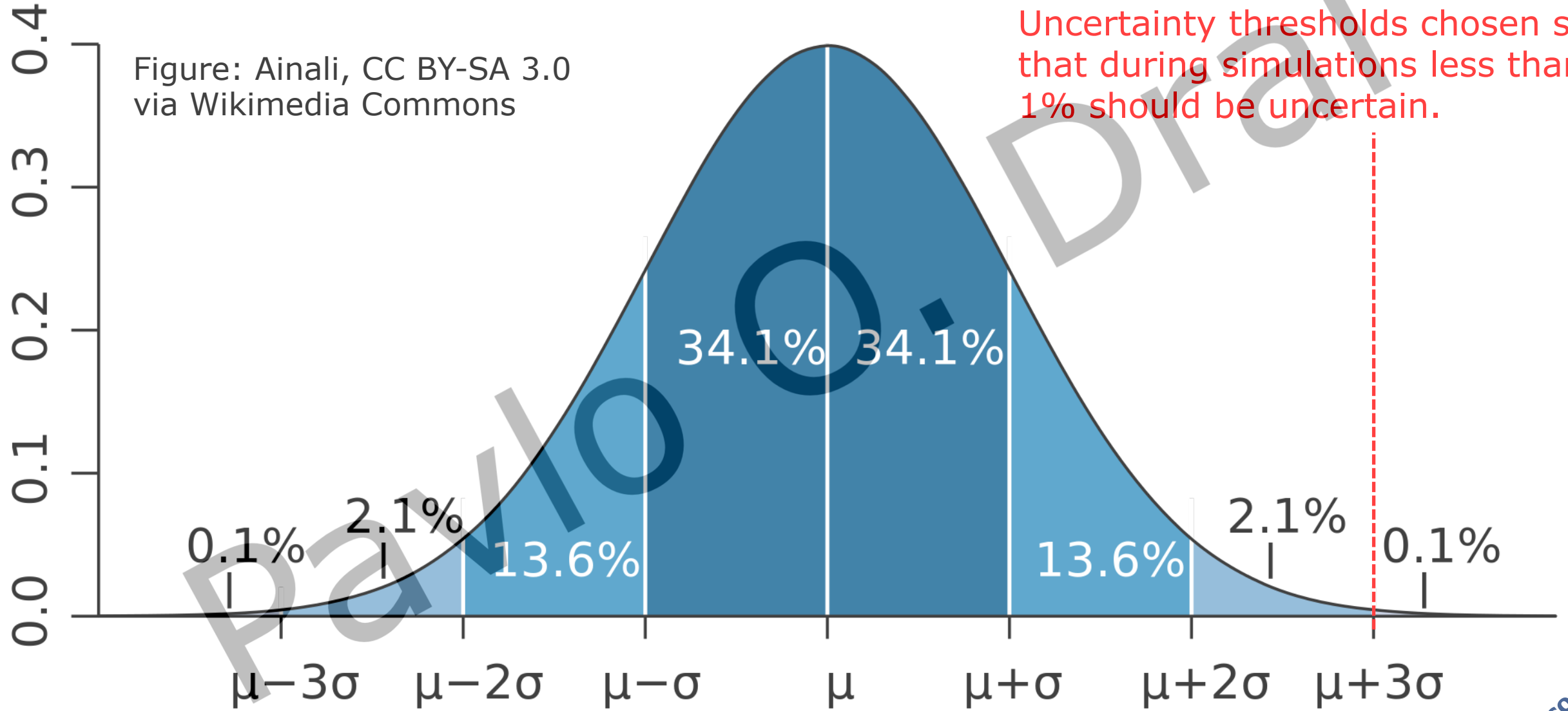
- HF, MP2, CC  
ADC(2), CASSCF  
...
- B3LYP  
ωB97x  
...
- PM6, GFN2-xTB  
AM1, PM3, OMx  
...
- AIQM1  
AIQM1@DFT  
AIQM1@DFT\*
- ANI-1x, ANI-2x  
ANI-1ccx  
...
- ANI, PhysNet, DPMD  
DeepPot-SE, MACE  
...
- KREG, sGDML,  
KRR-CM, GAP-SOAP  
...

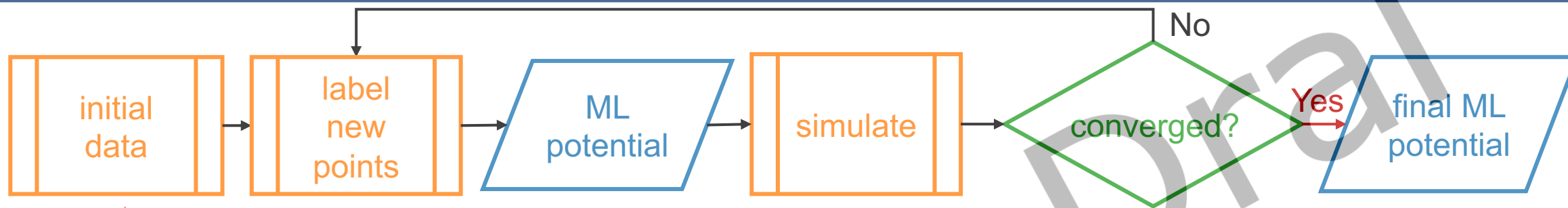


Pavlo O. Dral

## Utilizing different amount of physics-derived information for uncertainty quantification



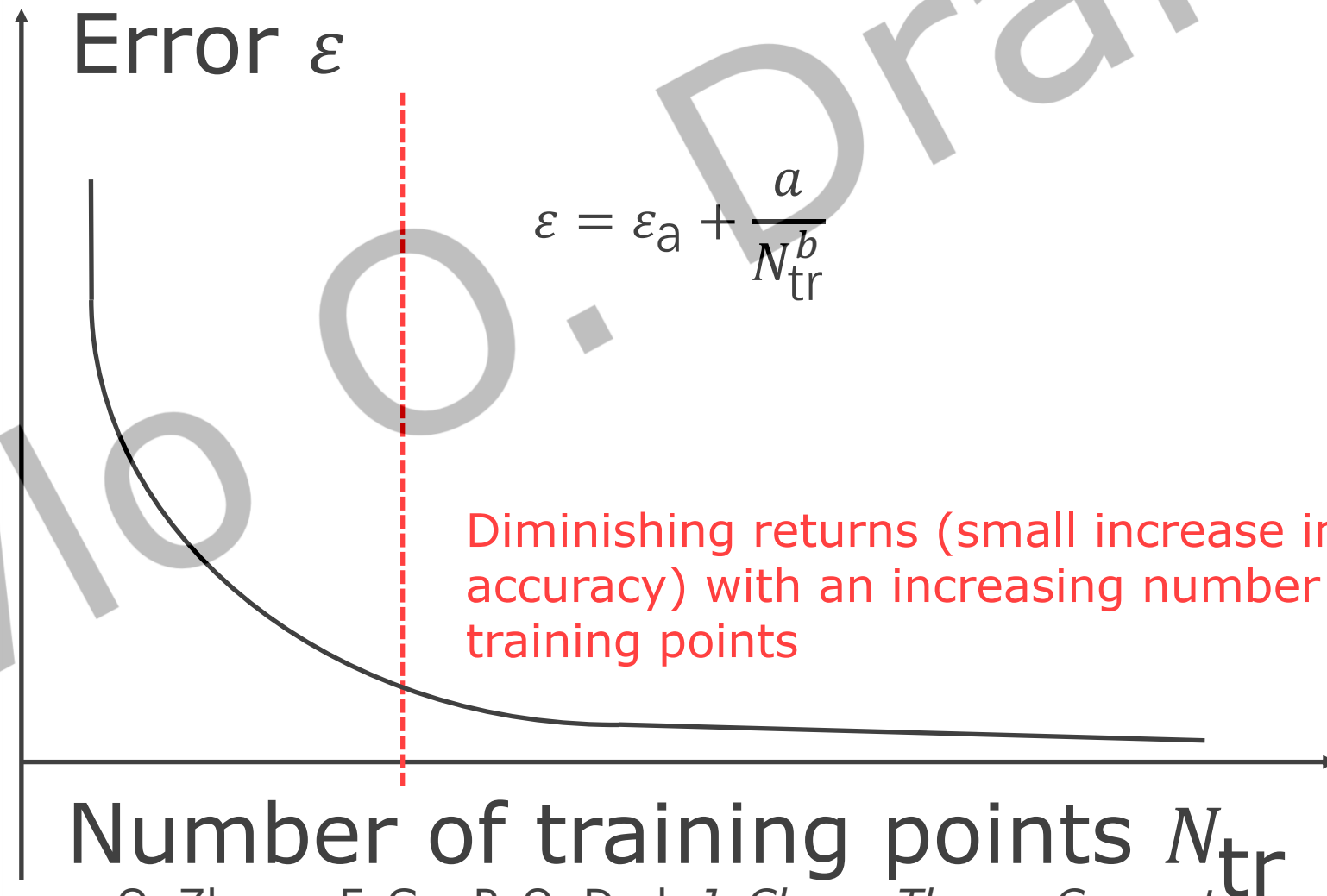
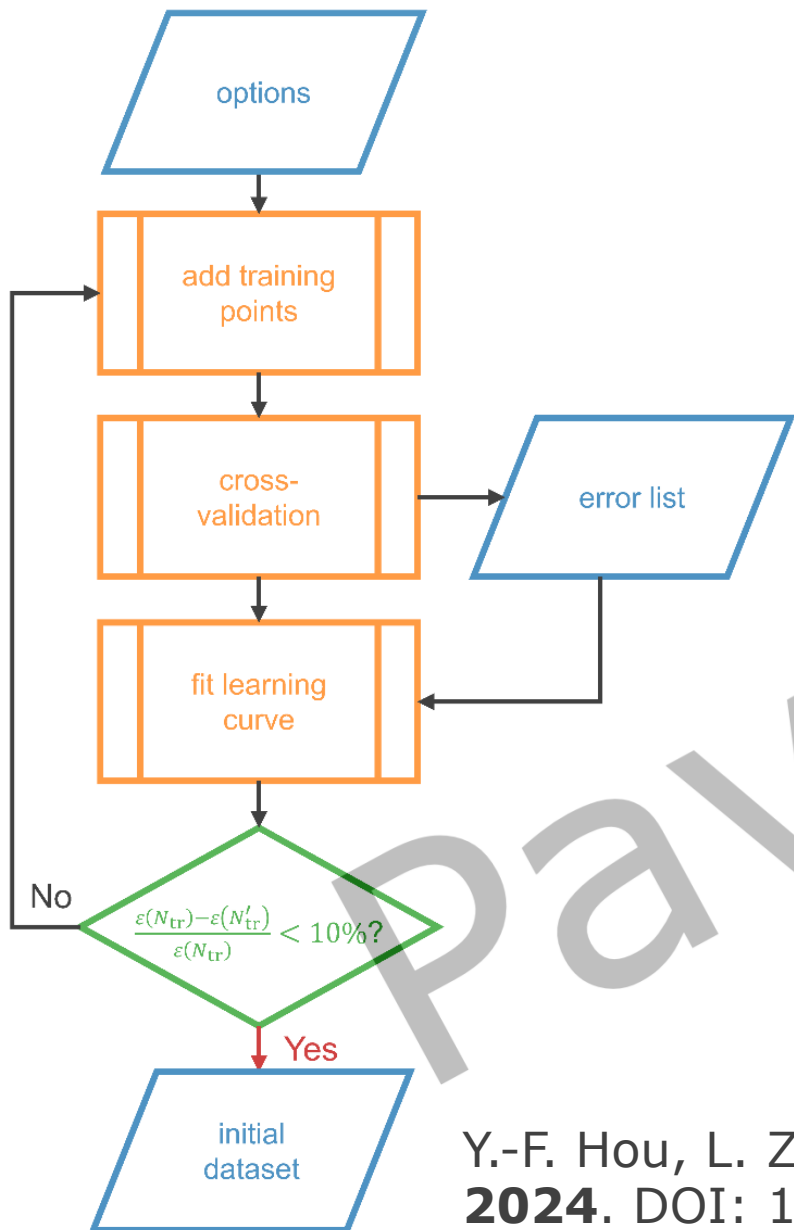




?

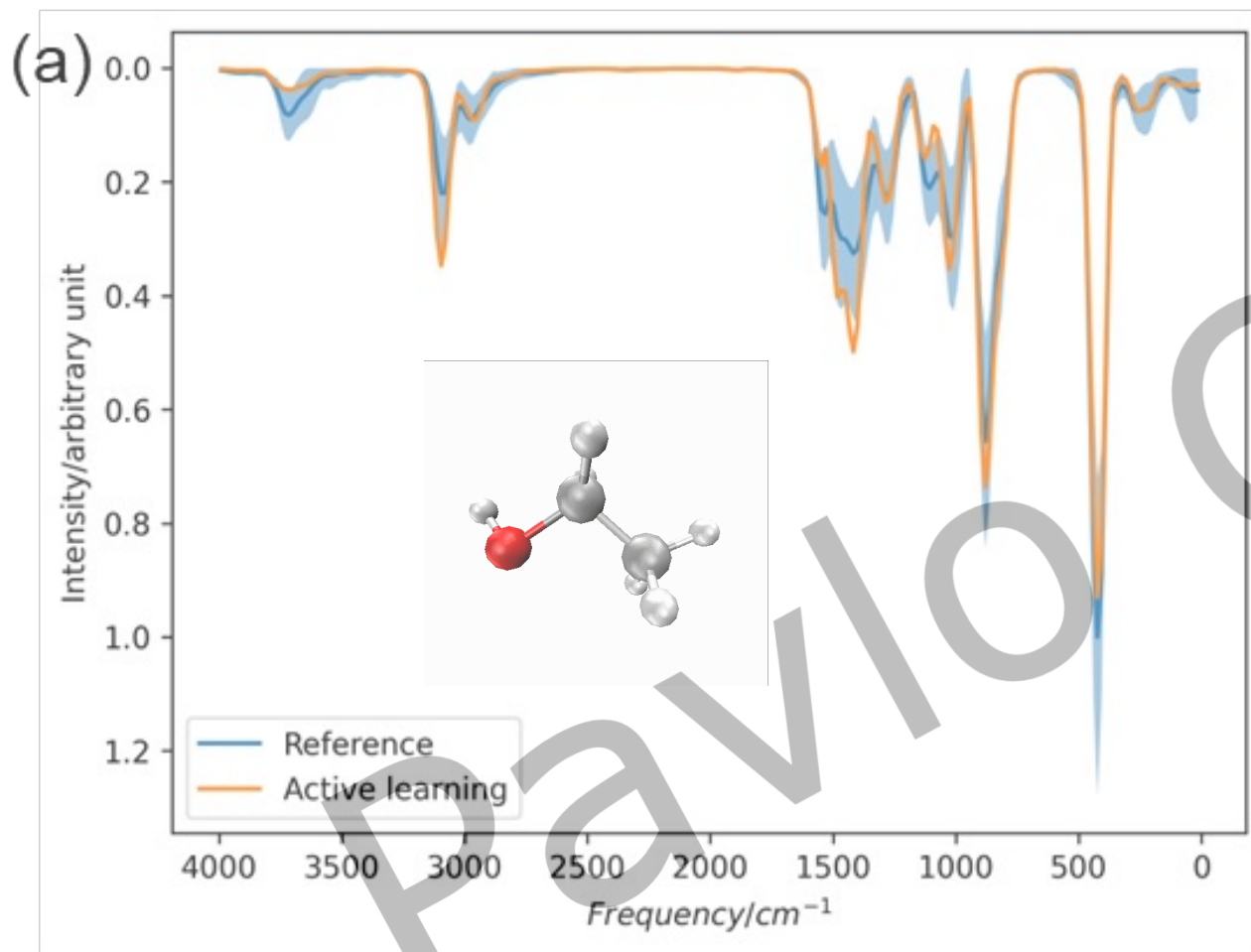
Pavlo O. Dral



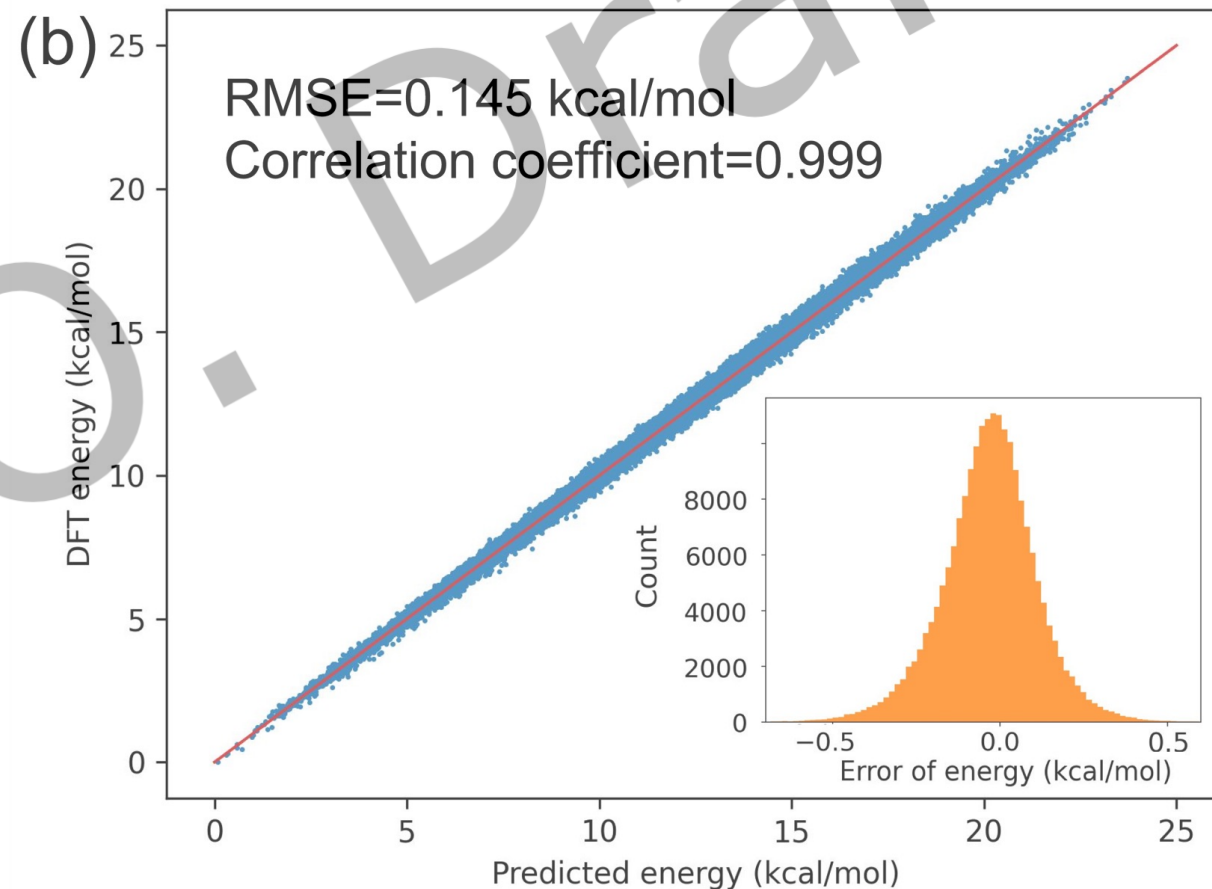


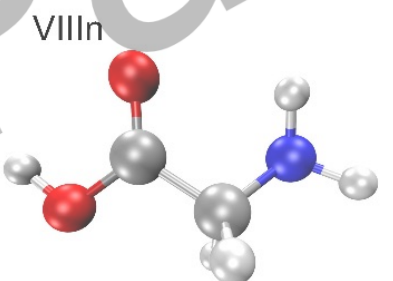
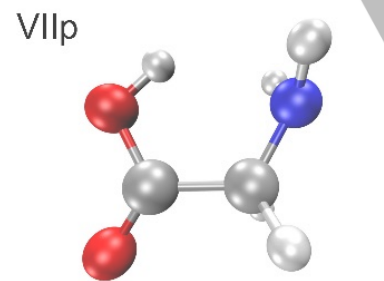
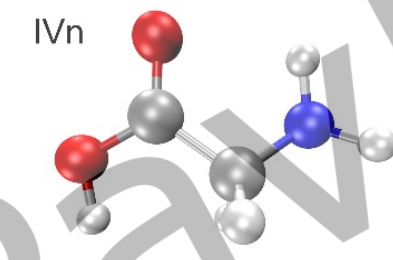
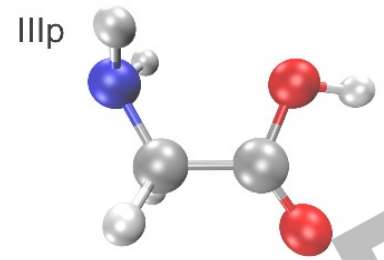
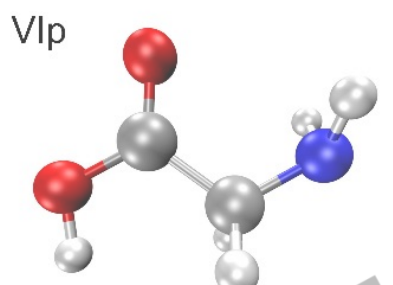
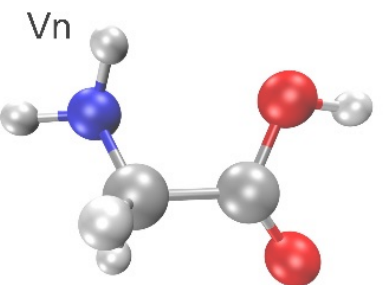
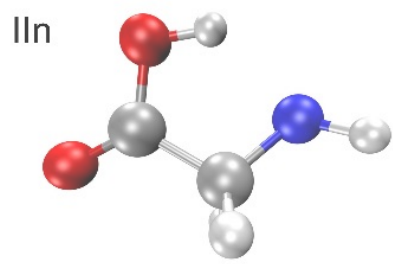
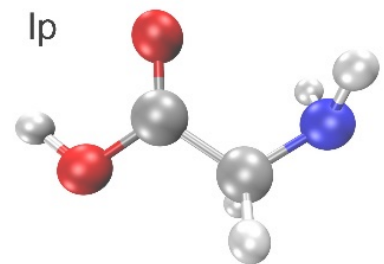
Diminishing returns (small increase in accuracy) with an increasing number of training points

< 1000 training points with ANI potential



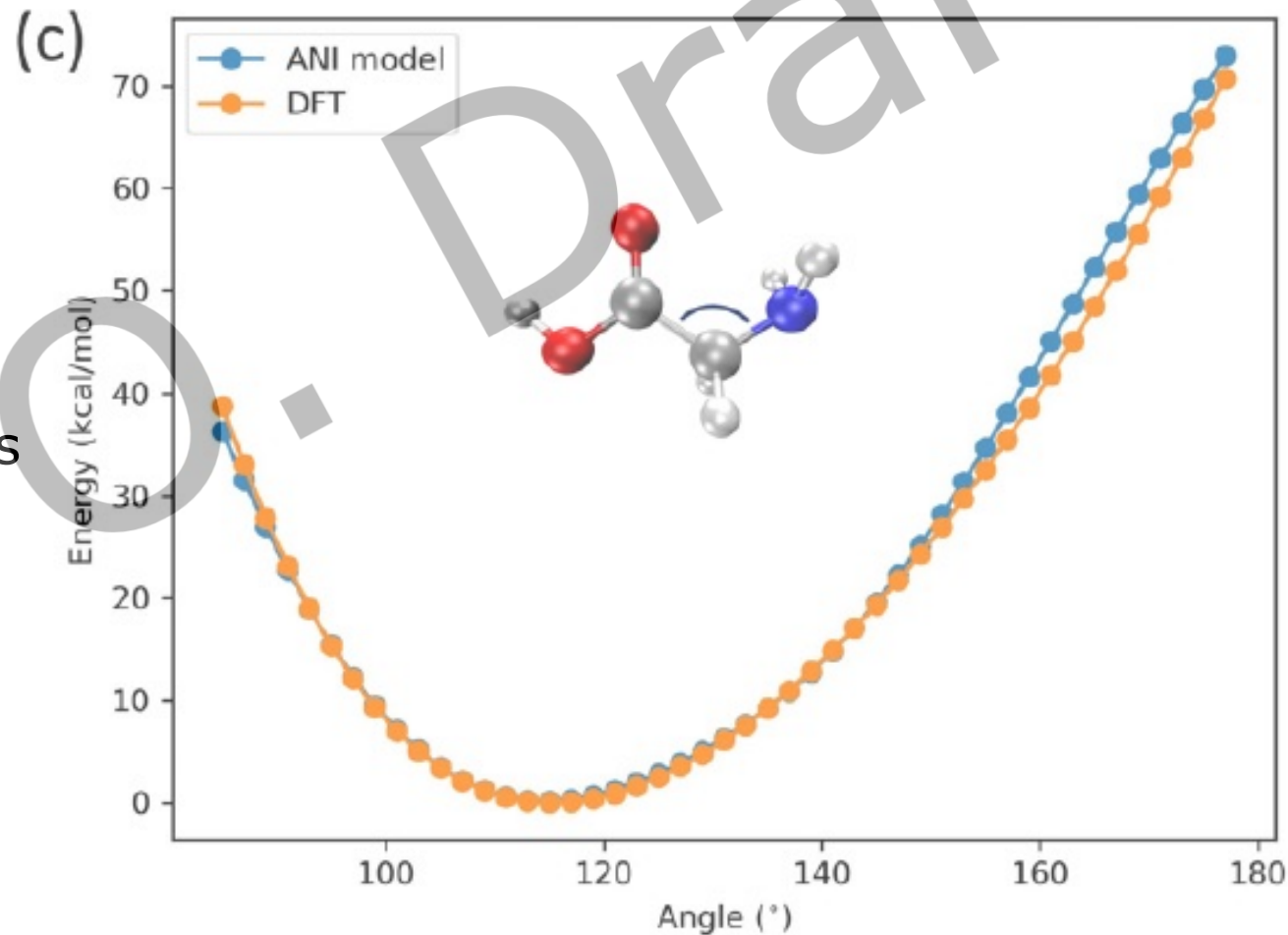
Ethanol spectra





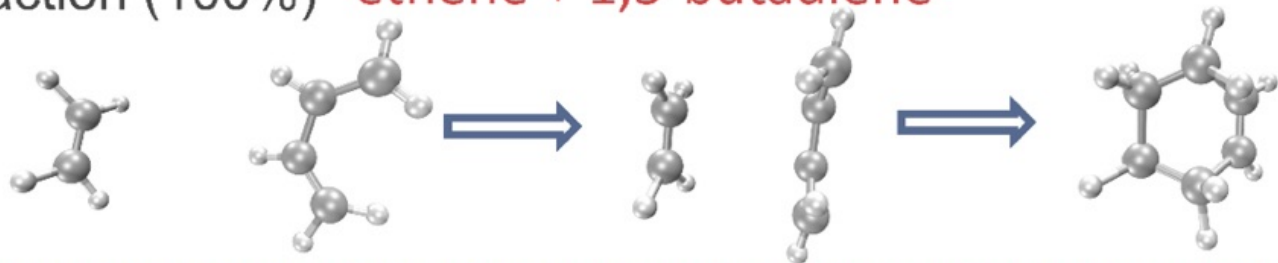
< 2000 training points with ANI potential

Glycine  
8  
conformers

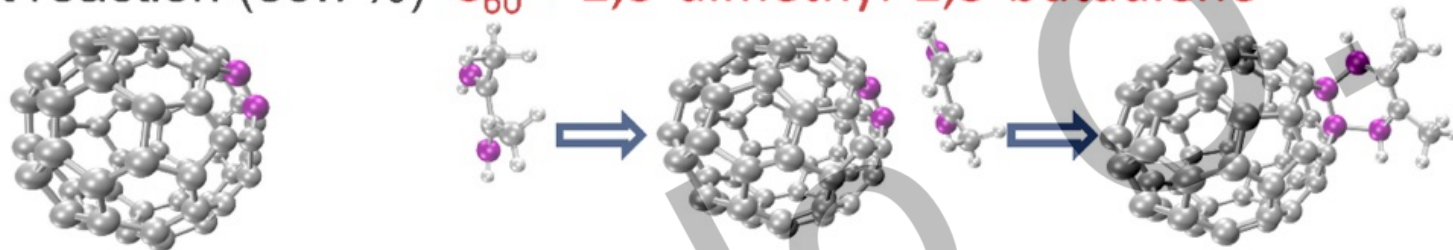


Direct reaction (100%) ethene + 1,3-butadiene

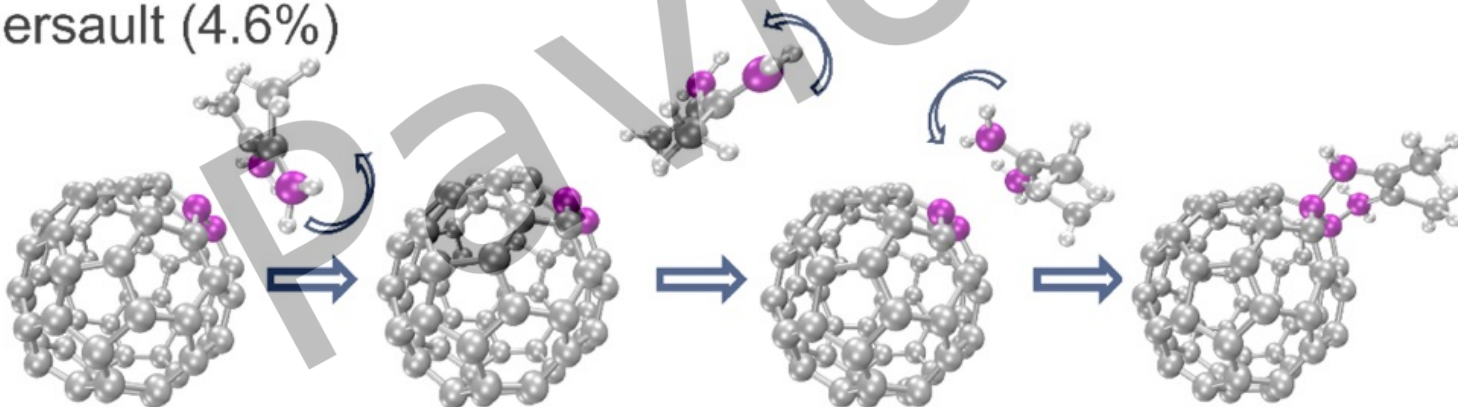
~3000 training points with ANI potential

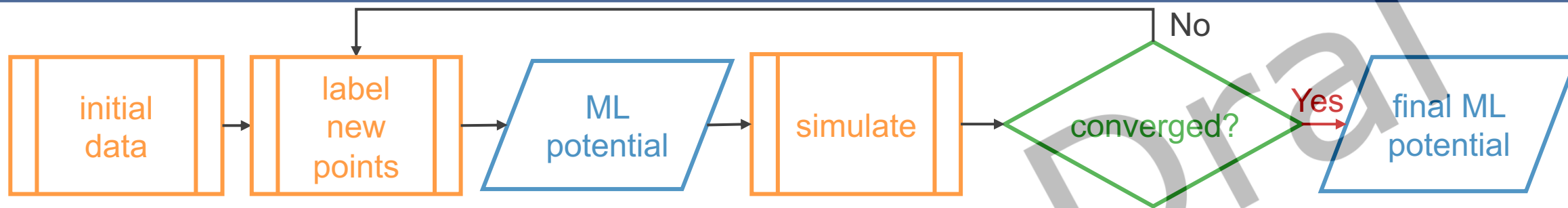


Direct reaction (89.7%)  $C_{60}$  + 2,3-dimethyl-1,3-butadiene



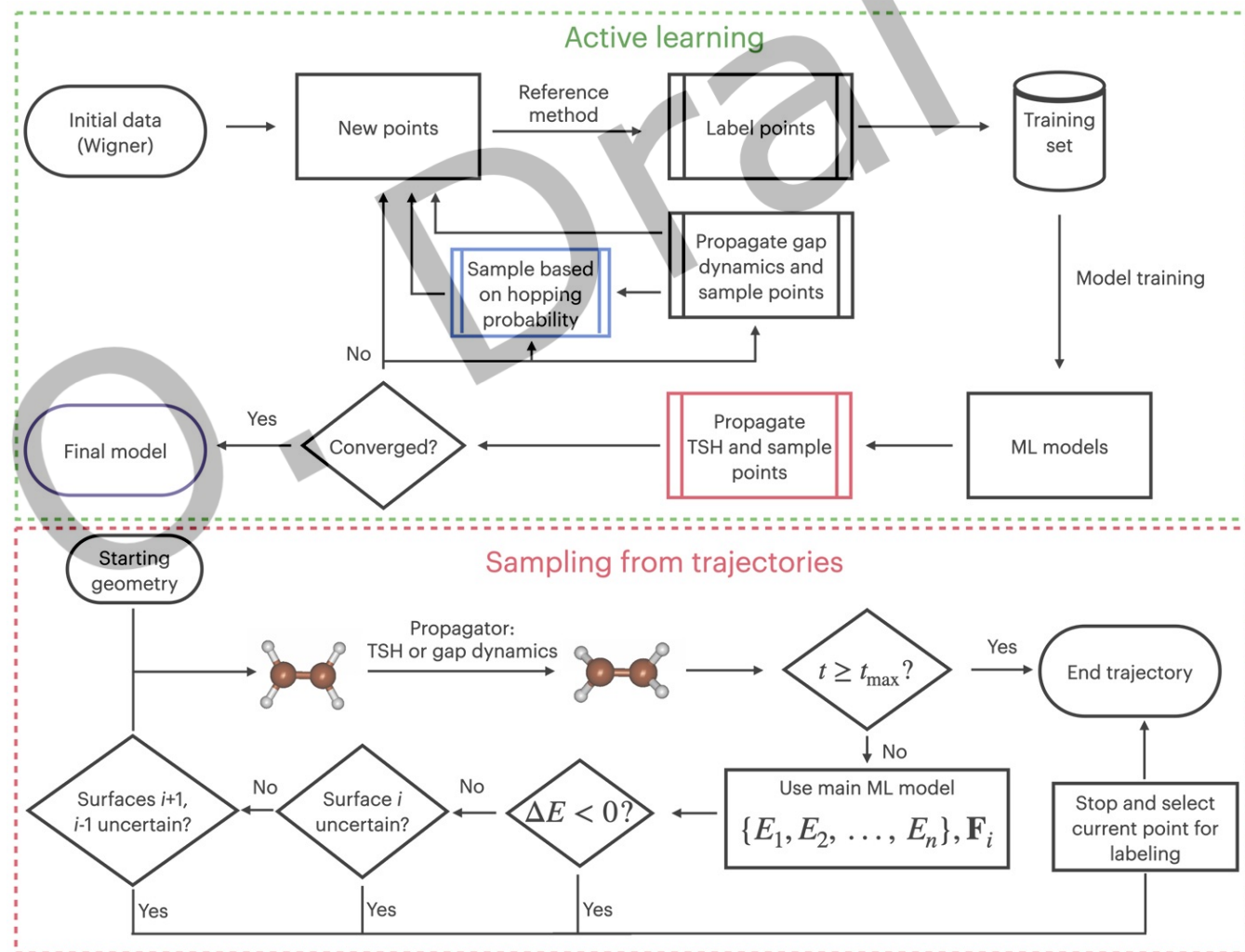
Somersault (4.6%)





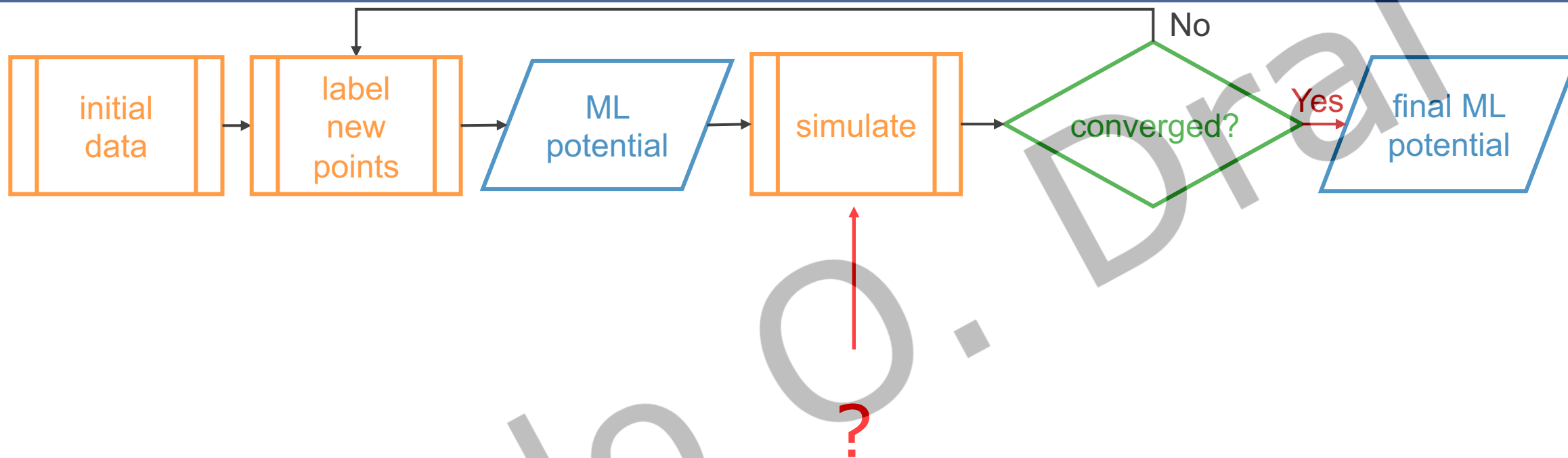
Does it work for NAMMD?

Pavlo O. Dral

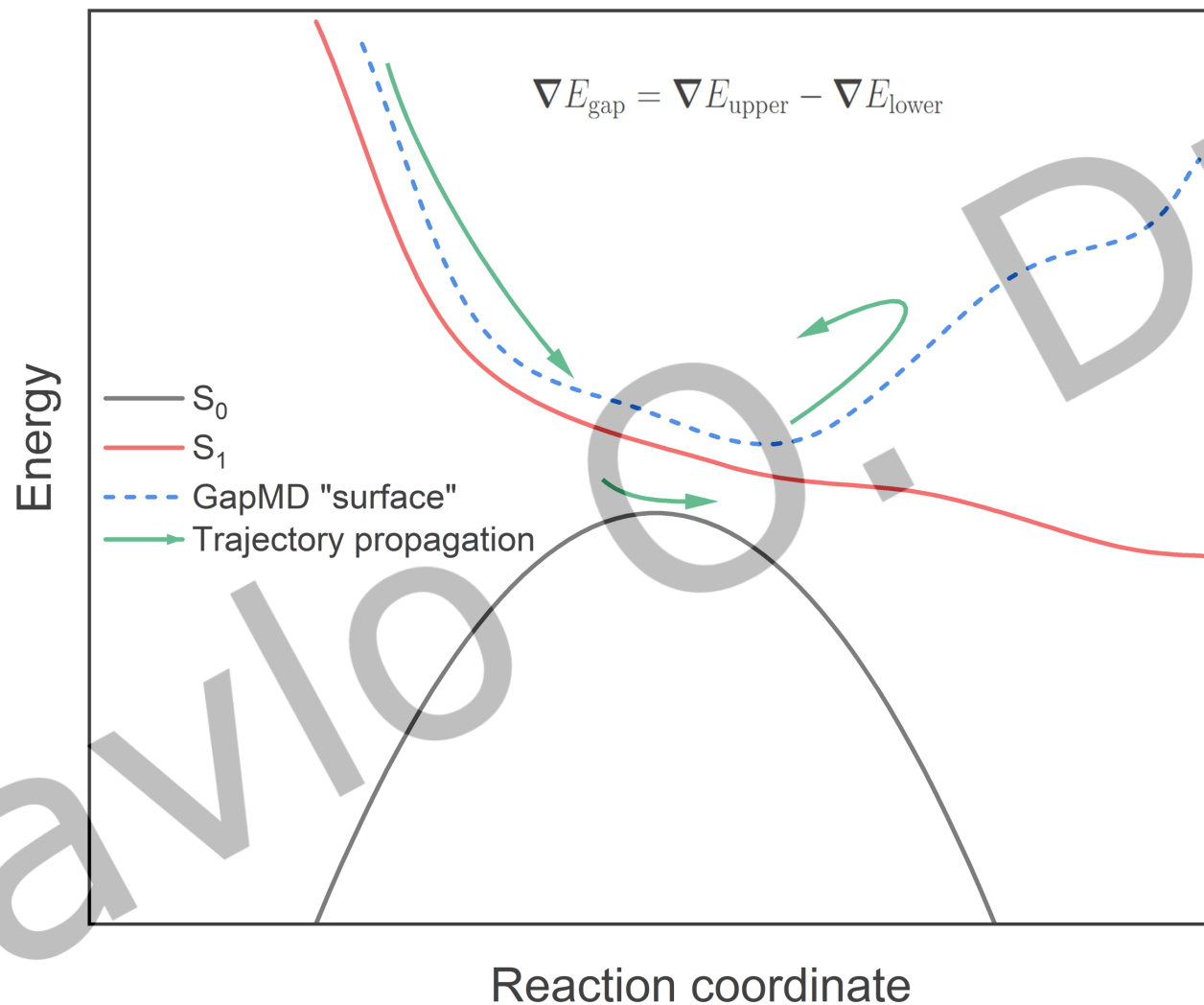


Not so simple for excited states:

Pavlo

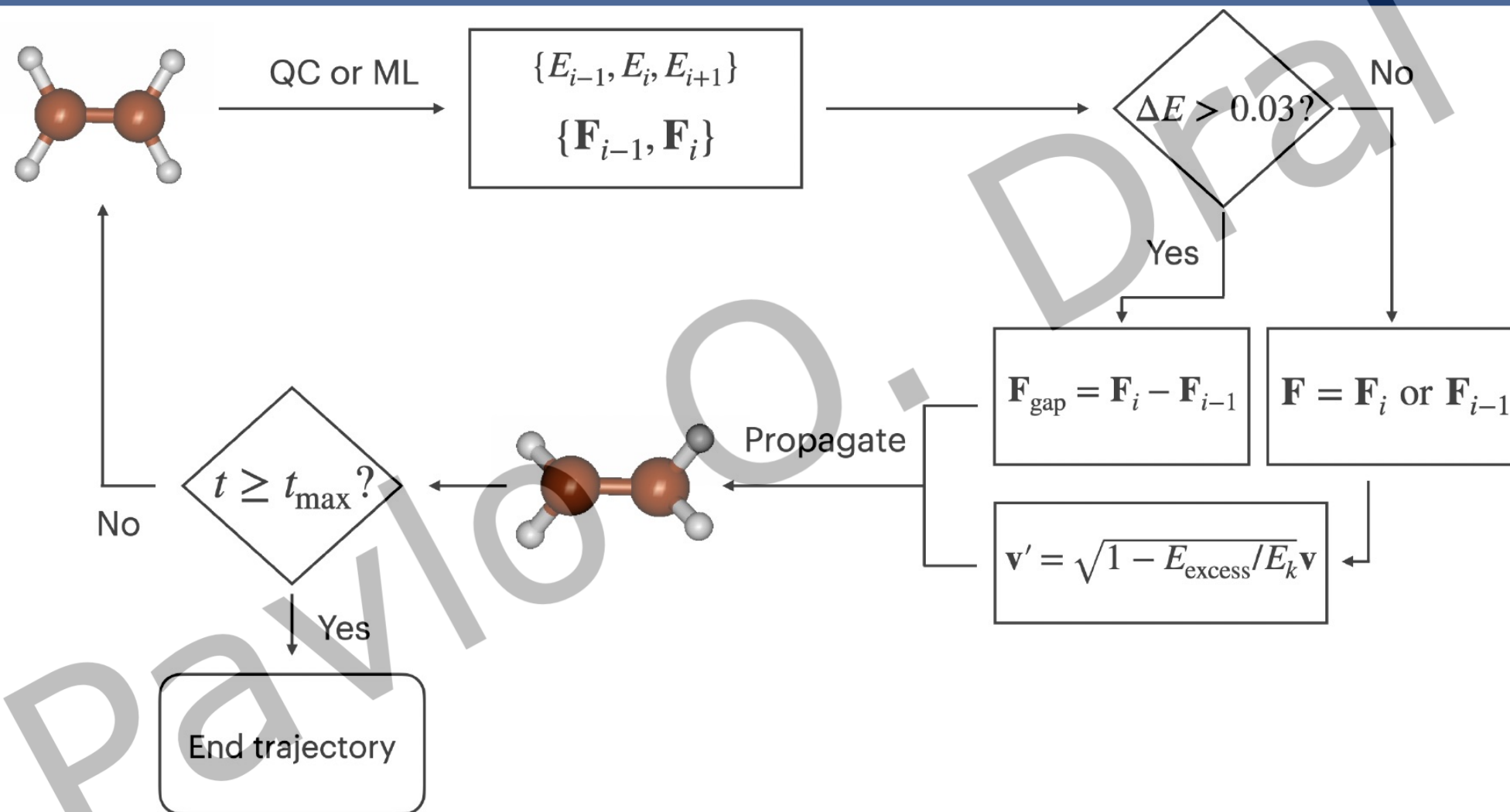


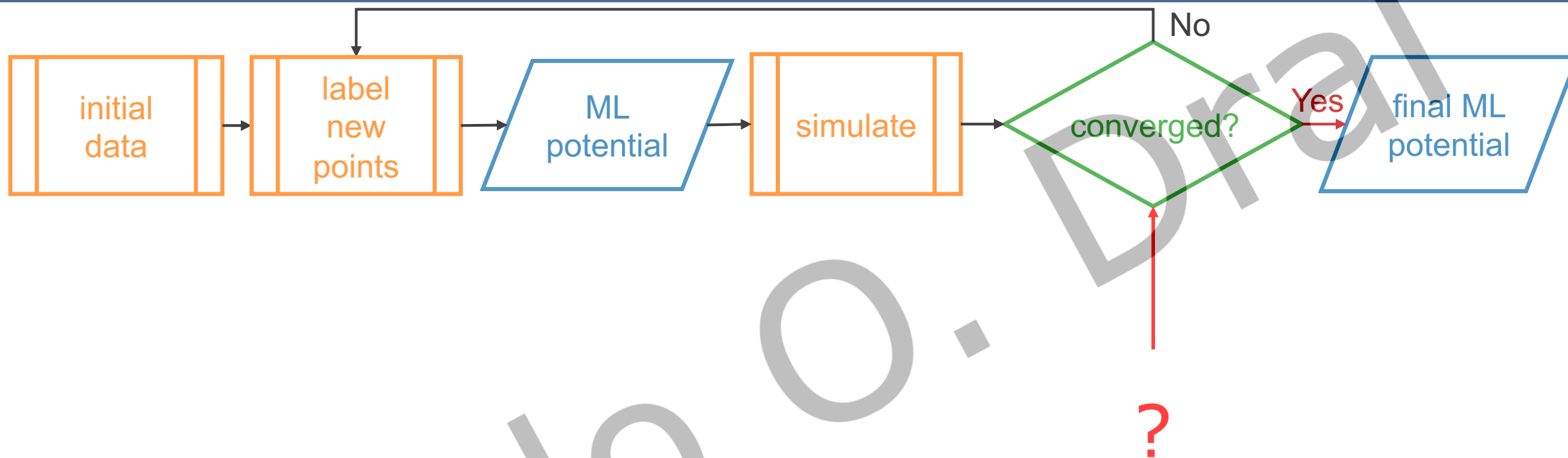
Pavlo



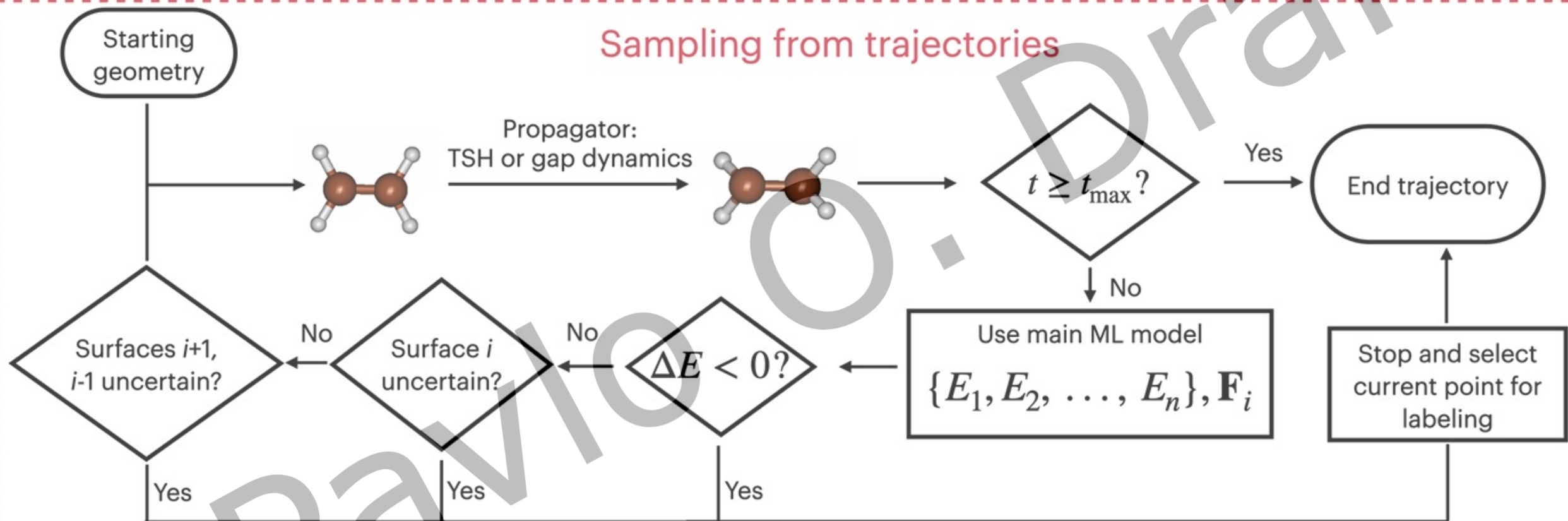
M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

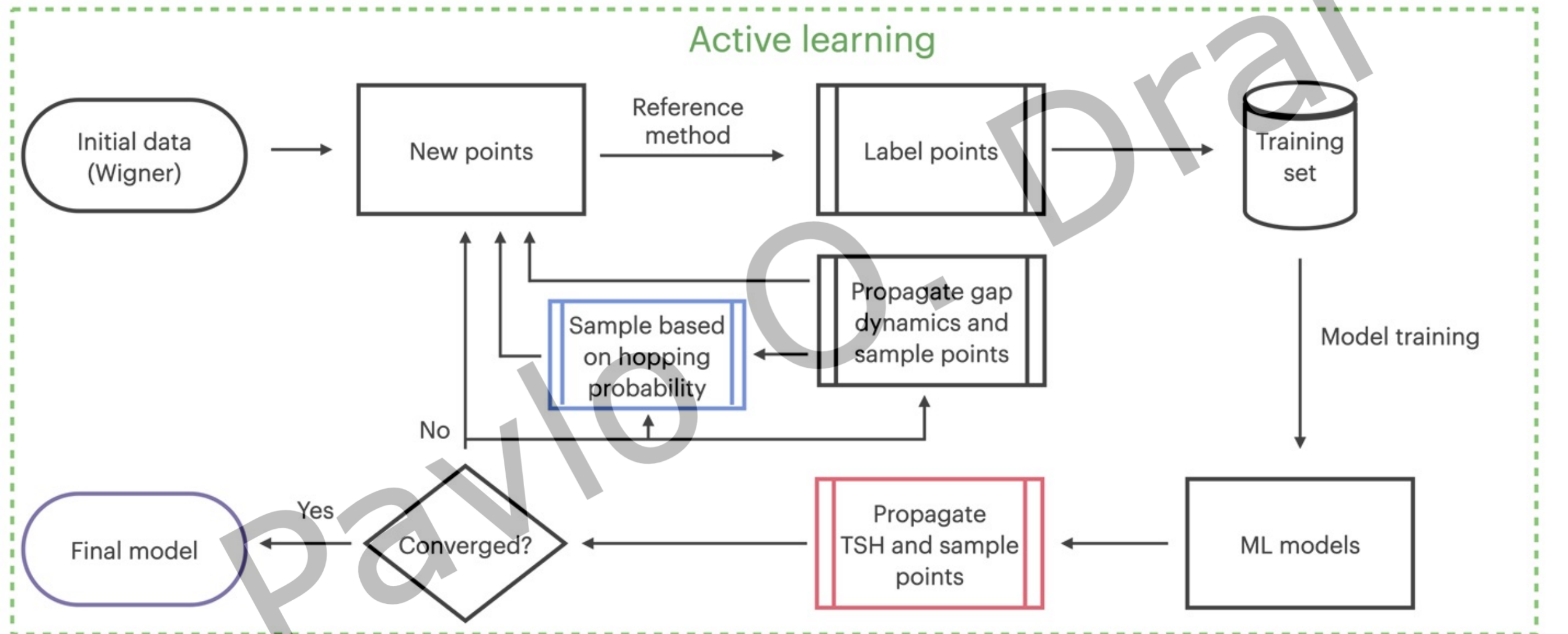






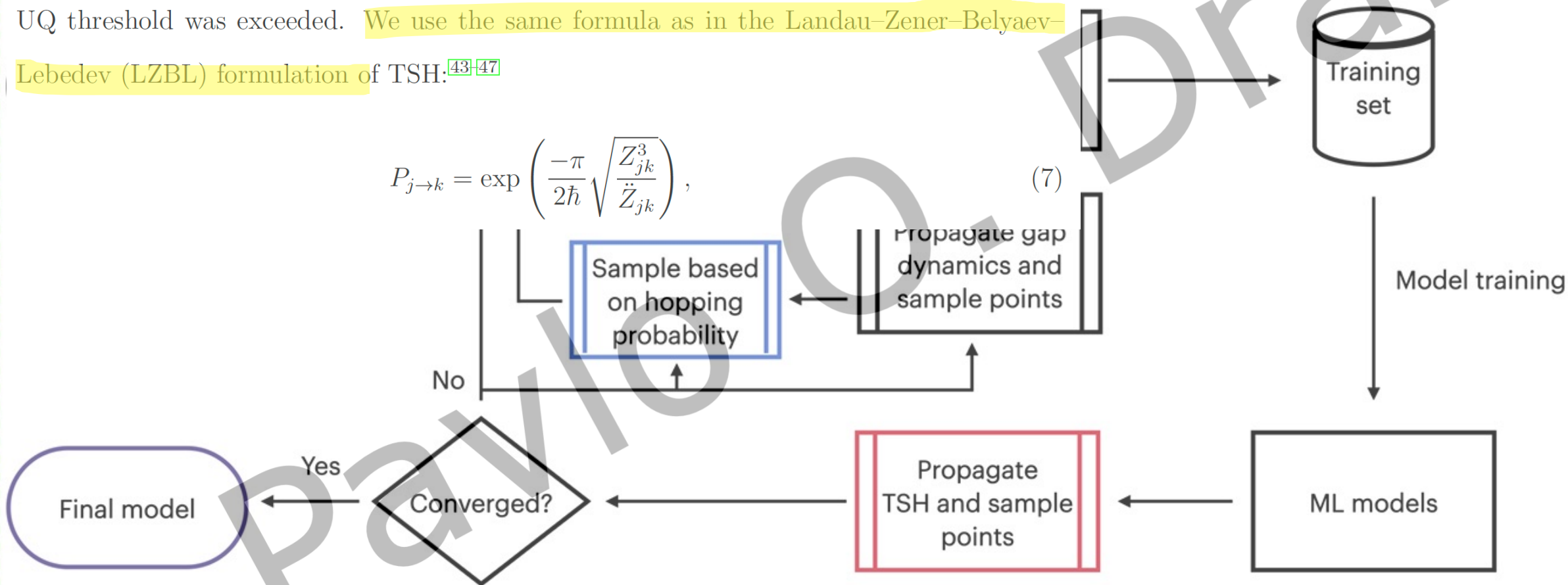
Pavlo O. Dral





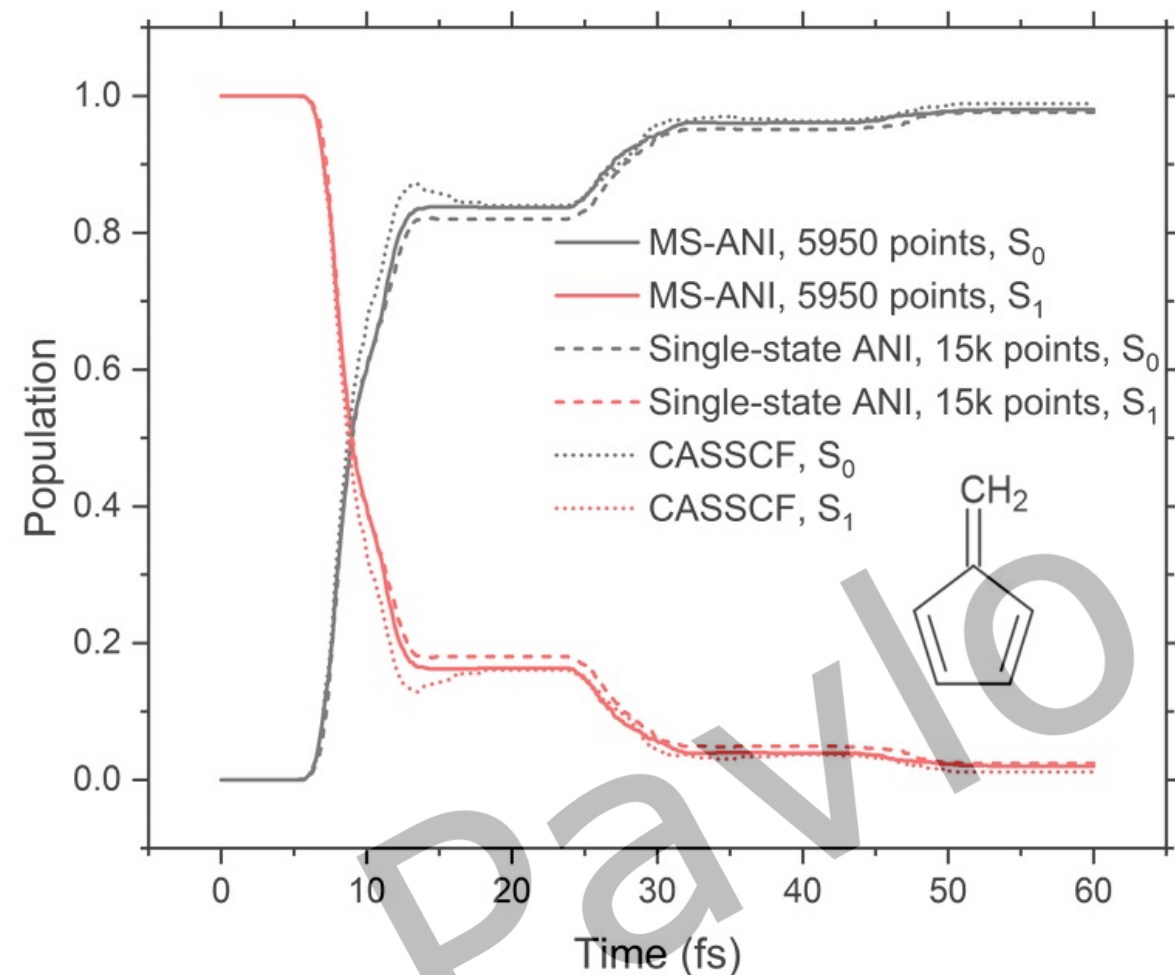
the robust performance in TSH. To further refine the model in terms of hopping probability, after the propagation of each TSH trajectory, we evaluate their uncertainties. The UQ metrics are computed as absolute deviations between the hopping probabilities evaluated using the energies predicted by the main, and auxiliary models at each time step before the UQ threshold was exceeded. We use the same formula as in the Landau-Zener-Belyaev-Lebedev (LZBL) formulation of TSH:<sup>43-47</sup>

$$P_{j \rightarrow k} = \exp\left(\frac{-\pi}{2\hbar} \sqrt{\frac{Z_{jk}^3}{\dot{Z}_{jk}}}\right), \quad (7)$$



dr NAMD





19 iterations of active learning

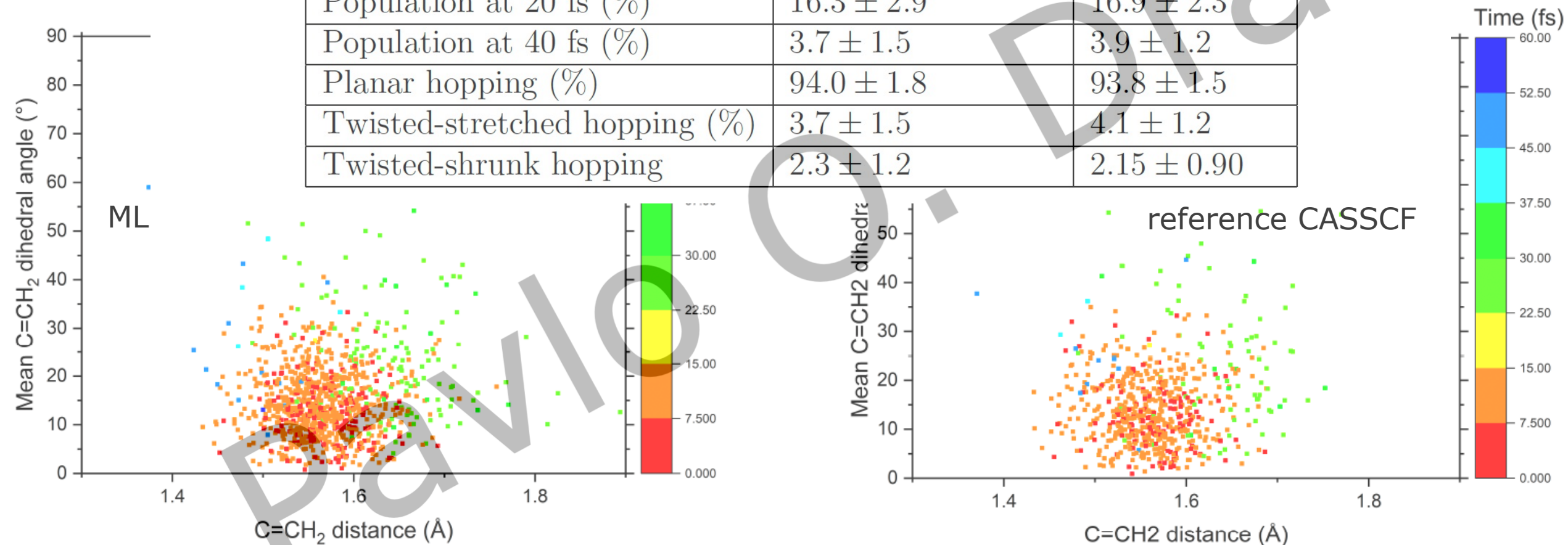
**Three days** on

RTX 4090 GPU and 16 Intel Xeon Gold 6226R CPUs for the entire procedure with trainings and labeling

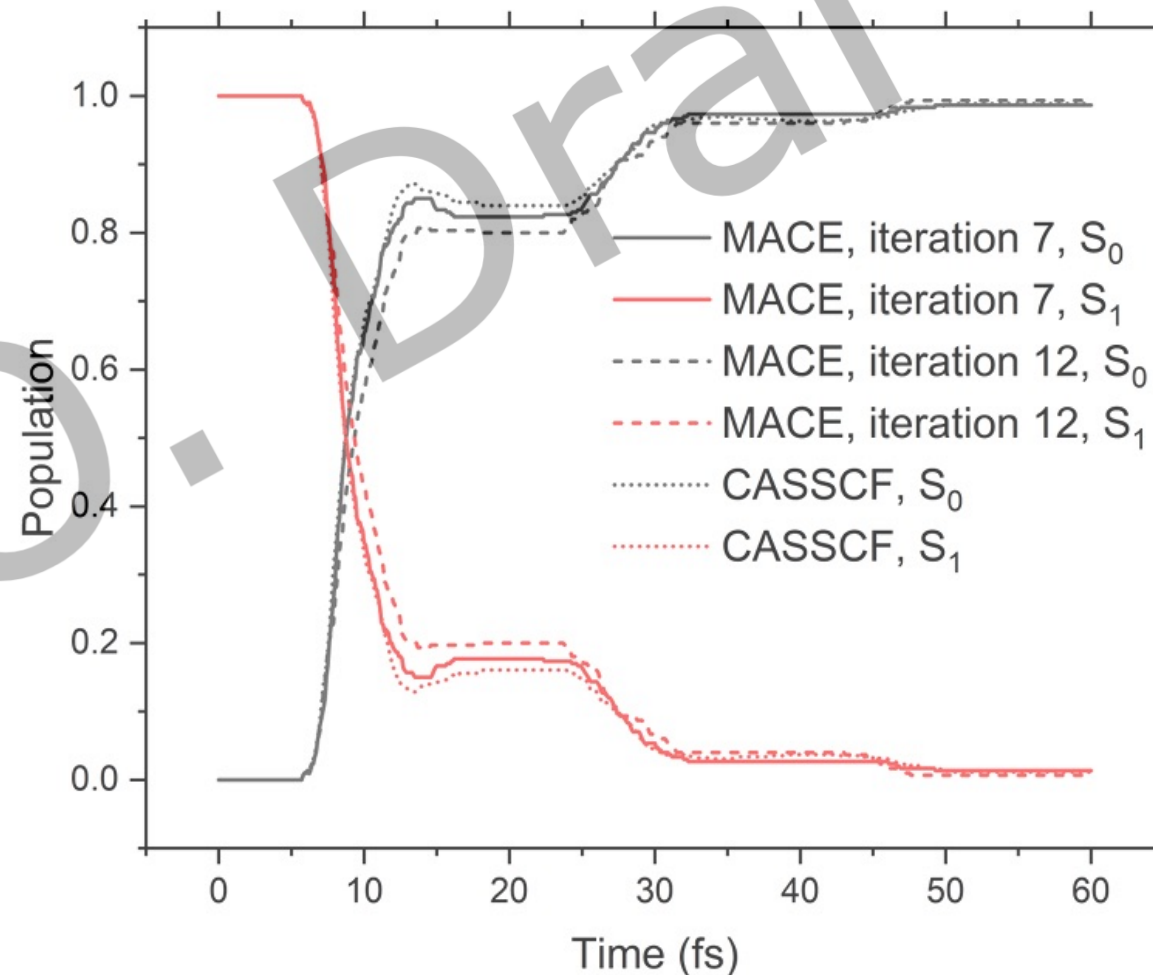
M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

Table 1: Mean value and error bars (95% confidence interval) of observables describing the deactivation channels and kinetics of fulvene for ML dynamics and reference CASSCF dynamics.

Observable	CASSCF dynamics	ML dynamics
Population at 20 fs (%)	$16.3 \pm 2.9$	$16.9 \pm 2.3$
Population at 40 fs (%)	$3.7 \pm 1.5$	$3.9 \pm 1.2$
Planar hopping (%)	$94.0 \pm 1.8$	$93.8 \pm 1.5$
Twisted-stretched hopping (%)	$3.7 \pm 1.5$	$4.1 \pm 1.2$
Twisted-shrunk hopping	$2.3 \pm 1.2$	$2.15 \pm 0.90$

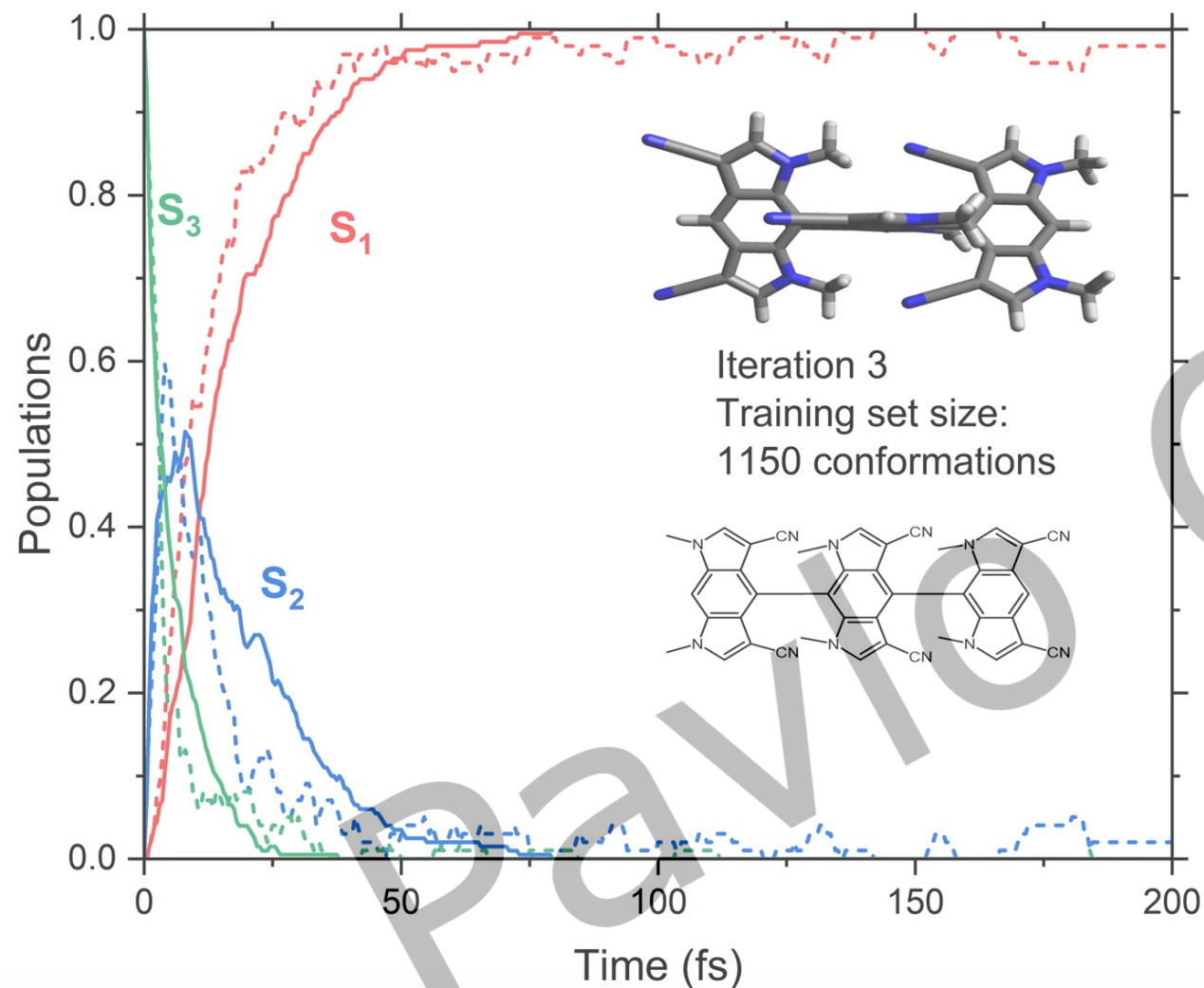


Crashed after 28 days of active learning after 13 iterations and producing 3850 training points

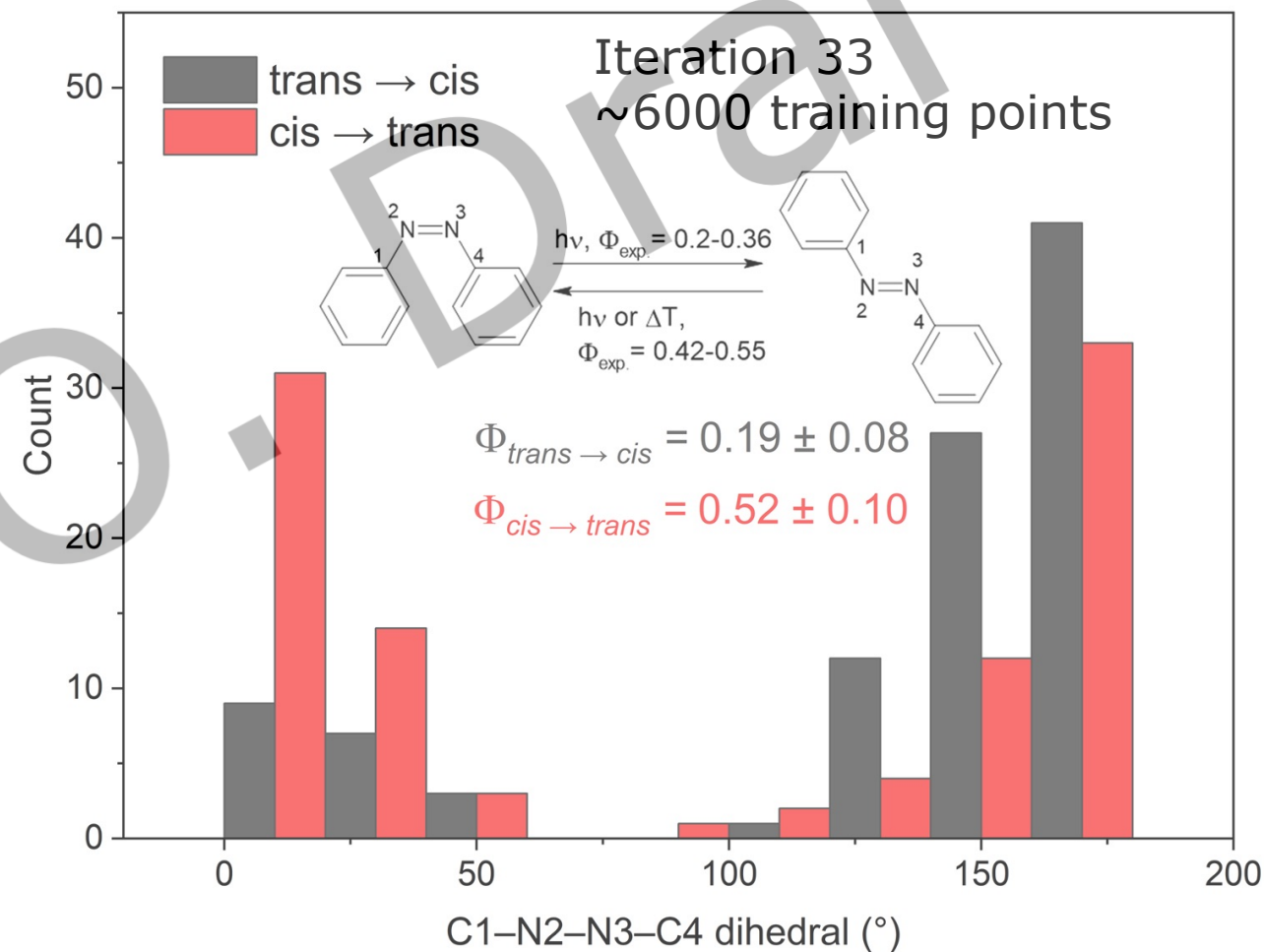
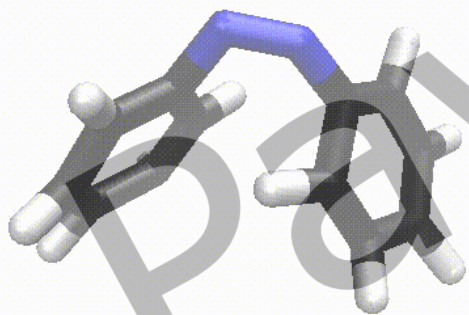
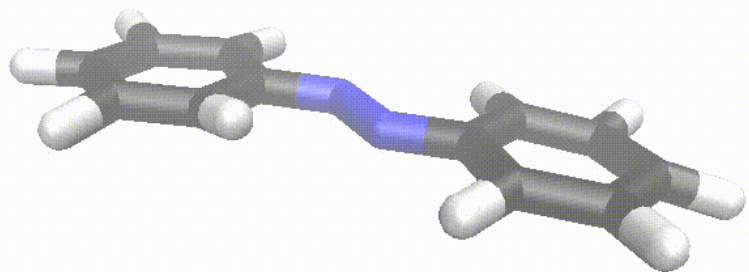


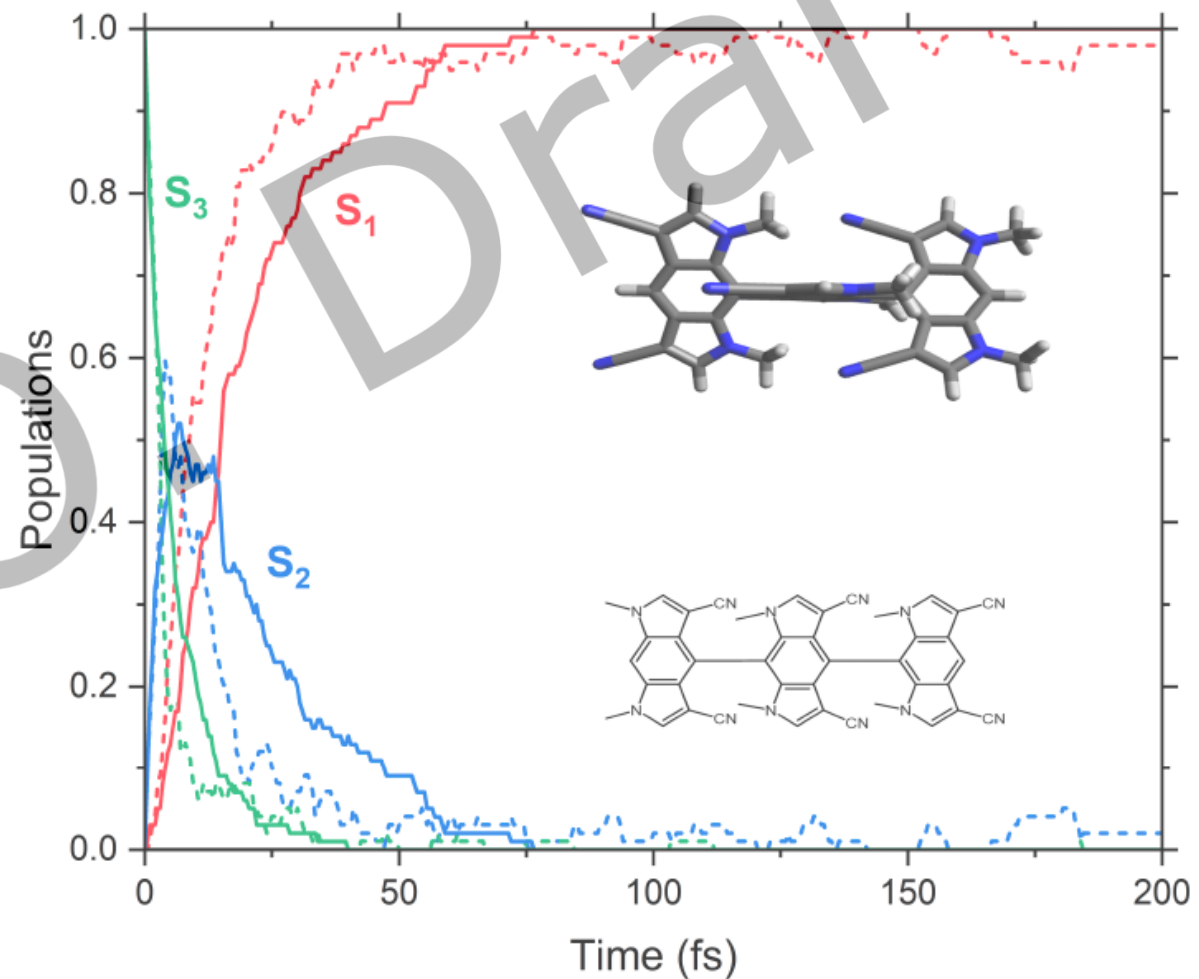
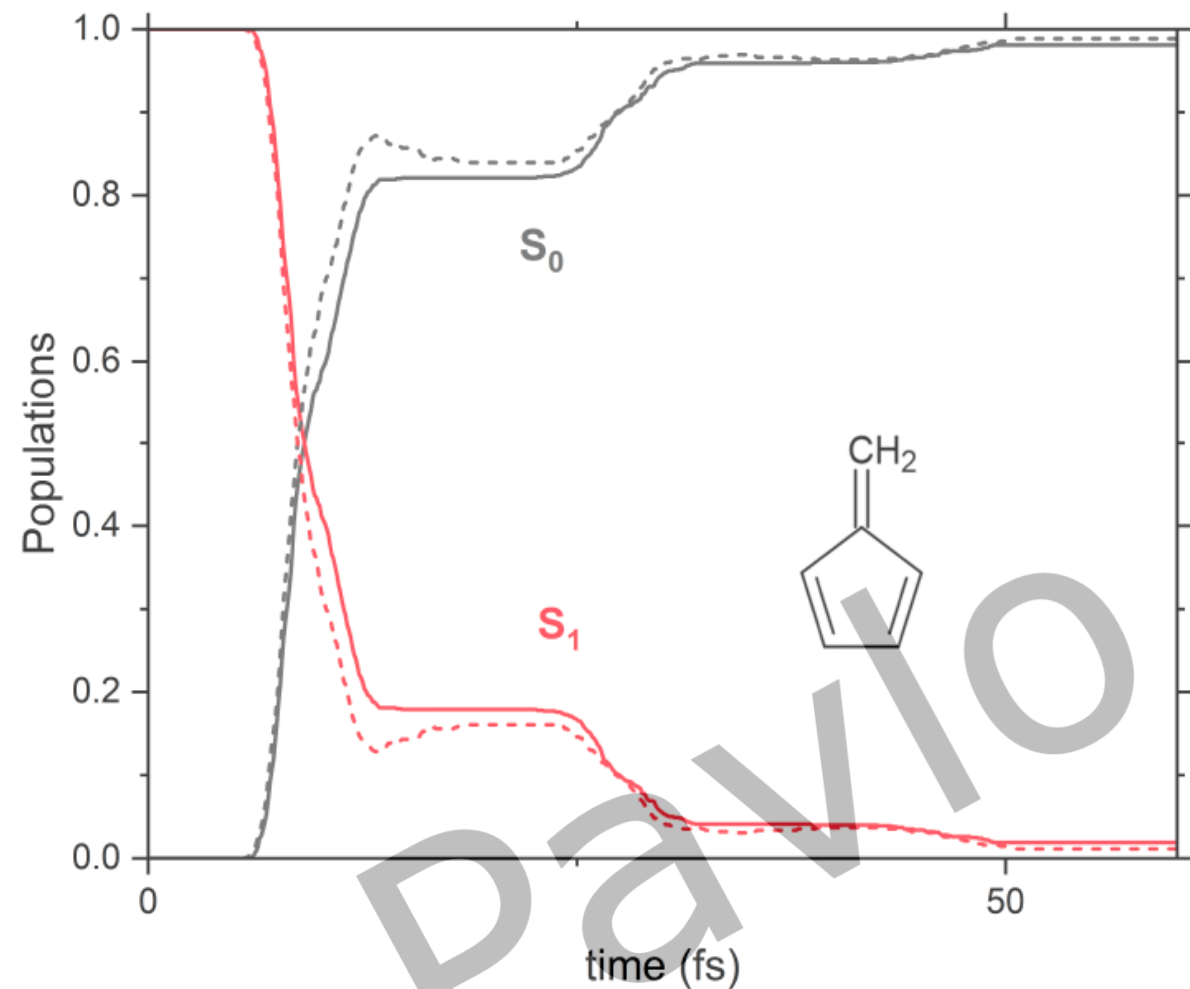
M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>





M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>



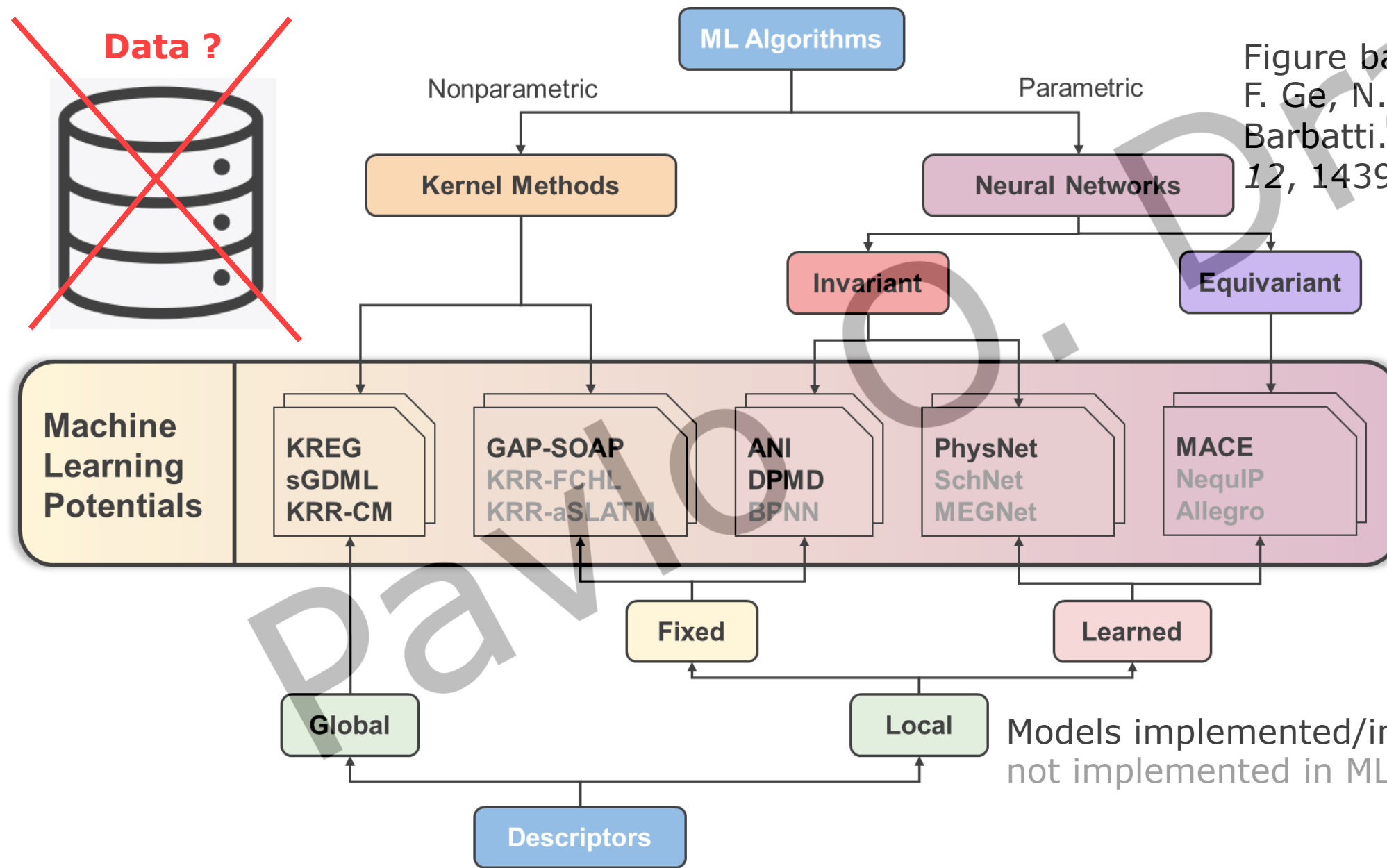


M. Martyka, L. Zhang, F. Ge, Y.-F. Hou, J. Jankowska, M. Barbatti, P. O. Dral. *Charting electronic-state manifolds across molecules with multi-state learning and gap-driven dynamics via efficient and robust active learning.* <https://doi.org/10.26434/chemrxiv-2024-dtc1w>

**Data ?**



Figure based on M. Pinheiro Jr, F. Ge, N. Ferré, P. O. Dral, M. Barbatti. *Chem. Sci.* **2021**, *12*, 14396–14413



Models implemented/interfaced in **MLatom**  
not implemented in MLatom

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

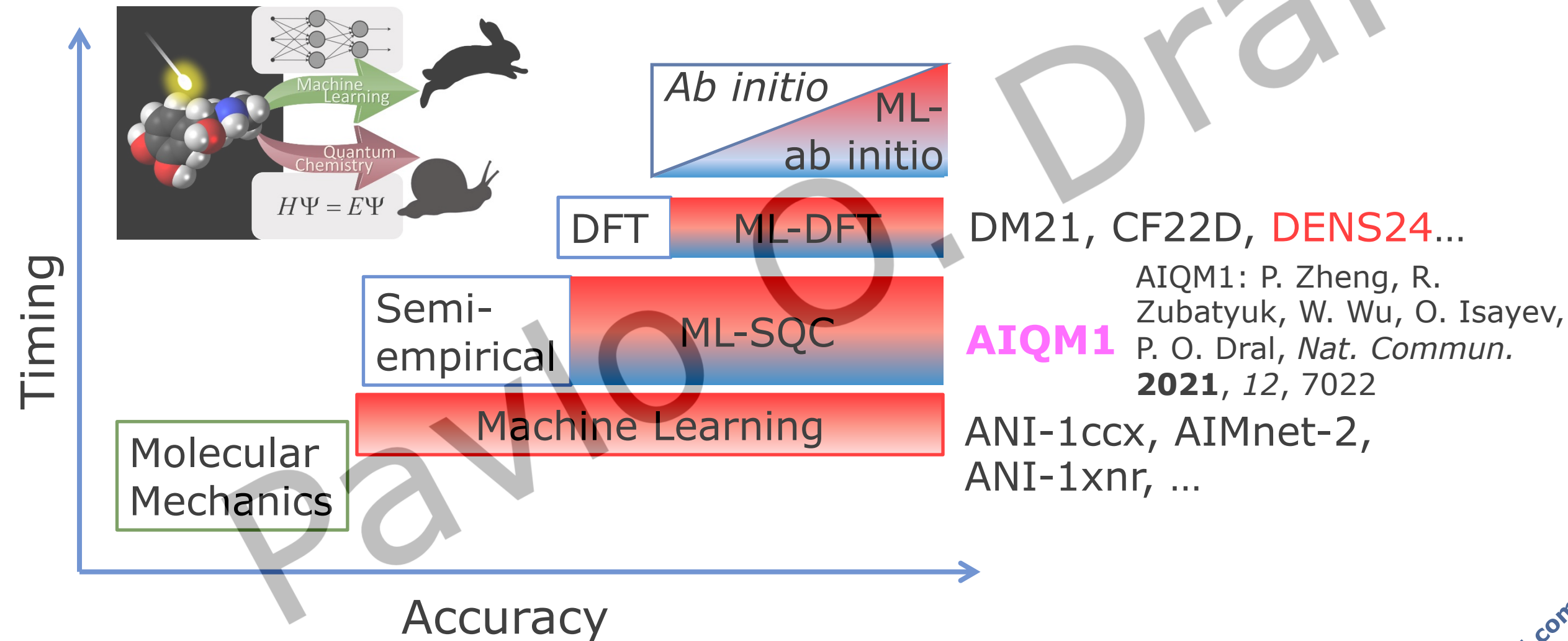
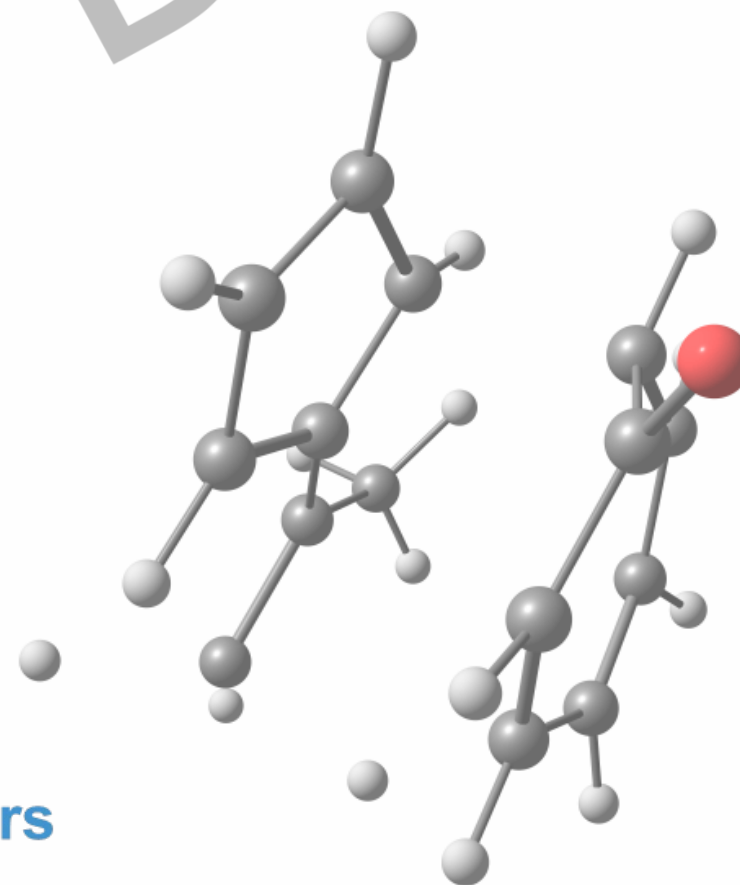
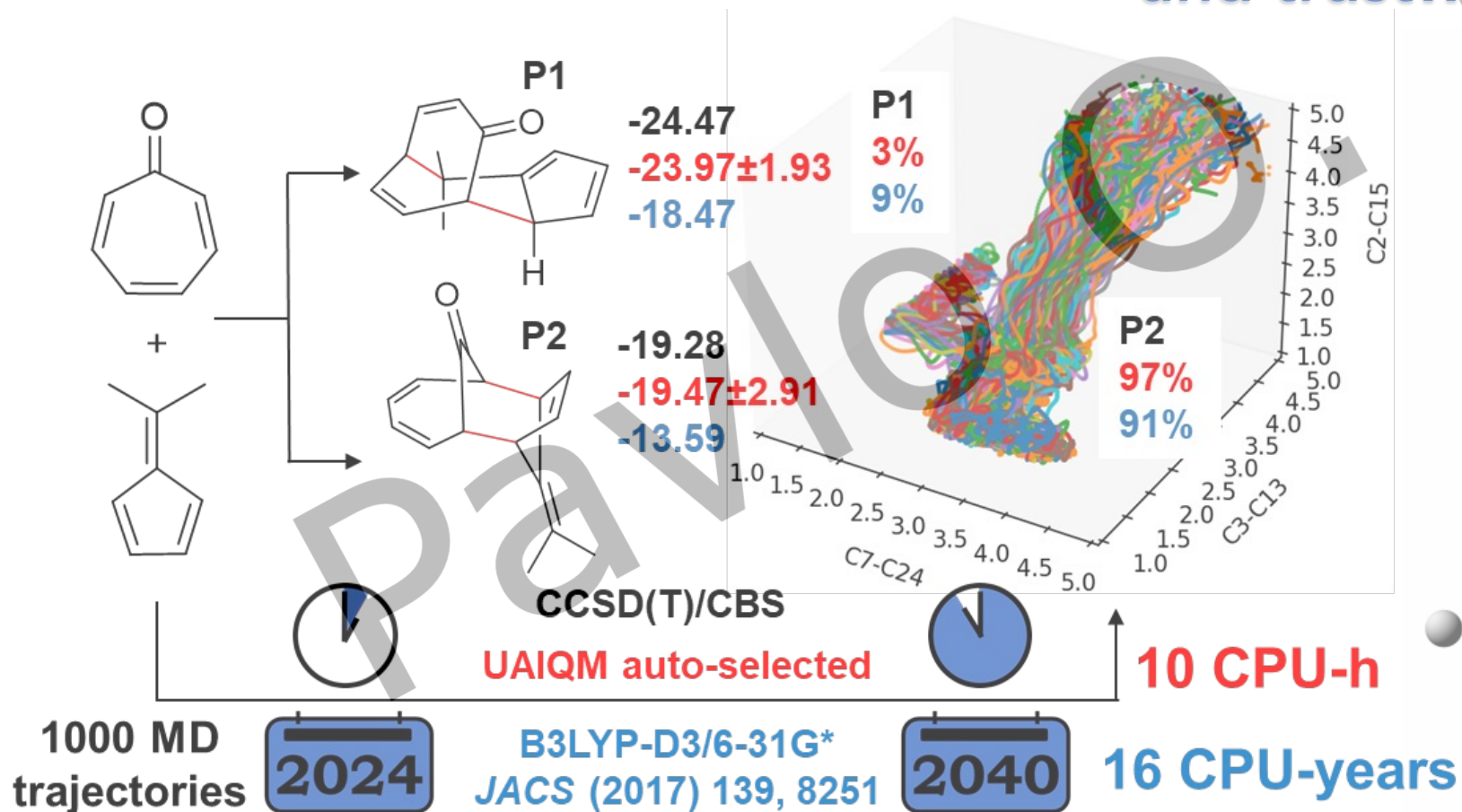


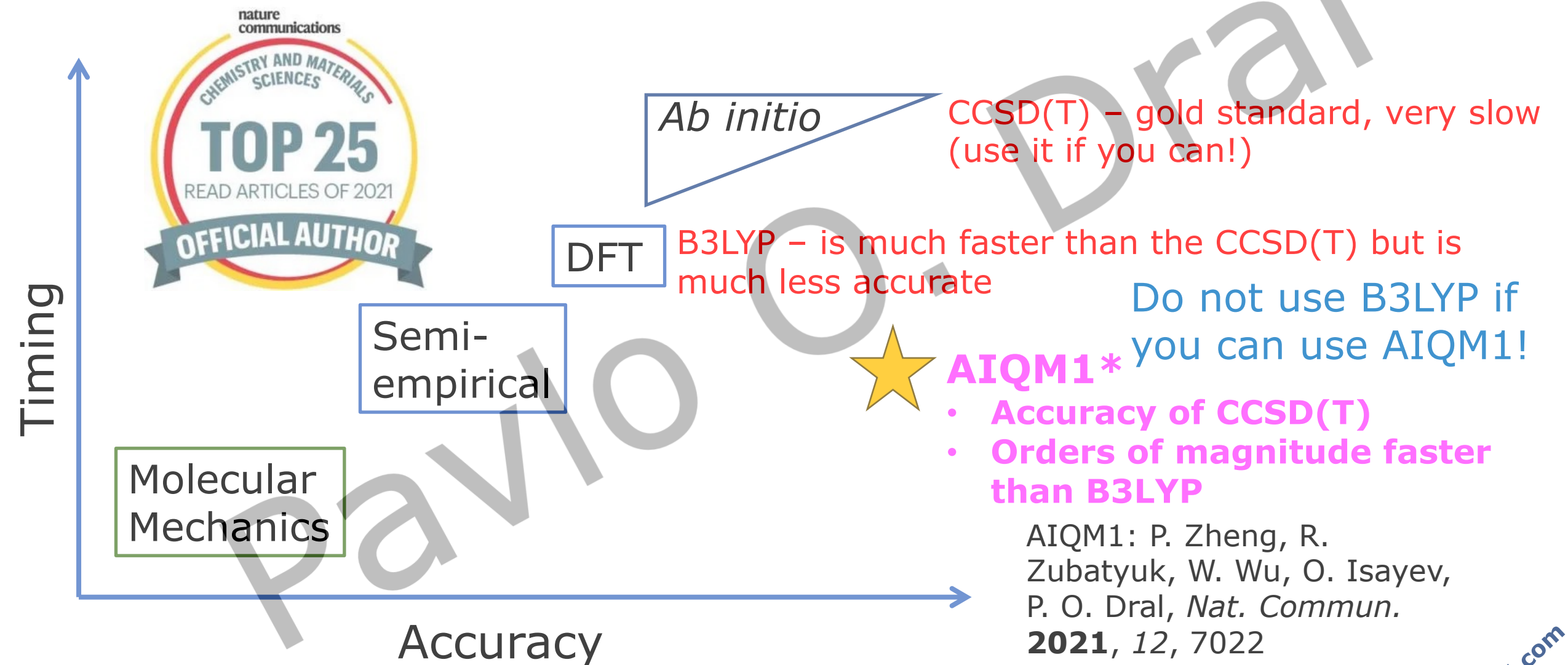
Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

*No reason to do non-ML computational chemistry!*

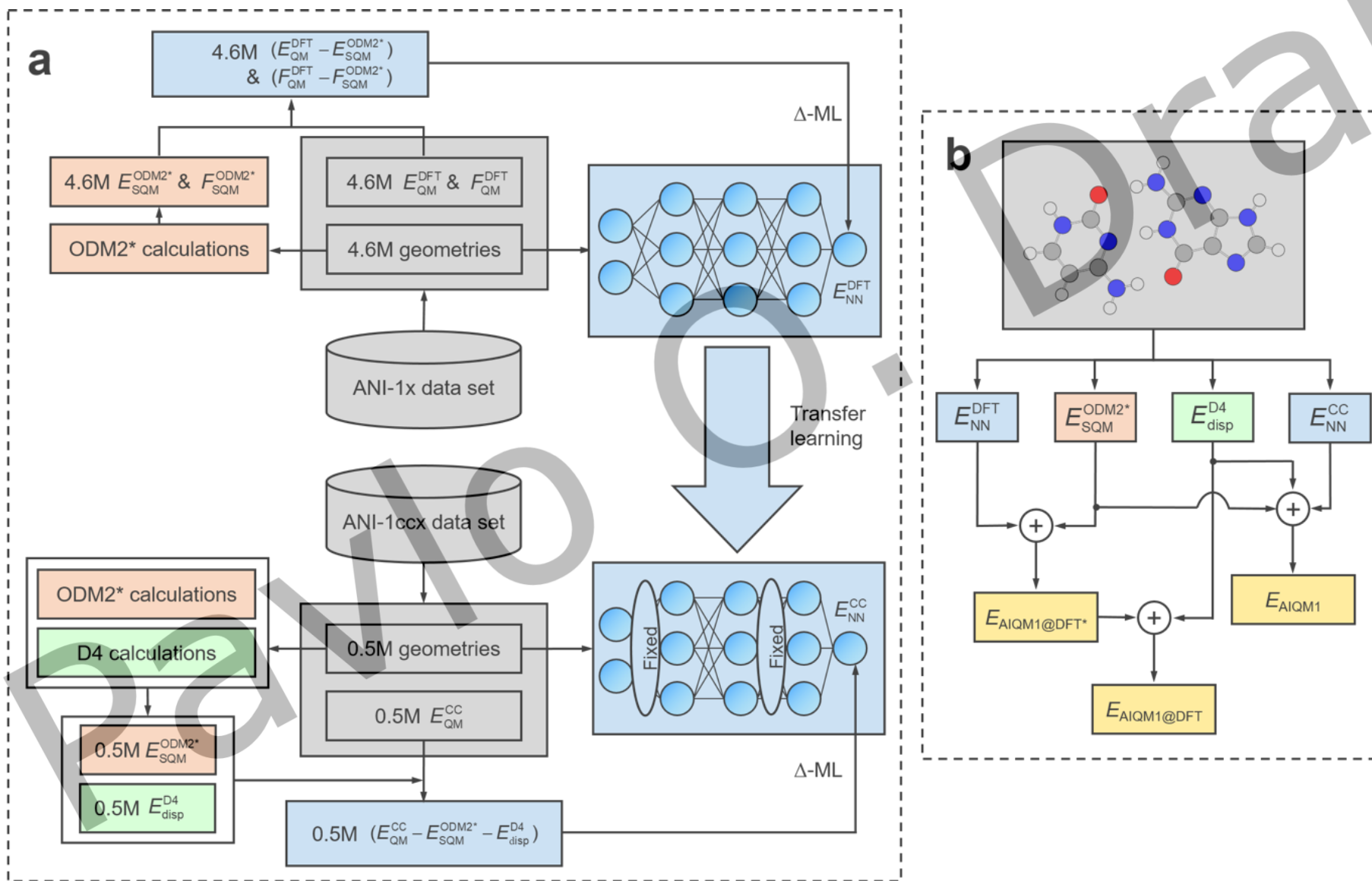
*AI methods: easy to use, also online*

*AI methods are faster, more accurate, and trustworthy*





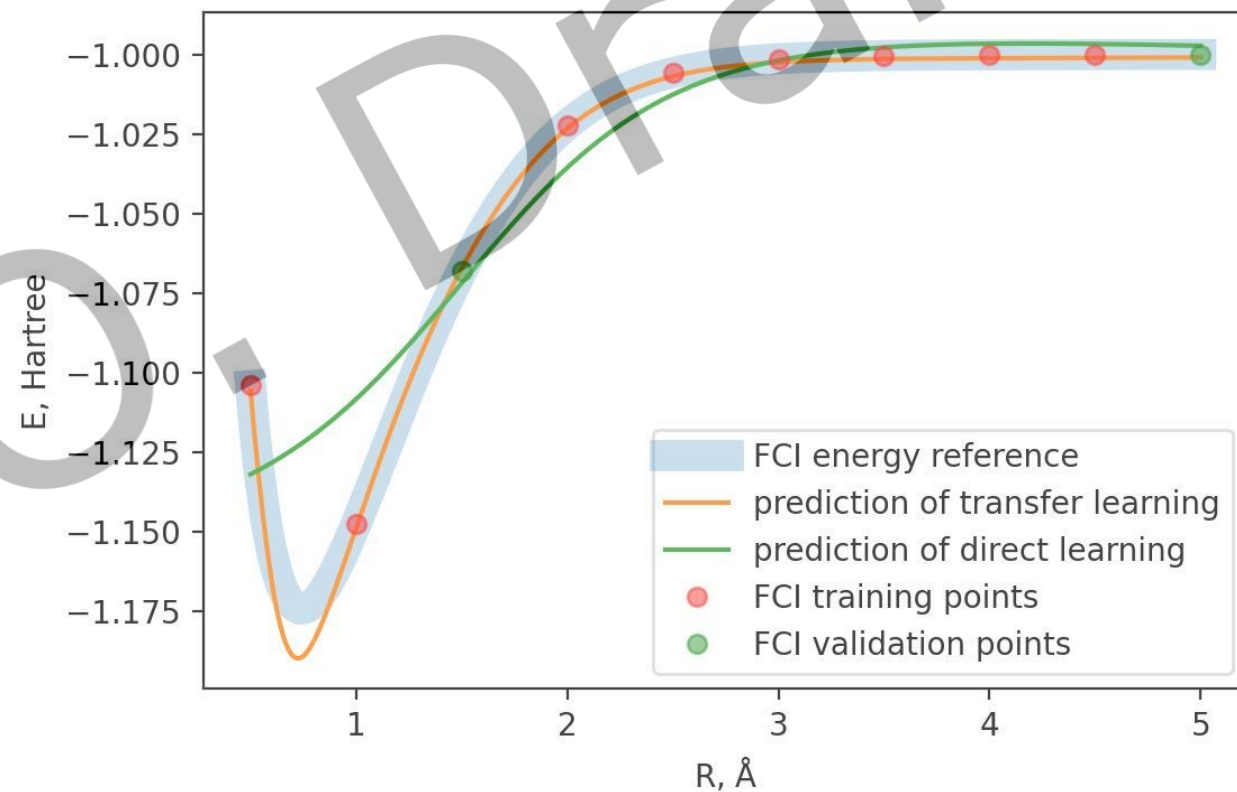
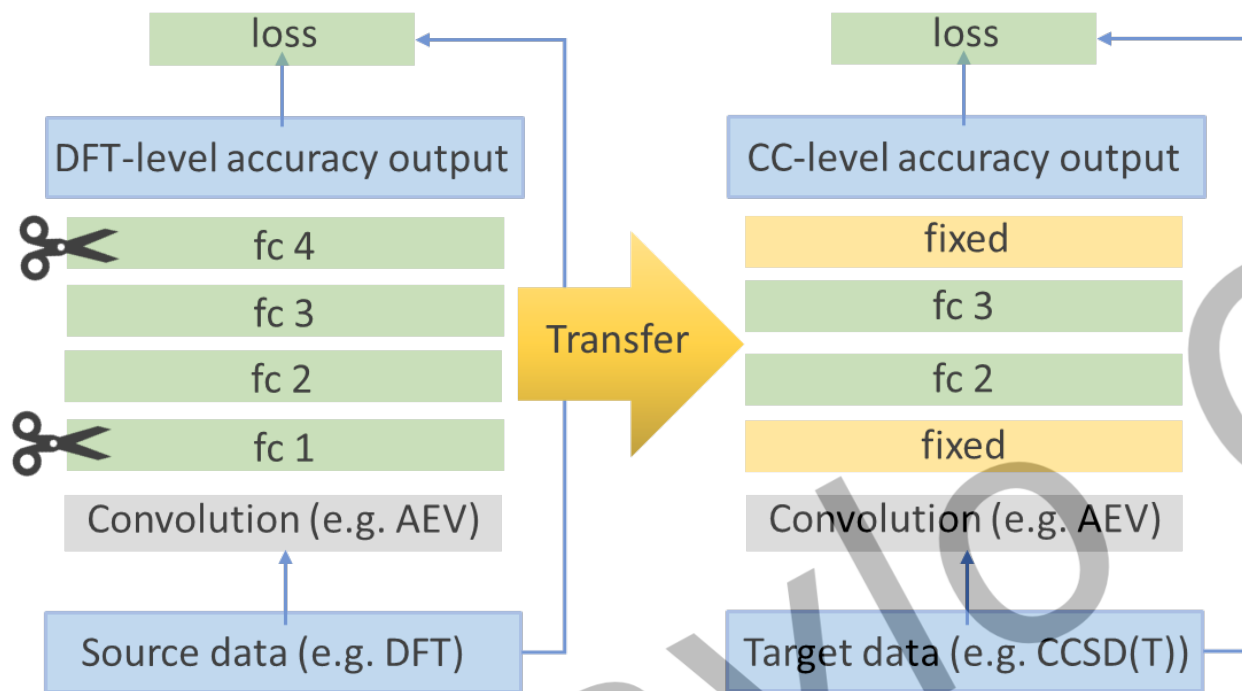
\*CHNO elements only – extensions on the way





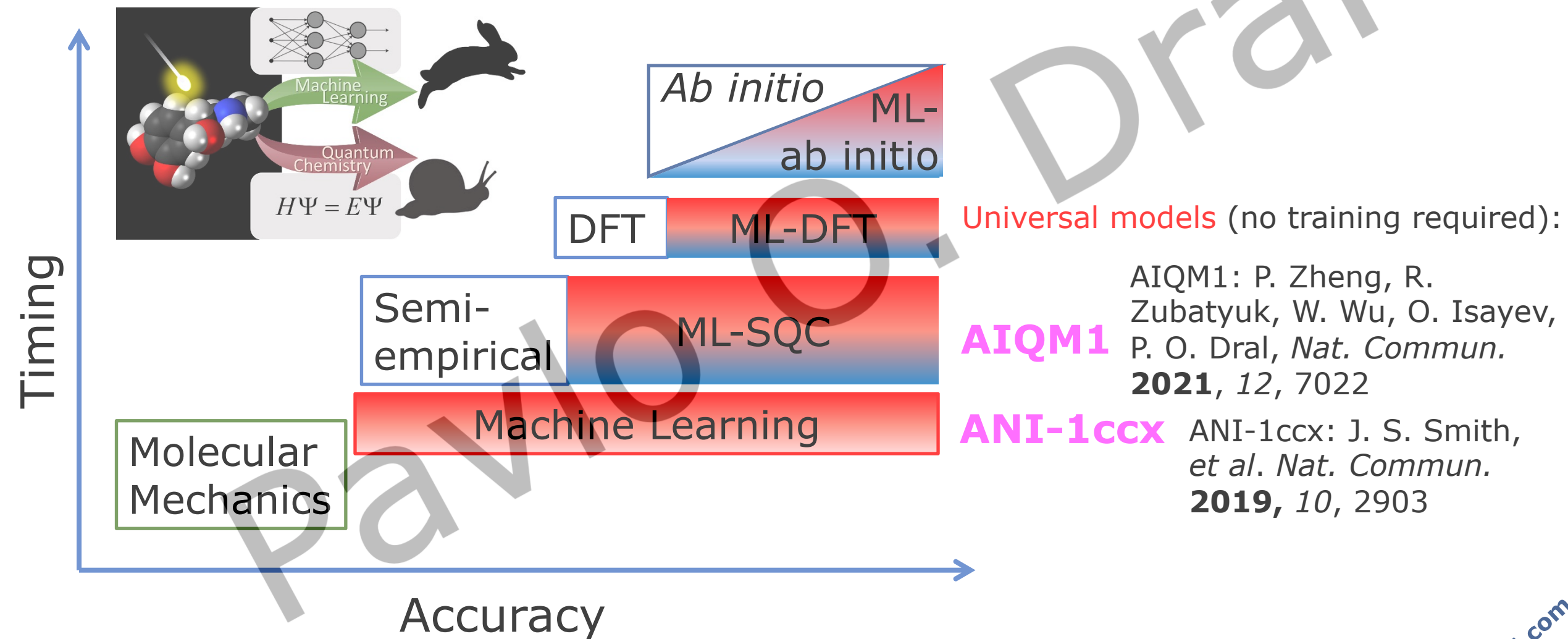
# *Dealing with multiple levels*

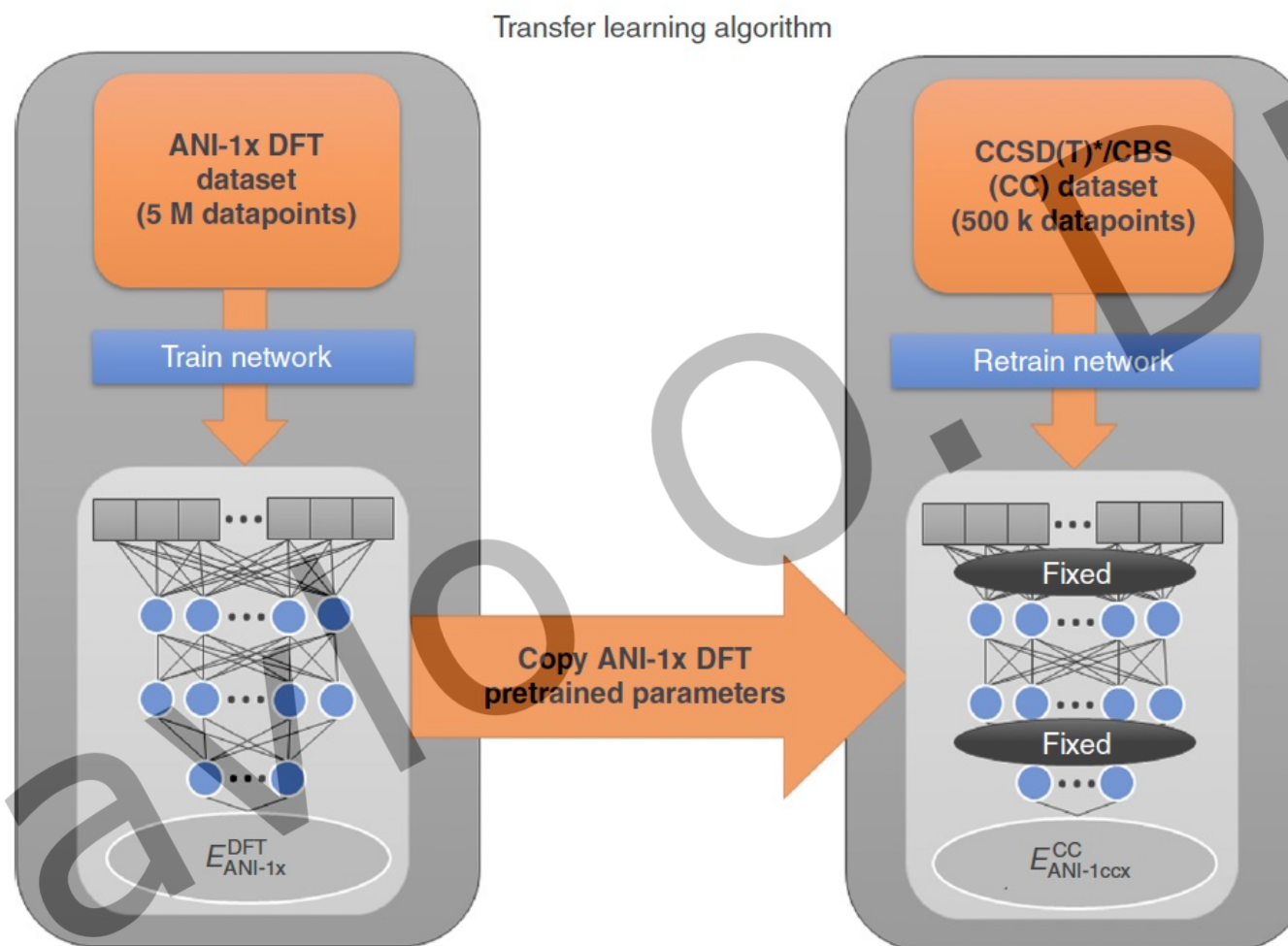
Pavlo O. Dral



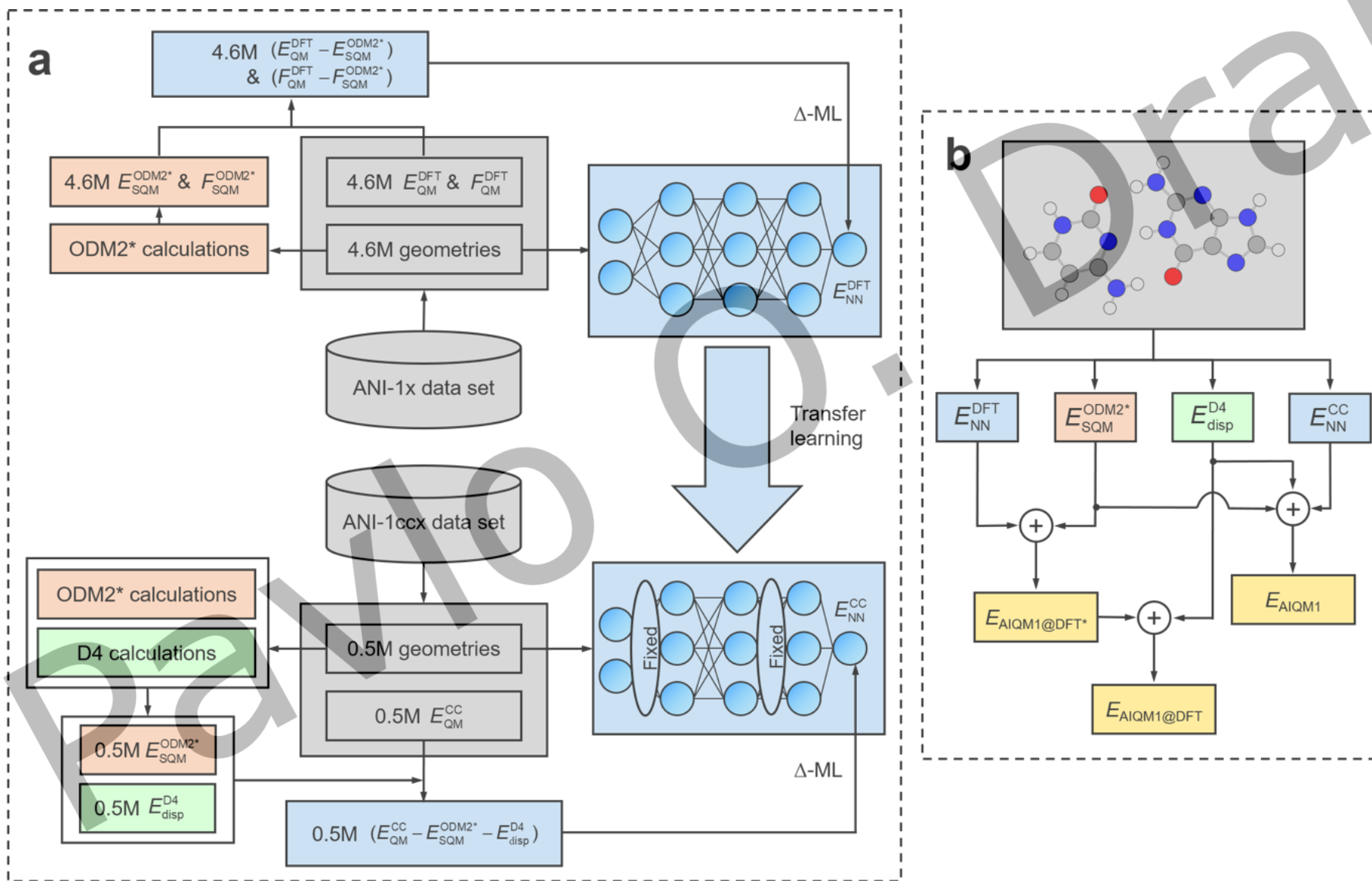
Figures: Pavlo O. Dral, Tetiana Zubatiuk, Bao-Xin Xue, Learning from multiple quantum chemical methods:  $\Delta$ -learning, transfer learning, co-kriging, and beyond. In *Quantum Chemistry in the Age of Machine Learning*, Pavlo O. Dral, Ed. Elsevier: **2023**. Paperback ISBN: 9780323900492

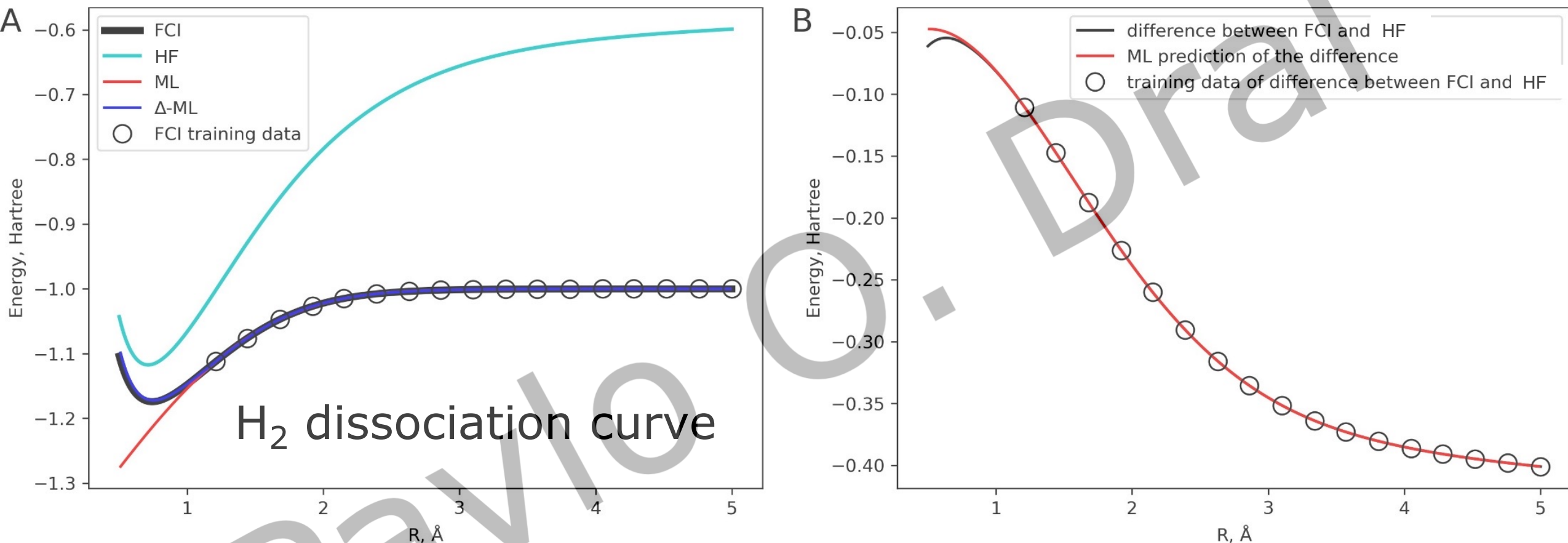
P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388





**Fig. 4** Diagram of the transfer learning technique evaluated in this work. Transfer learning starts from a pretrained ANI-1x DFT model, then retrains to higher accuracy CCSD(T)\* / CBS data with some parameters fixed during training





$\Delta$ -learning: R. Ramakrishnan, P. O. Dral, M. Rupp, O. A. von Lilienfeld,  
*J. Chem. Theory Comput.* **2015**, *11*, 2087

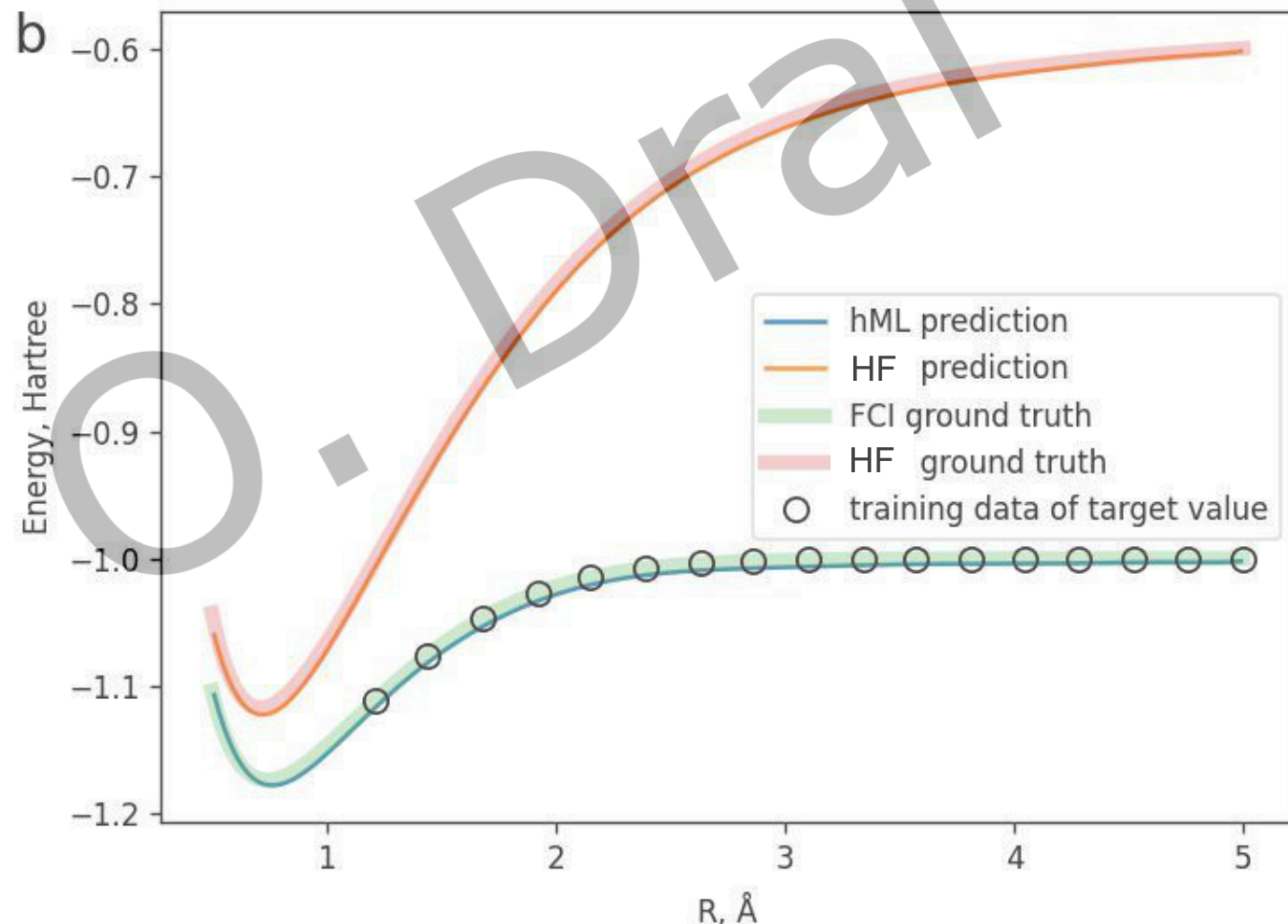
Figure: Pavlo O. Dral, Tetiana Zubatiuk, Bao-Xin Xue, Learning from multiple quantum chemical methods:  $\Delta$ -learning, transfer learning, co-kriging, and beyond. In *Quantum Chemistry in the Age of Machine Learning*, Pavlo O. Dral, Ed. Elsevier: **2023**. Paperback ISBN: 9780323900492

Automatic procedure to find optimal training points for each delta-model

$$\hat{y}_{\Sigma,i} = \sum_M \hat{y}_{M,i}(N_{\text{tr},M})$$

hML for  $\text{H}_2$

$$\text{hML} = \Delta_0^{\text{HF}} + \Delta_{\text{HF}}^{\text{FCI}}$$



Hierarchical ML: P. O. Dral, A. Owens, A. Dral, G. Csányi, *J. Chem. Phys.* **2020**, *152*, 204110

Figure: Pavlo O. Dral, Tetiana Zubatiuk, Bao-Xin Xue, Learning from multiple quantum chemical methods:  $\Delta$ -learning, transfer learning, co-kriging, and beyond. In *Quantum Chemistry in the Age of Machine Learning*, Pavlo O. Dral, Ed. Elsevier: **2023**. Paperback ISBN: 9780323900492

**Automatic** procedure  
to find optimal  
training points for each  
 $\Delta$ -model

$$\hat{y}_{\Sigma,i} = \sum_M \hat{y}_{M,i}(N_{\text{tr},M})$$

$$\frac{e_{\Sigma}}{e_{\Sigma,\text{th}}} + \frac{t_{\Sigma}/t_{\Sigma,\text{max}}}{1 - s_{\text{th}}} + \frac{p}{e_{\Sigma,\text{th}}}$$

$$\frac{\hat{y}_{M,i}(N_{\text{tr},M})}{\hat{y}_{1,i}(N_{\text{tr},M})} \approx r_{M/1,i} = \frac{y_{M,i}}{y_{1,i}}$$

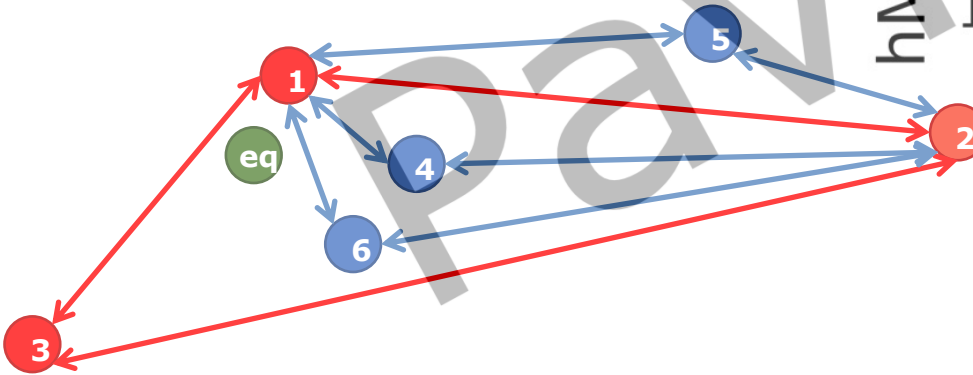
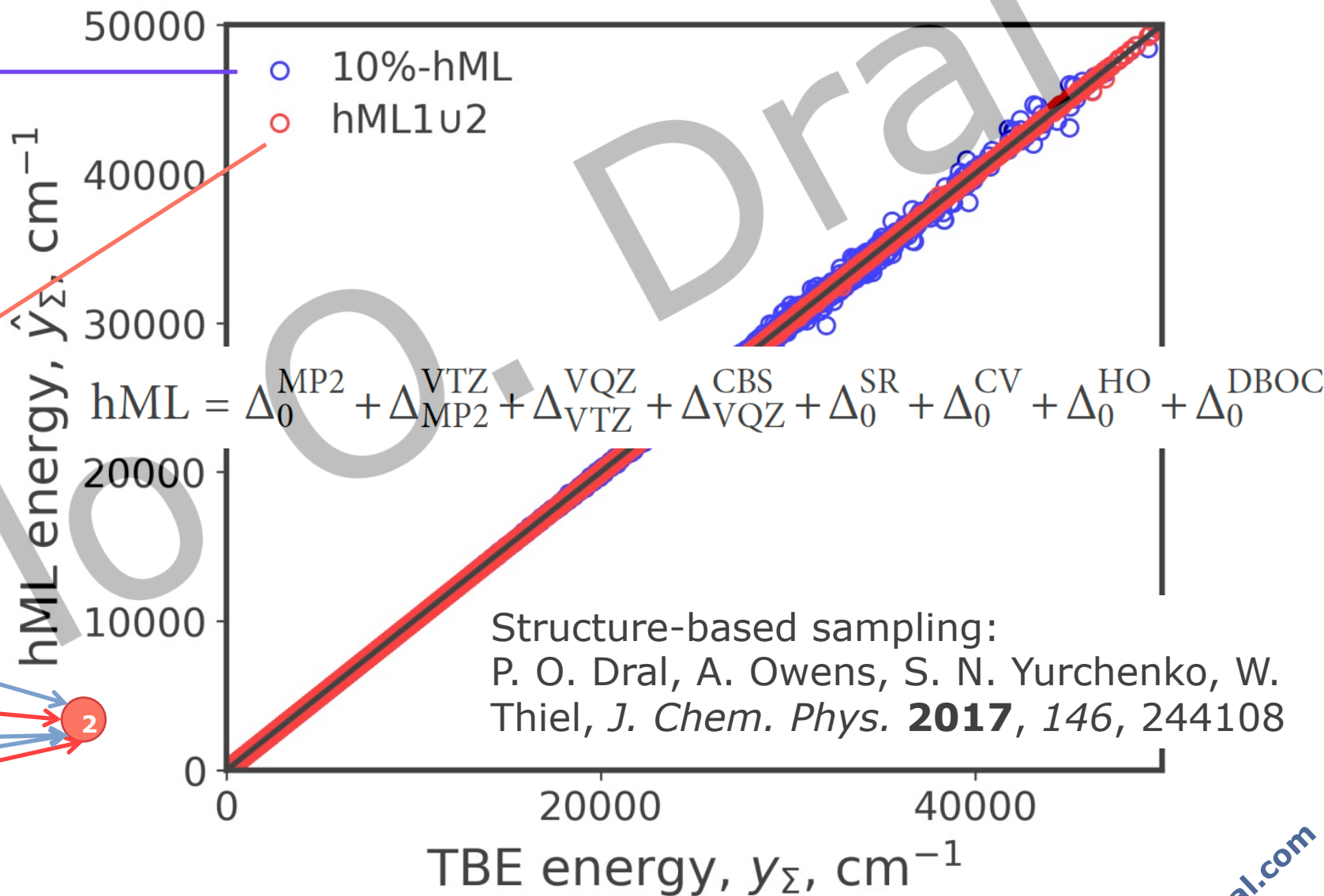
$$\text{hML} = \Delta_0^{\text{MP2}} + \Delta_{\text{MP2}}^{\text{VTZ}} + \Delta_{\text{VTZ}}^{\text{VQZ}} + \Delta_{\text{VQZ}}^{\text{CBS}} + \Delta_0^{\text{SR}} + \Delta_0^{\text{CV}} + \Delta_0^{\text{HO}} + \Delta_0^{\text{DBOC}}$$



Pure ML-model trained on **saved 90% of CPU-time**

Weighted RMSE:  
 **$3.49 \text{ cm}^{-1} = 0.01 \text{ kcal/mol}$**

$\Delta$ -ML models:  
**saved 99% of CPU-time**  
Weighted RMSE:  
 **$1.12 \text{ cm}^{-1} = 0.003 \text{ kcal/mol}$**



data set	ODM2	B3LYP/ 6-31G*	$\omega$ B97X/ 6-31G*	$\omega$ B97X-D/ 6-31G*	$\omega$ B97X/ def2-TZVPP	$\omega$ B97X-D4/ def2-TZVPP	ANI- 1ccx	AIQM1 @DFT*	AIQM1 @DFT	AIQM1	CCSD(T)* /CBS
energies, kcal/mol											
CHNO	2.64	6.71	4.10	3.84	3.21	2.76	—	2.49	2.12	0.87	—
G3/99	3.04	8.53	3.46	3.22	4.18	3.20	—	2.83	2.06	0.88	—
ISOMERS44 ( $\Delta H_f$ )	1.16	8.08	3.57	3.53	4.52	3.78	—	3.00	2.27	0.42	—
ISOMERS44 ( $\Delta H_r$ )	0.70	2.29	1.45	1.31	1.19	1.10	1.68	0.95	0.89	0.50	—
IsoL6/11	1.48	5.26	3.83	3.36	1.75	1.64	1.46	1.65	1.55	0.62	0.47
HC7/11	5.37	6.44	16.90	13.98	6.83	7.10	2.53	8.89	9.16	1.43	1.57

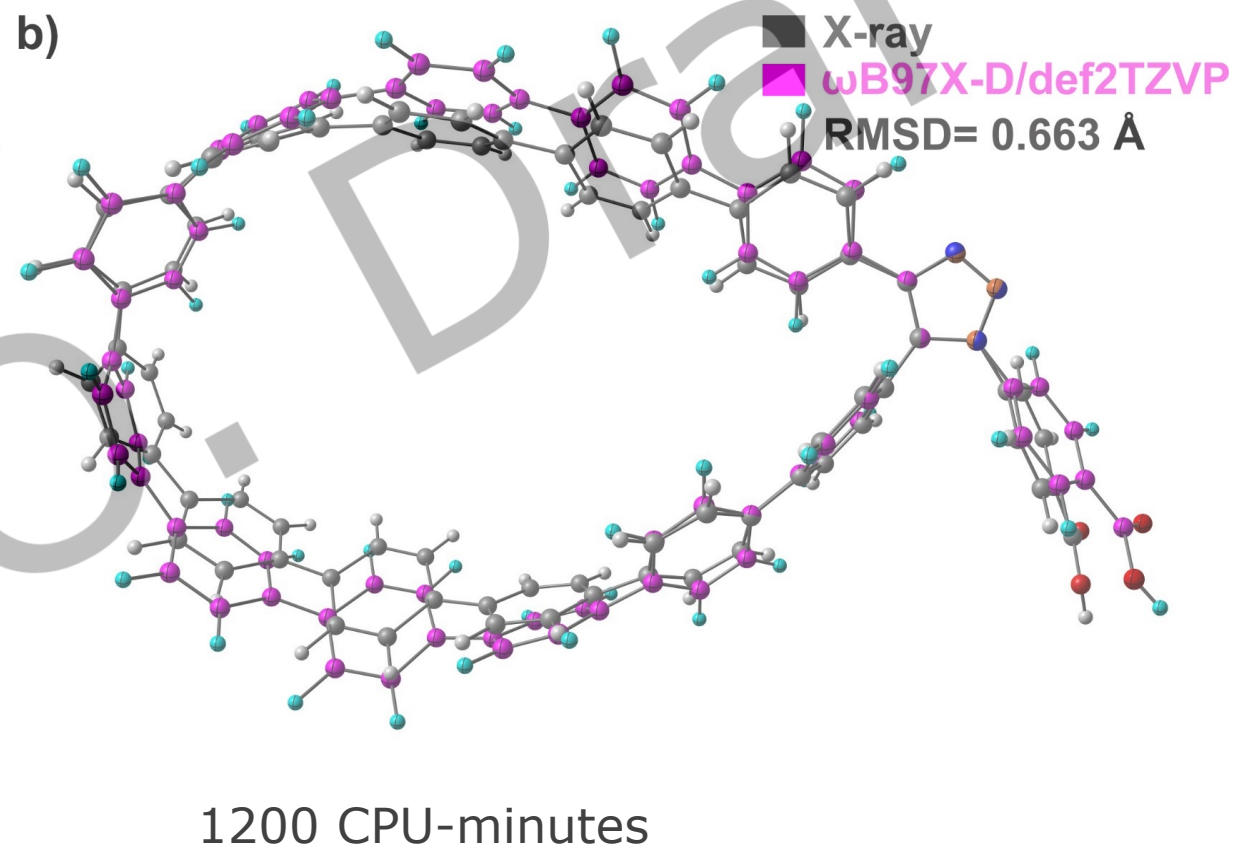
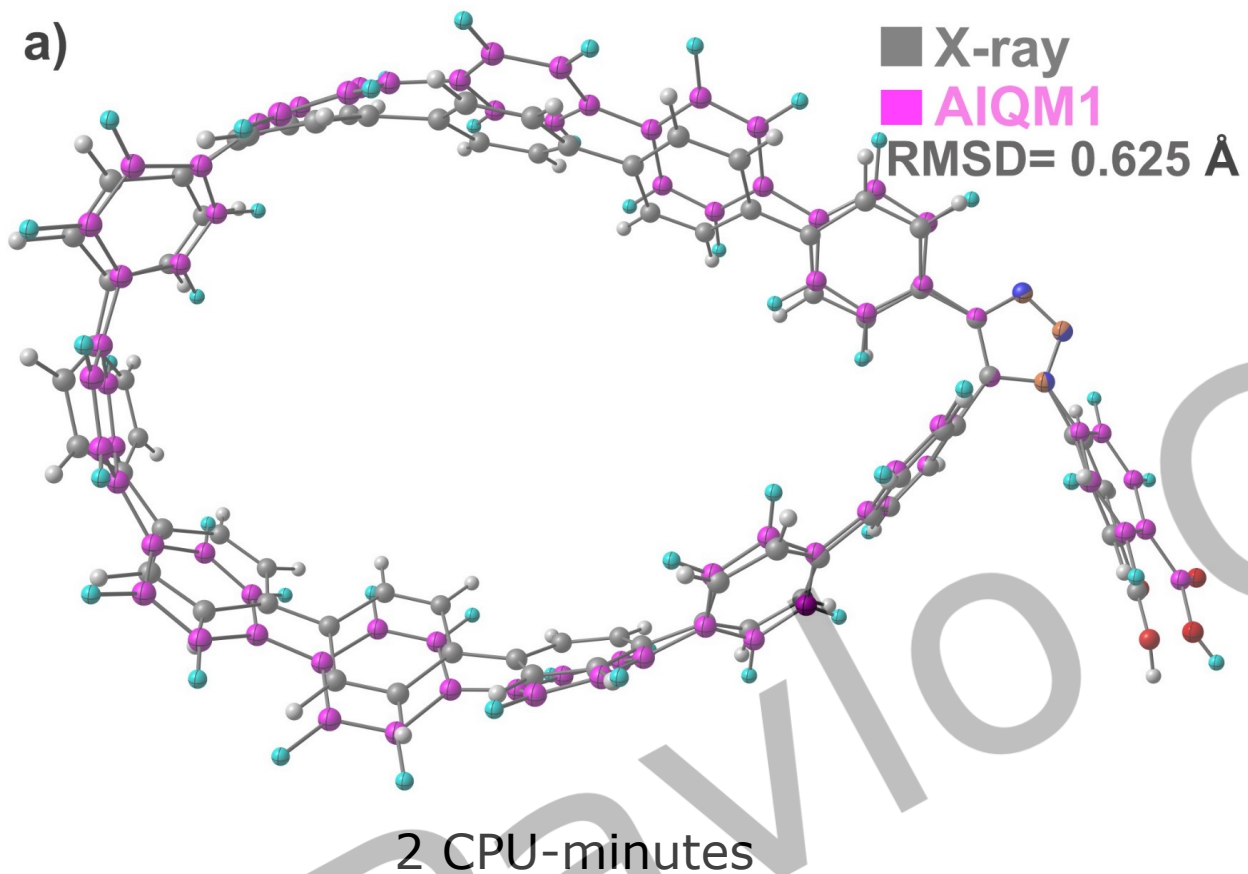
Ground-state properties of neutral, closed-shell compounds  
(heats of formation, reaction enthalpies, and ZPVE-exclusive reaction energies)

Torsion	0.74	0.55	0.30	0.29	0.20	0.19	0.23	0.23	0.23	0.19	0.05
---------	------	------	------	------	------	------	------	------	------	------	------

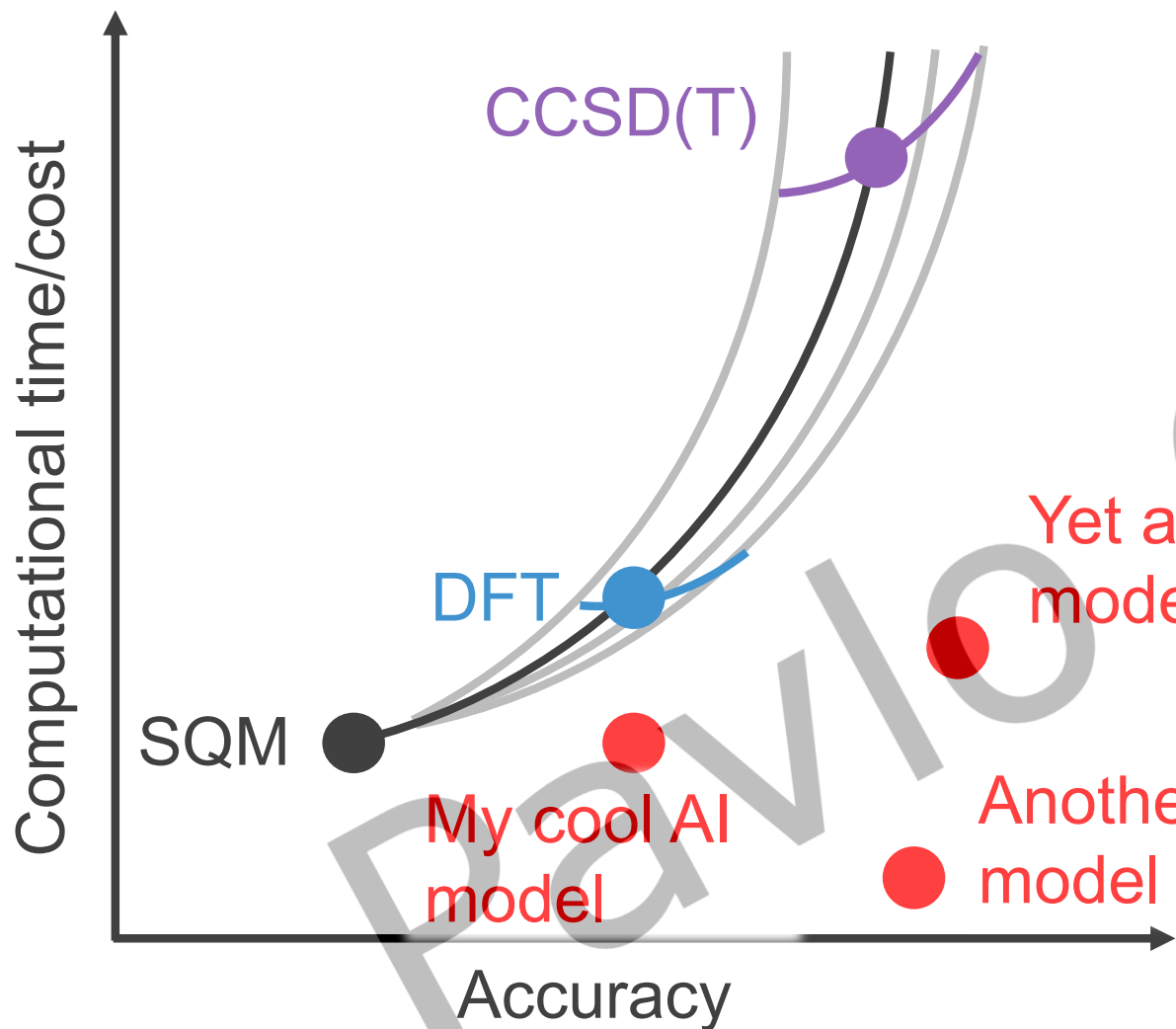
bond lengths, Å											
CHNO	0.015	0.006	0.008	0.007	0.010	0.010	0.011	0.010	0.010	0.007	—
MGHBL9	0.023	0.007	0.006	0.005	0.002	0.002	0.047	0.011	0.011	0.004	—
MGNHBL11	0.026	0.006	0.003	0.002	0.008	0.008	0.004	0.008	0.008	0.002	—

bond angles, °											
CHNO	2.04	0.70	0.68	0.64	0.68	0.68	1.00	0.77	0.77	0.70	—

dihedral angles, °											
CHNO	4.07	5.20	4.68	6.10	7.12	7.11	5.86	2.14	2.14	2.31	—



T. A. Schaub, A. Zieleniewska, R. Kaur, M. Minameyer, W. Yang, C. M. Schüßlbauer, L. Zhang, M. Freiberger, L. N. Zakharov, T. Drewello, P. O. Dral, D. Guldi, R. Jasti. Tunable Macrocyclic Polyparaphenylene Nanolassos via Copper-Free Click Chemistry. *Chem. Eur. J.* **2023**, 29, e202300668



Try to create more  
better models...

# Can we do better?

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

**Lots of models!  
How to choose?**

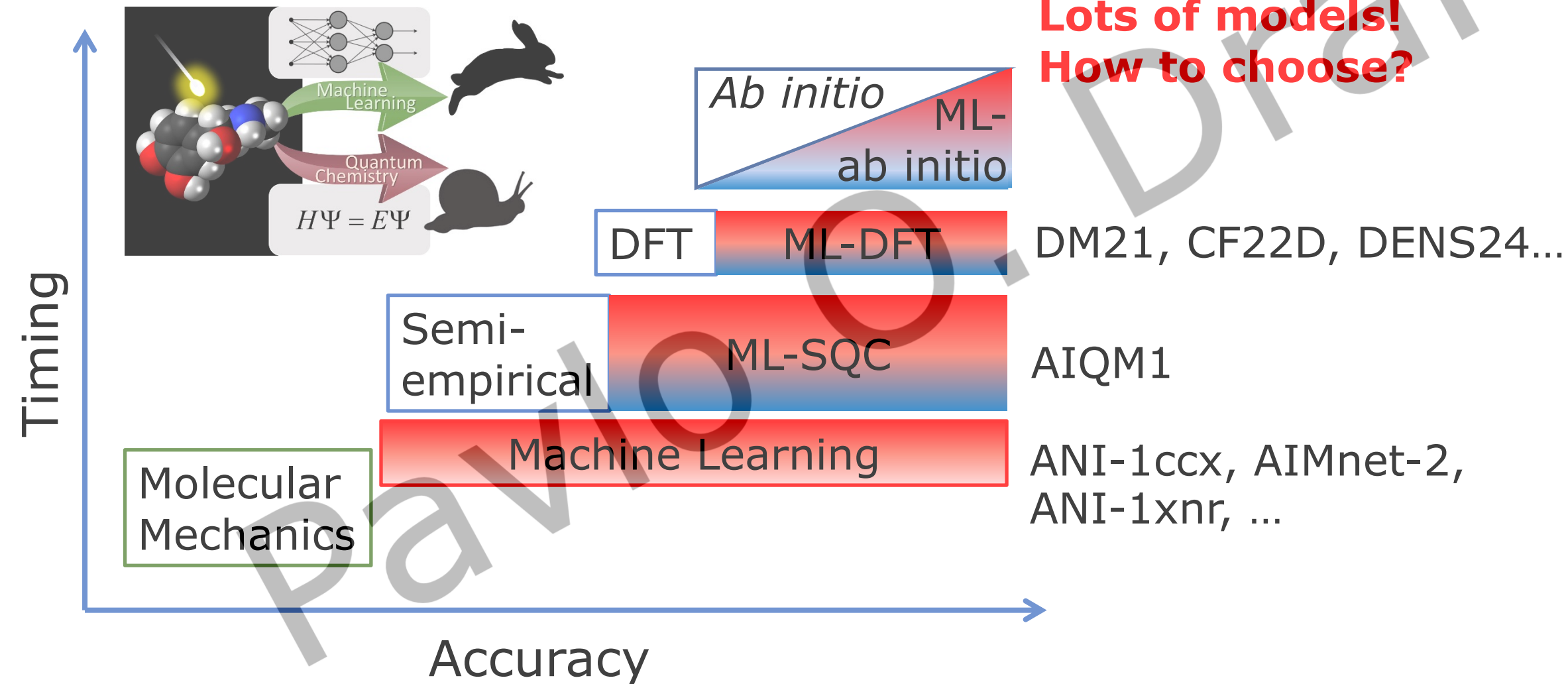


Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

# Can we do better?

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

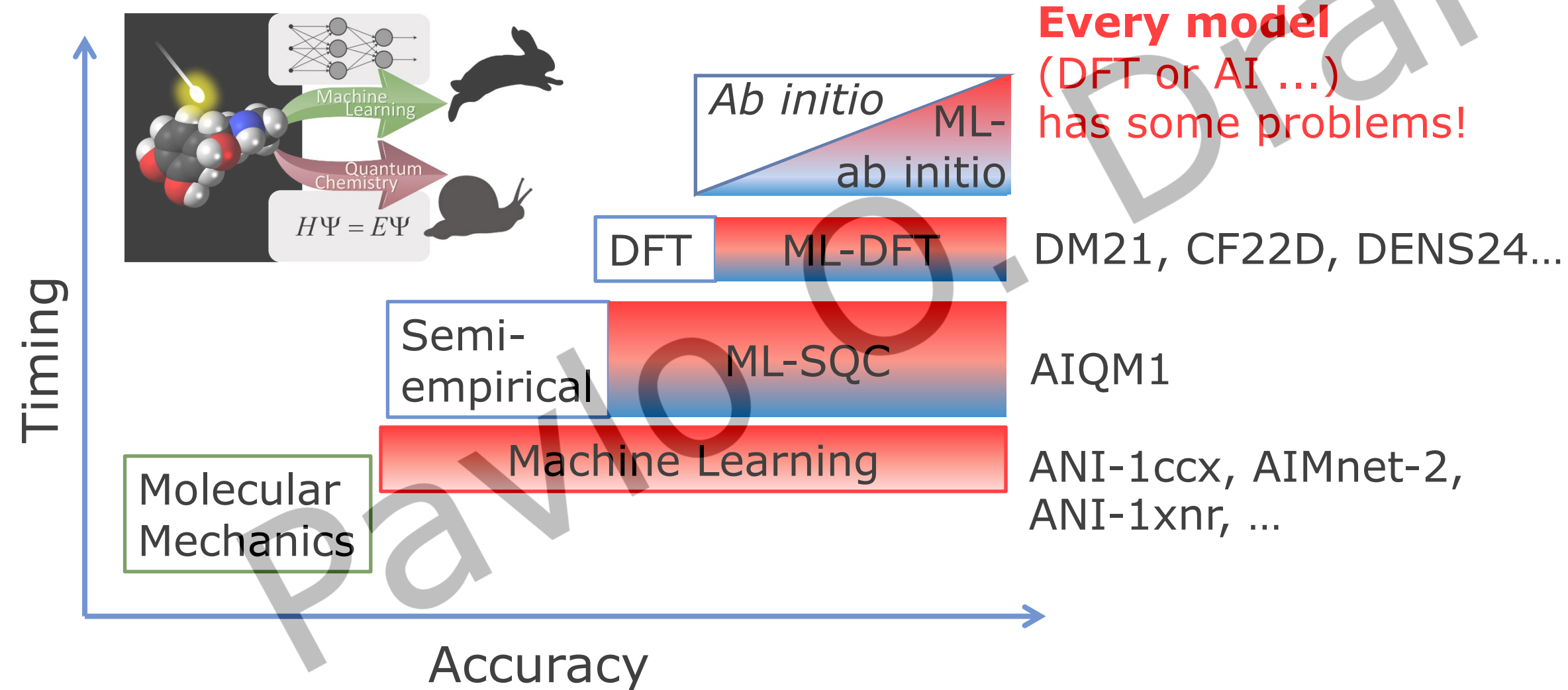
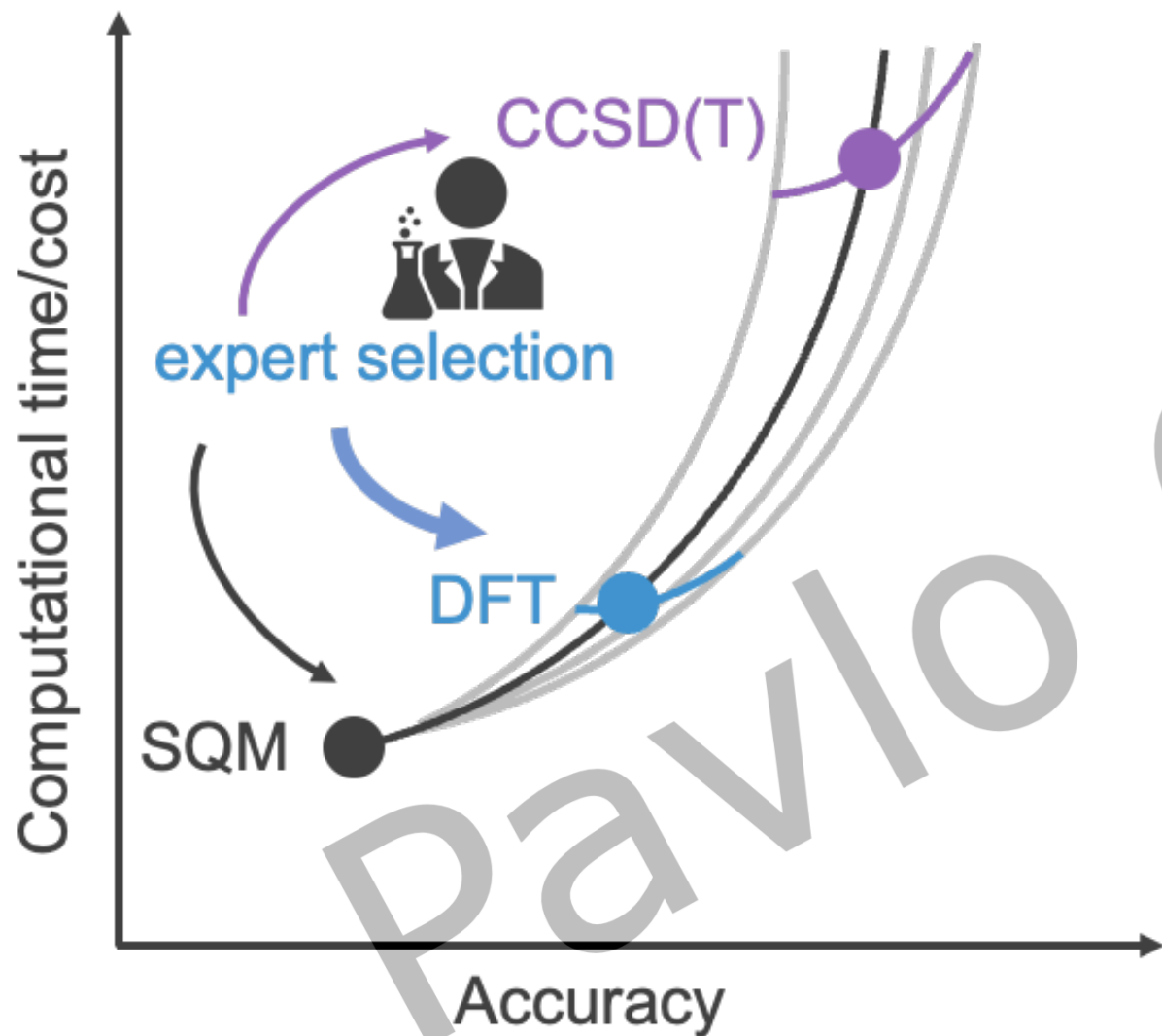
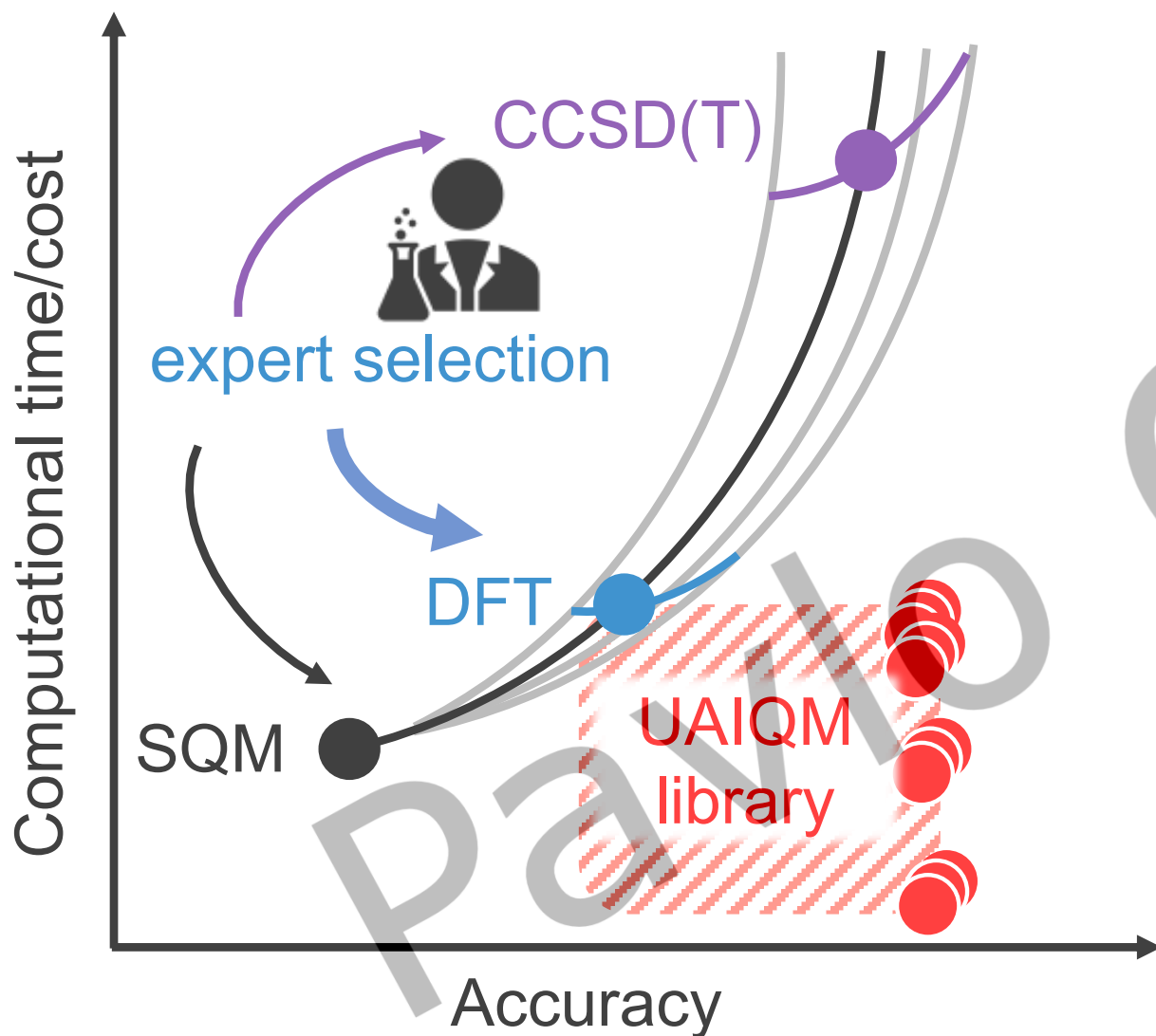


Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336



We can use **human expertise to select models**



We put together a dozen of state-of-the-art models in a library and just give new models **version numbers!**



nature communications

Explore content ▾ About the journal ▾ Publish with us ▾

nature > nature communications > articles > article

Article | [Open access](#) | Published: 02 December 2021

## Artificial intelligence-enhanced quantum chemical method with broad applicability

Peikun Zheng, Roman Zubatyuk, Wei Wu, Olexandr Isayev  & Pavlo O. Dral 

*Nature Commu*

15k Accesses | **Journal of XXX**

AIQM2 : the 2<sup>nd</sup> generation of AI models  
By Dral's group

**Journal of XXX**

AIQM3 : the 3<sup>rd</sup> generation of AI models  
By Dral's group

.....

O. Dral

Reviewers



\* We might end up doing this...

nature communications

Explore content ▾ About the journal ▾ Publish with us ▾

nature > nature communications > articles > article

Article | [Open access](#) | Published: 02 December 2021

## Artificial intelligence-enhanced quantum chemical method with broad applicability

Peikun Zheng, Roman Zubatyuk, Wei Wu, Olexandr Isayev  & Pavlo O. Dral 

Nature Commur

15k Accesses

### Journal of XXX

AIQM2 : the 2<sup>nd</sup> generation of AI models


By Dral's group


### Journal of XXX

AIQM3 : the 3<sup>rd</sup> generation of AI models

By Dral's group

.....







 Pinned

 **Pavlo Dral** @PavloDral · Jul 12

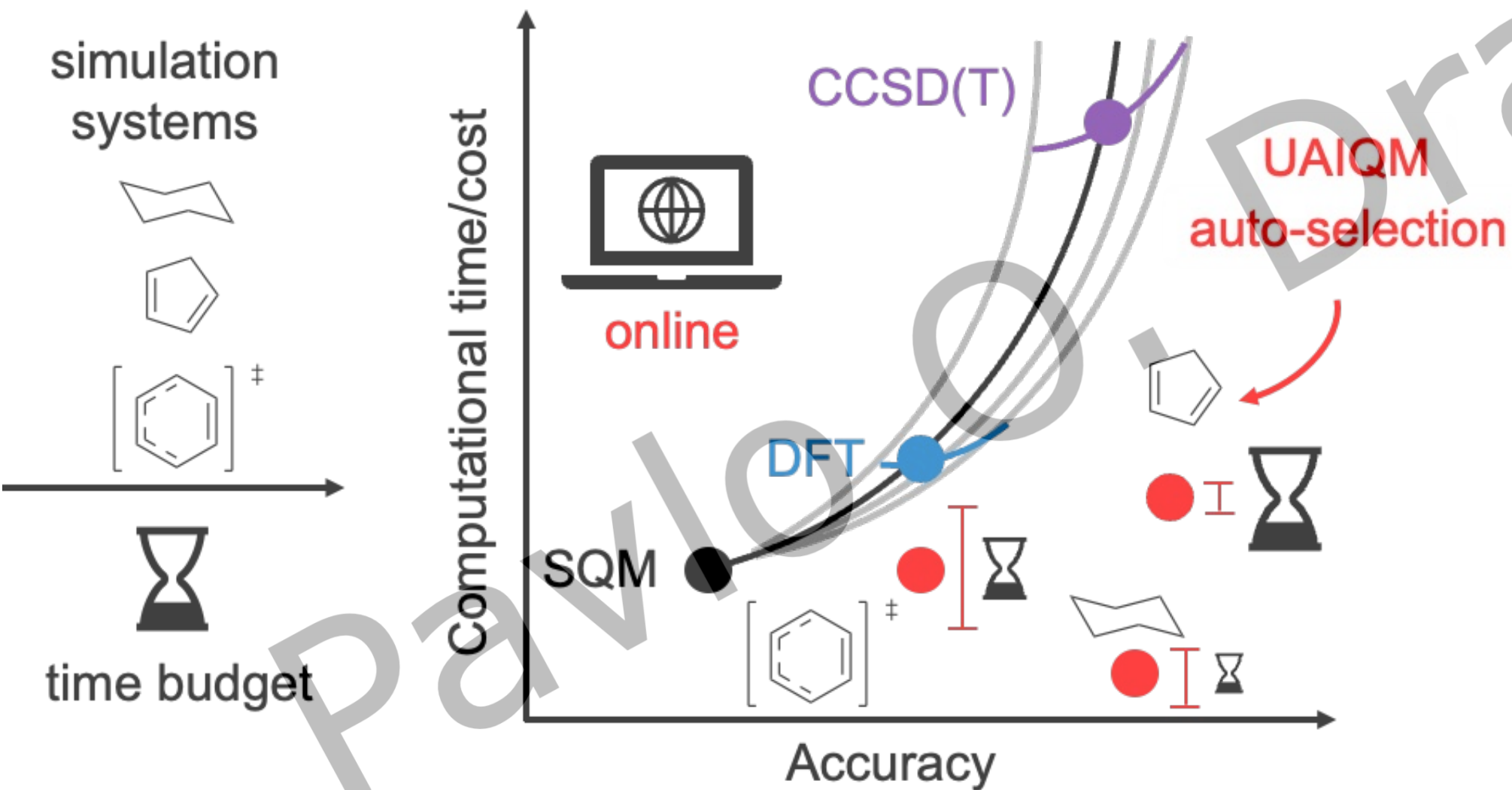
It seems that putting a dozen of state of the art #AI methods in one big paper is not appreciated by some journals. Slice or Not to Slice: That is the Question 😊 [youtube.com/watch?v=wwYDPq...](https://youtube.com/watch?v=wwYDPq...)  
Preprint: [doi.org/10.26434/chemr...](https://doi.org/10.26434/chemr...)

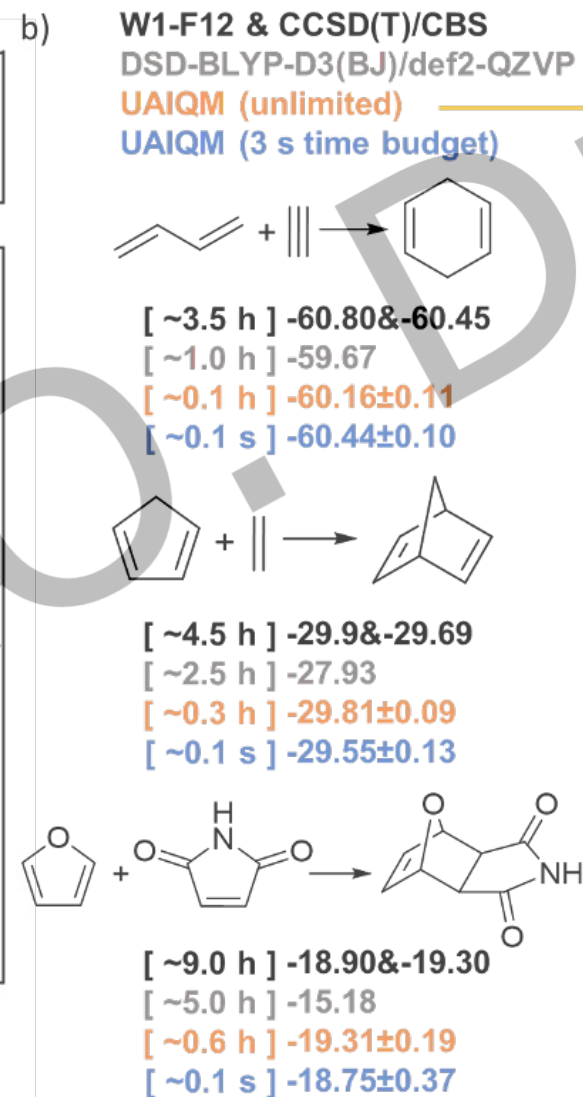
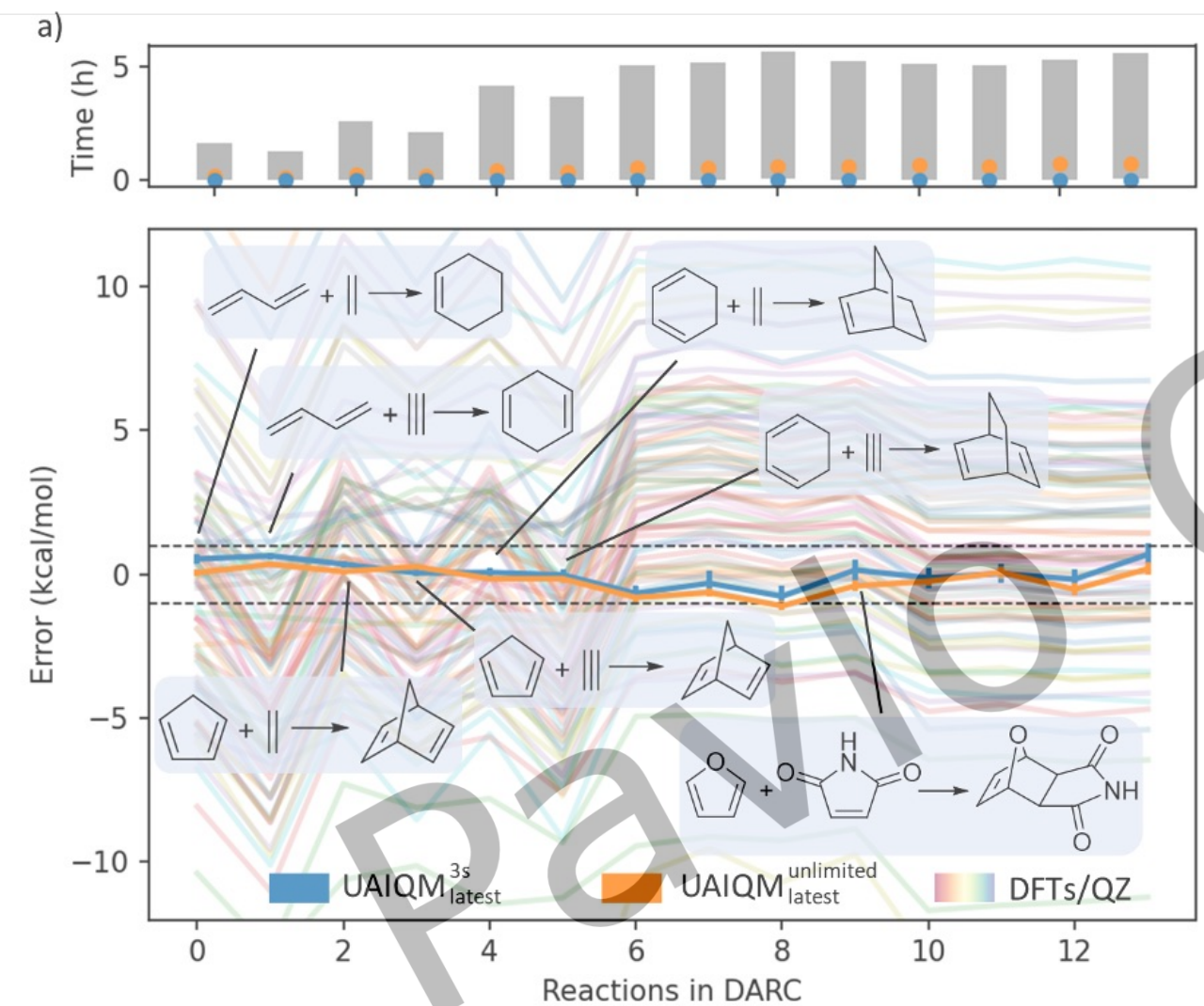
Slice	50%
Not slice	50%

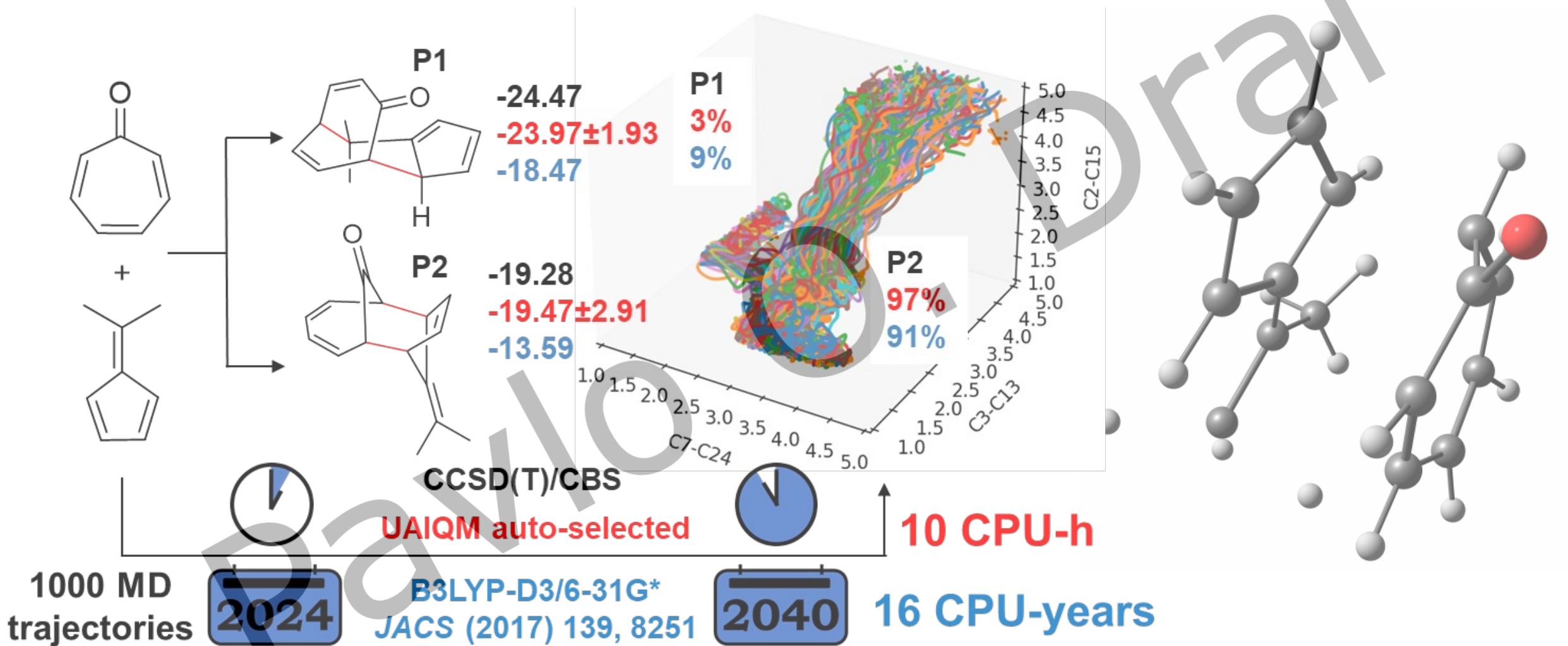
12 votes · Final results

 3   3  1K  

\* We might end up doing this...







=====

WARNING: Uncertainty is too high for selected UAIQM method

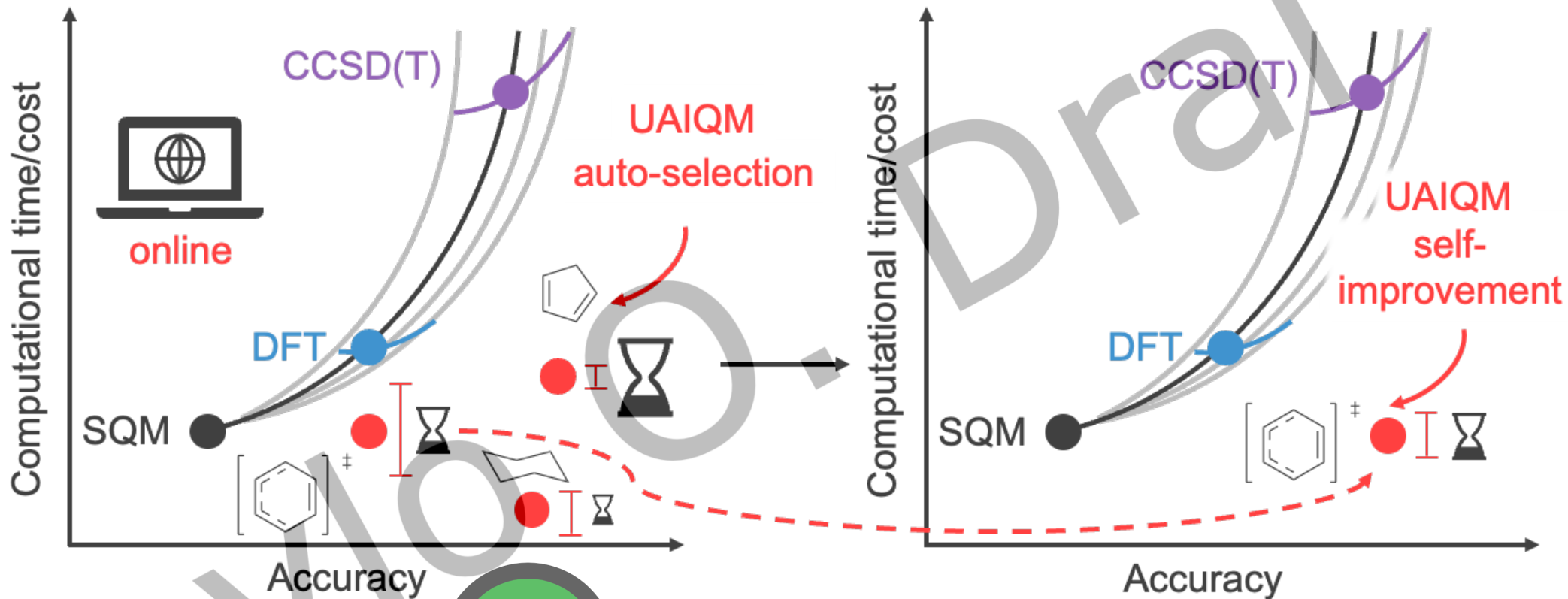
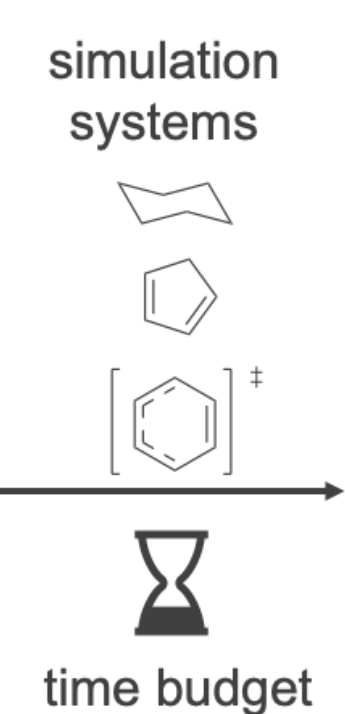
Properties of molecule 1

Selected UAIQM method: uaiqm\_gfn2xtbstar@cc  
Selected version: 20240106

Standard deviation of ML contribution	:	0.00546625 Hartree	3.43012 kcal/mol
Baseline contribution	:	-2.73848388 Hartree	
NN contribution	:	-0.27210359 Hartree	
D4 contribution	:	-0.00007046 Hartree	
Total energy	:	-3.01065793 Hartree	

=====







Updating UAIQM Library...



# Element availability

IA												VIII A					
1 <b>H</b> Hydrogen 1.01	IIA										2 <b>He</b> Helium 4.00						
3 <b>Li</b> Lithium 6.94	4 <b>Be</b> Beryllium 9.01											10 <b>Ne</b> Neon 20.18					
11 <b>Na</b> Sodium 22.99	12 <b>Mg</b> Magnesium 24.31											18 <b>Ar</b> Argon 39.95					
		III B	IV B	VB	VIB	VII B	VIII B	VIII B	IB	IIB	III A	IV A	V A	VIA	VII A		
19 <b>K</b> Potassium 39.10	20 <b>Ca</b> Calcium 40.08	21 <b>Sc</b> Scandium 44.96	22 <b>Ti</b> Titanium 47.87	23 <b>V</b> Vanadium 50.94	24 <b>Cr</b> Chromium 52.00	25 <b>Mn</b> Manganese 54.94	26 <b>Fe</b> Iron 55.85	27 <b>Co</b> Cobalt 58.93	28 <b>Ni</b> Nickel 58.69	29 <b>Cu</b> Copper 63.55	30 <b>Zn</b> Zinc 65.38	31 <b>Ga</b> Gallium 69.72	32 <b>Ge</b> Germanium 72.63	33 <b>As</b> Arsenic 74.92	34 <b>Se</b> Selenium 78.97	35 <b>Br</b> Bromine 79.90	36 <b>Kr</b> Krypton 83.80
37 <b>Rb</b> Rubidium 85.47	38 <b>Sr</b> Strontium 87.62	39 <b>Y</b> Yttrium 88.91	40 <b>Zr</b> Zirconium 91.22	41 <b>Nb</b> Niobium 92.91	42 <b>Mo</b> Molybdenum 95.95	43 <b>Tc</b> Technetium (98)	44 <b>Ru</b> Ruthenium 101.07	45 <b>Rh</b> Rhodium 102.91	46 <b>Pd</b> Palladium 106.42	47 <b>Ag</b> Silver 107.87	48 <b>Cd</b> Cadmium 112.41	49 <b>In</b> Indium 114.82	50 <b>Sn</b> Tin 118.71	51 <b>Sb</b> Antimony 121.76	52 <b>Te</b> Tellurium 127.60	53 <b>I</b> Iodine 126.90	54 <b>Xe</b> Xenon 131.29
55 <b>Cs</b> Cesium 132.91	56 <b>Ba</b> Barium 137.33	57 - 71 Lanthanides	72 <b>Hf</b> Hafnium 178.49	73 <b>Ta</b> Tantalum 180.95	74 <b>W</b> Tungsten 183.84	75 <b>Re</b> Rhenium 186.21	76 <b>Os</b> Osmium 190.23	77 <b>Ir</b> Iridium 192.22	78 <b>Pt</b> Platinum 195.08	79 <b>Au</b> Gold 196.97	80 <b>Hg</b> Mercury 200.59	81 <b>Tl</b> Thallium 204.38	82 <b>Pb</b> Lead 207.20	83 <b>Bi</b> Bismuth 208.98	84 <b>Po</b> Polonium (209)	85 <b>At</b> Astatine (210)	86 <b>Rn</b> Radon (222)
87 <b>Fr</b> Francium (223)	88 <b>Ra</b> Radium (226)	89 - 103 Actinides	104 <b>Rf</b> Rutherfordium (261)	105 <b>Db</b> Dubnium (268)	106 <b>Sg</b> Seaborgium (271)	107 <b>Bh</b> Bohrium (270)	108 <b>Hs</b> Hassium (277)	109 <b>Mt</b> Meitnerium (276)	110 <b>Ds</b> Darmstadtium (281)	111 <b>Rg</b> Roentgenium (280)	112 <b>Cn</b> Copernicium (285)	113 <b>Nh</b> Nihonium (284)	114 <b>Fl</b> Flerovium 289	115 <b>Mc</b> Moscovium (288)	116 <b>Lv</b> Livermorium (293)	117 <b>Ts</b> Tennessine (294)	118 <b>Og</b> Oganesson (294)

 Excellent  
 Supported  
 Unsupported

~CCSD(T) level    max. DFT level

57 <b>La</b> Lanthanum 138.91	58 <b>Ce</b> Cerium 140.12	59 <b>Pr</b> Praseodymium 140.91	60 <b>Nd</b> Neodymium 144.24	61 <b>Pm</b> Promethium (145)	62 <b>Sm</b> Samarium 150.36	63 <b>Eu</b> Europium 151.96	64 <b>Gd</b> Gadolinium 157.25	65 <b>Tb</b> Terbium 158.93	66 <b>Dy</b> Dysprosium 162.50	67 <b>Ho</b> Holmium 164.93	68 <b>Er</b> Erbium 167.26	69 <b>Tm</b> Thulium 168.93	70 <b>Yb</b> Ytterbium 173.05	71 <b>Lu</b> Lutetium 174.97
89 <b>Ac</b> Actinium (227)	90 <b>Th</b> Thorium 232.04	91 <b>Pa</b> Protactinium 231.04	92 <b>U</b> Uranium 238.03	93 <b>Np</b> Neptunium (237)	94 <b>Pu</b> Plutonium (244)	95 <b>Am</b> Americium (243)	96 <b>Cm</b> Curium (247)	97 <b>Bk</b> Berkelium (247)	98 <b>Cf</b> Californium (251)	99 <b>Es</b> Einsteinium (252)	100 <b>Fm</b> Fermium (257)	101 <b>Md</b> Mendelevium (258)	102 <b>No</b> Nobelium (259)	103 <b>Lr</b> Lawrencium (262)

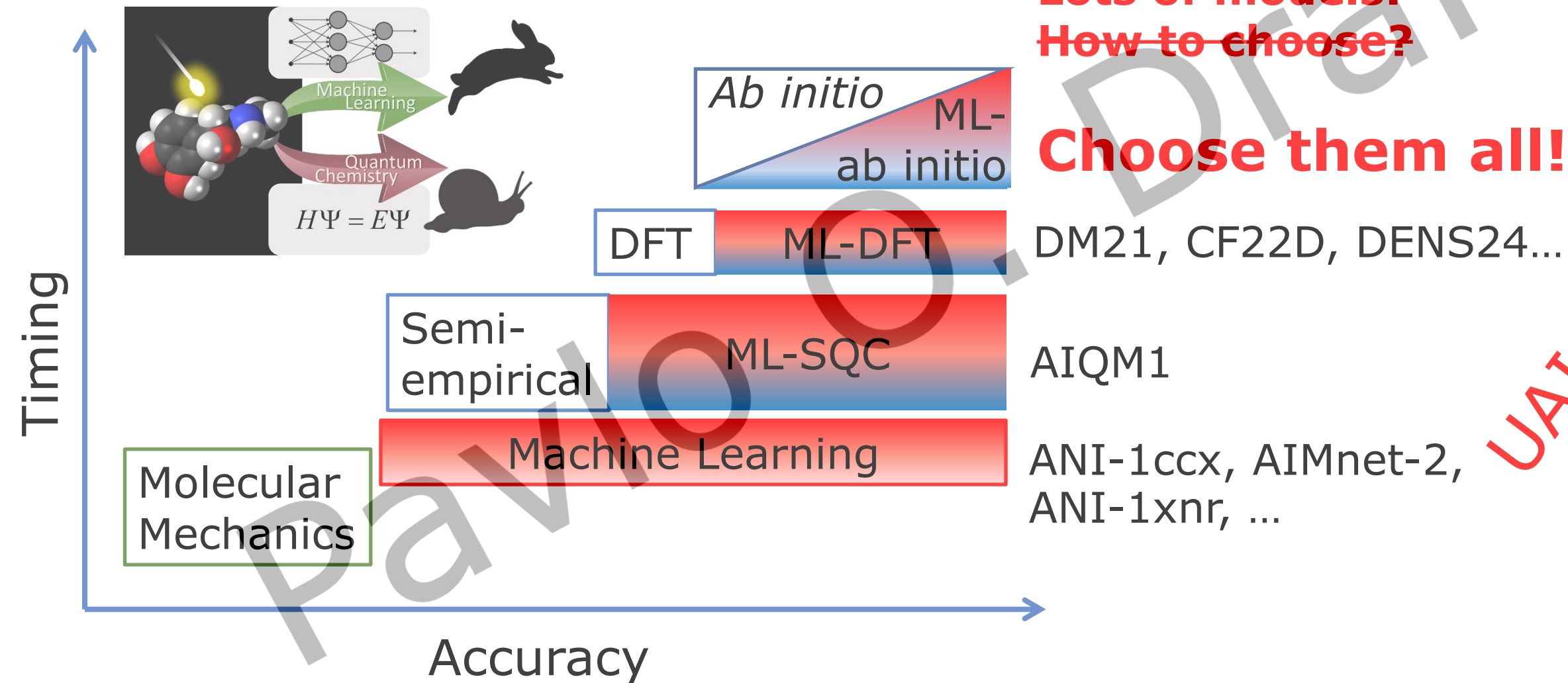


# Can we do better?

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

**Lots of models!**  
**How to choose?**

**Choose them all!**



UAIQM

Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

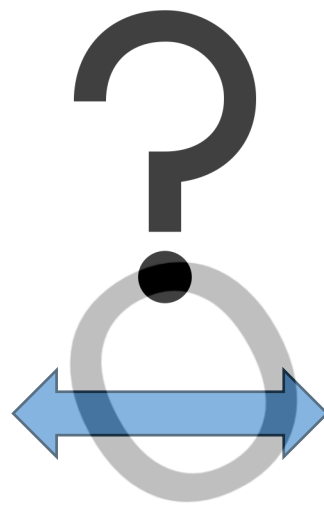
Basic properties and reaction energies of small systems

Reaction barrier heights

Intermolecular noncovalent interactions

Intramolecular noncovalent interactions

Reaction energies for large systems and isomerization reactions



MN12SX

BLYP

B3LYP

SCAN

DSD-BLYP

PBE0

revPBE

$\omega$ B97X-V

.....



Which one is the best?

P. O. Dral, M. Barbatti, *Nat. Rev. Chem.* **2021**, 5, 388

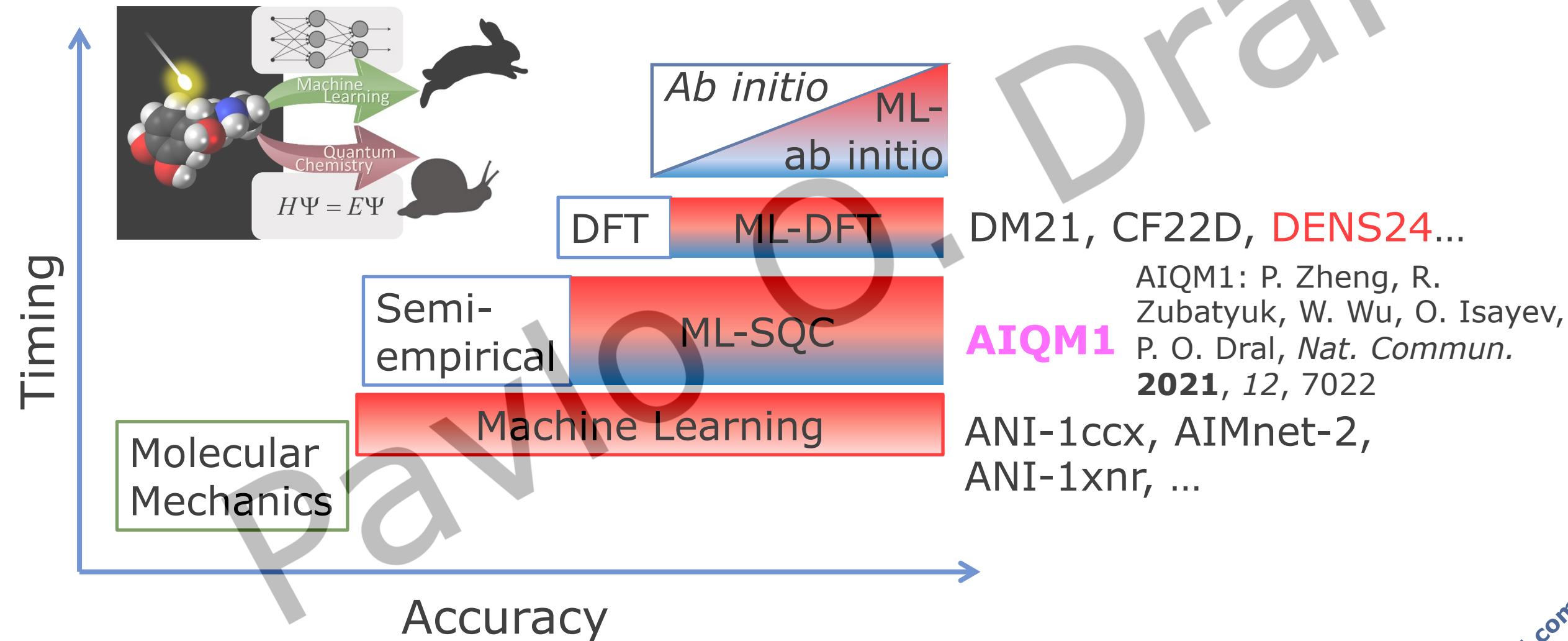


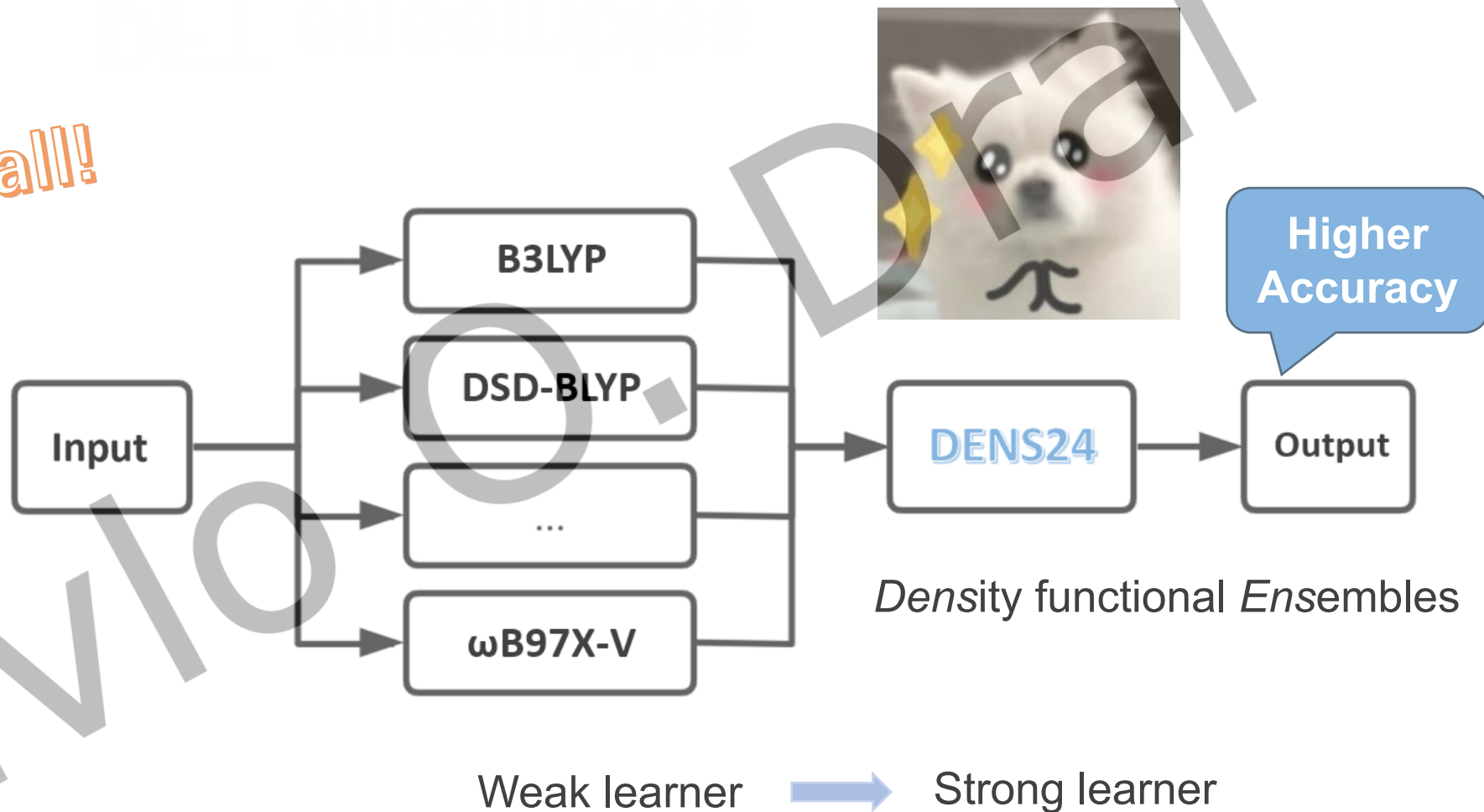
Figure: P. O. Dral, *J. Phys. Chem. Lett.* **2020**, 11, 2336

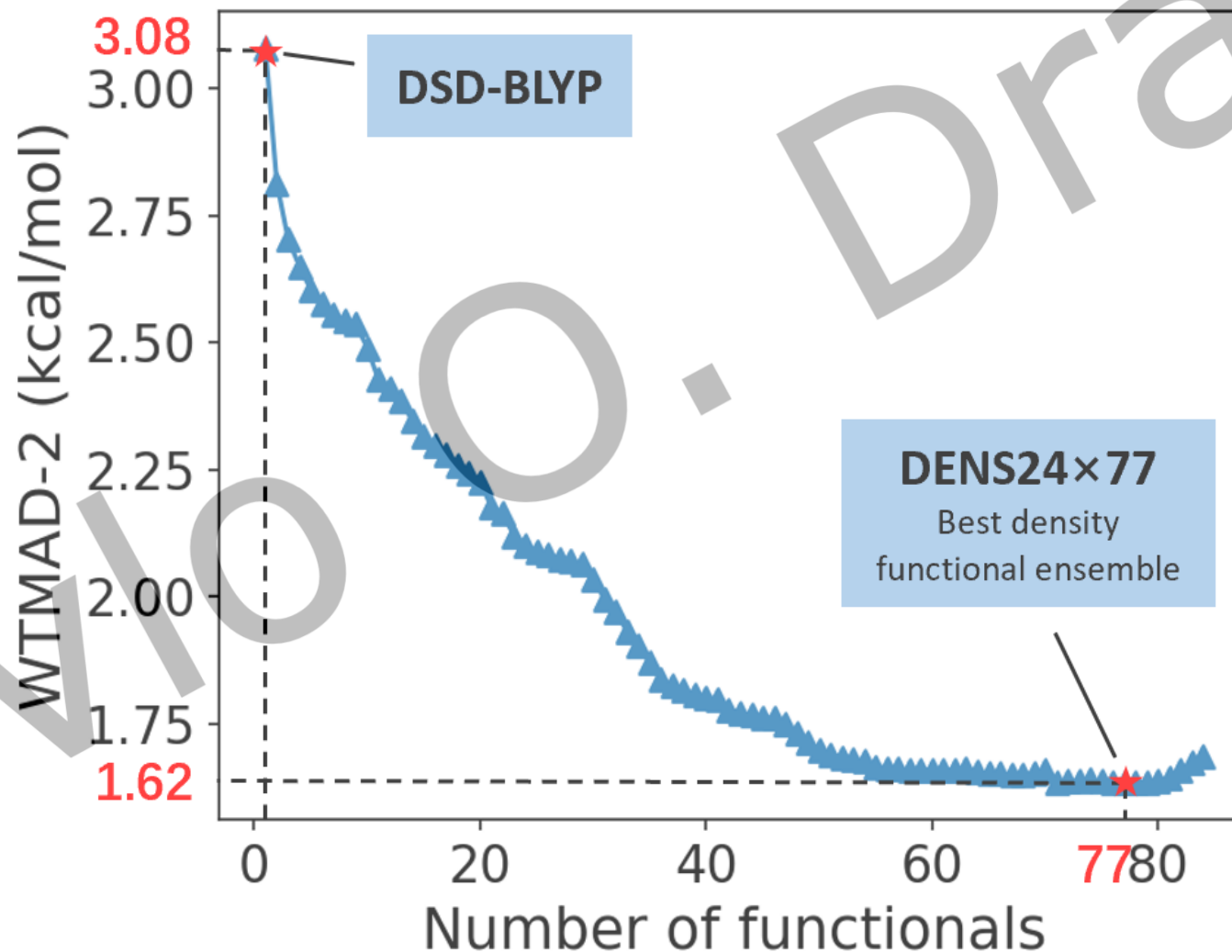
GMTKN55 benchmark:

- Double-hybrid variant: 1.62 kcal/mol
- Hybrid variant: 2.86 kcal/mol

Choose them all!

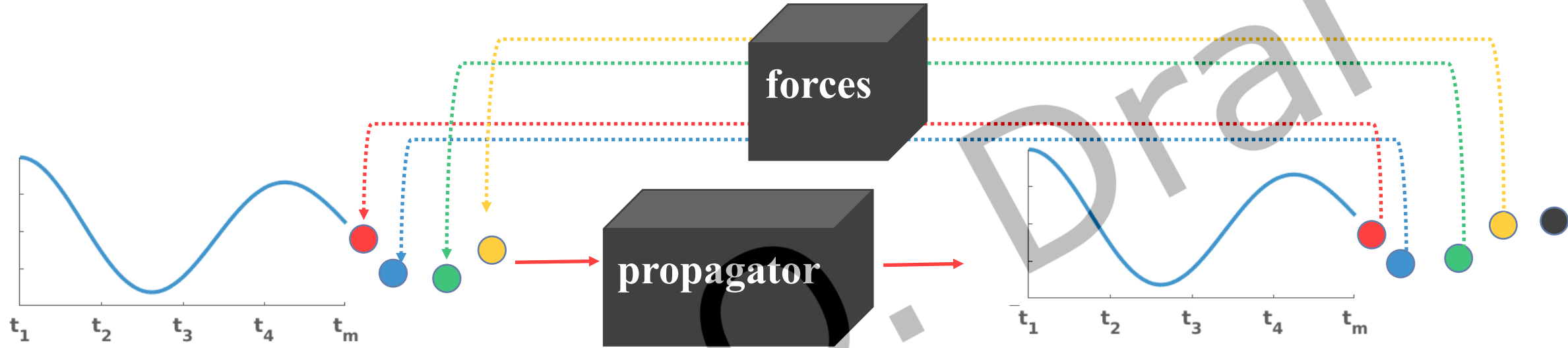
Ensemble





***What else can we do differently in computational chemistry?***

Pavlo O. Dral



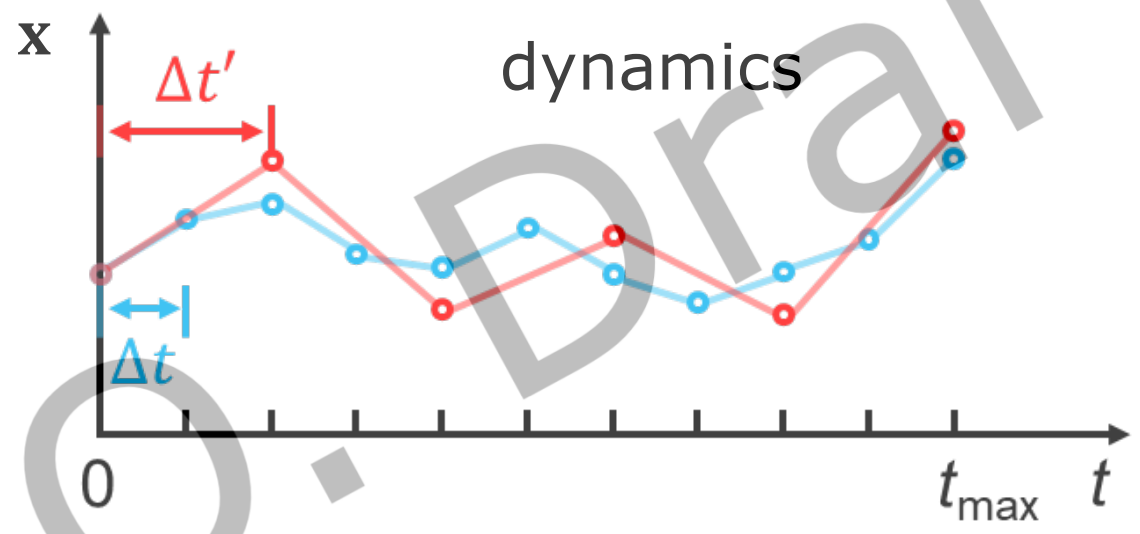
$$F_{A,d} = - \frac{\partial E}{\partial x_{A,d}} = - \frac{\partial E_{\text{machine learning potential}}(\mathbf{x})}{\partial x_{A,d}}$$



# Can we do better?

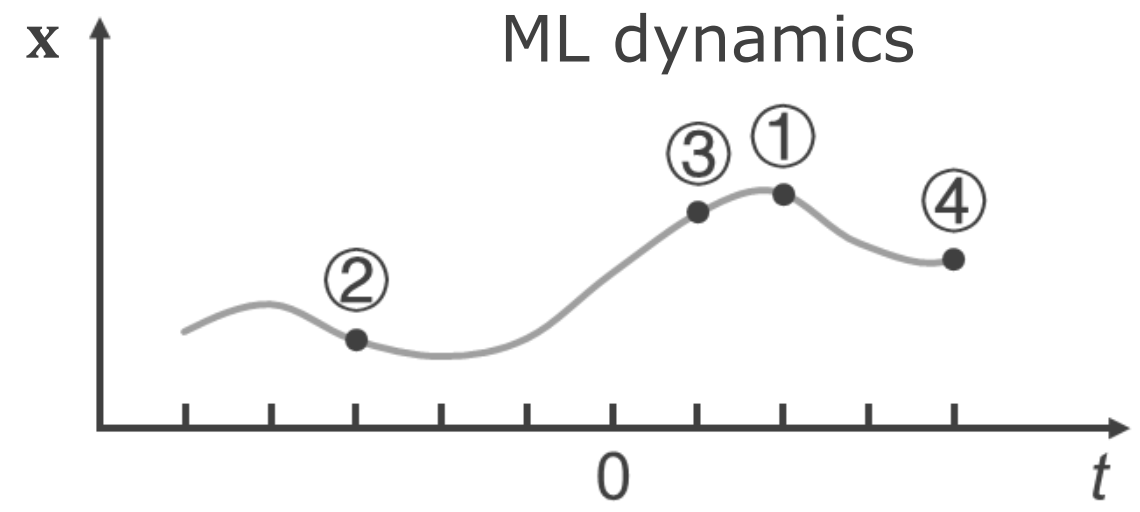
dynamics propagation is

- Iterative (non-parallelizable)
- Depends on time step
- Discrete



Directly learning dynamics

$$\mathbf{x}_t = f(\mathbf{x}_0, \mathbf{v}_0, t)$$



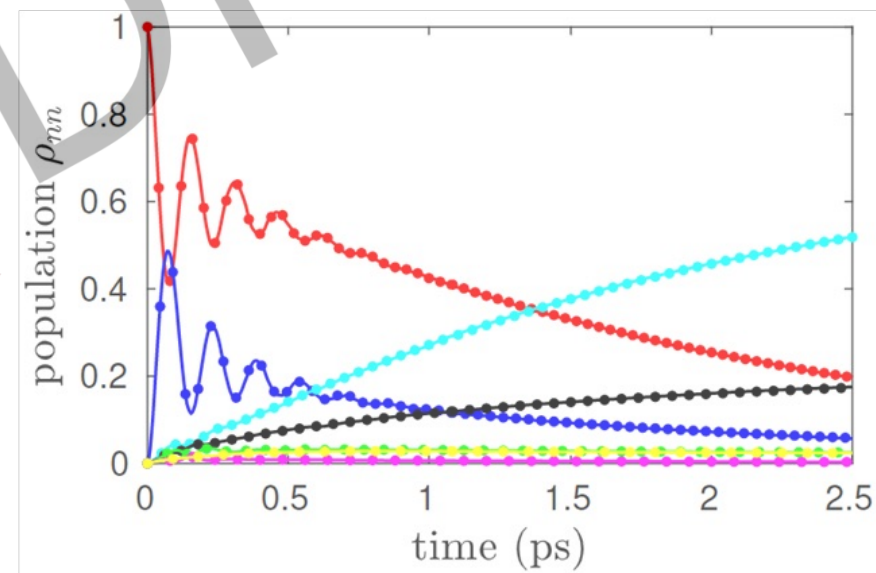
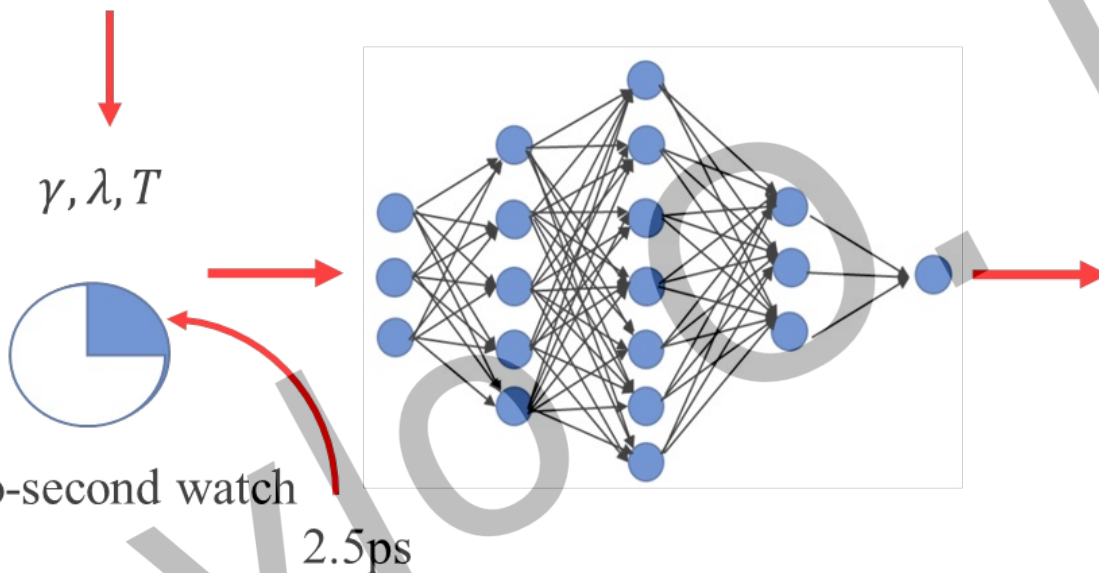
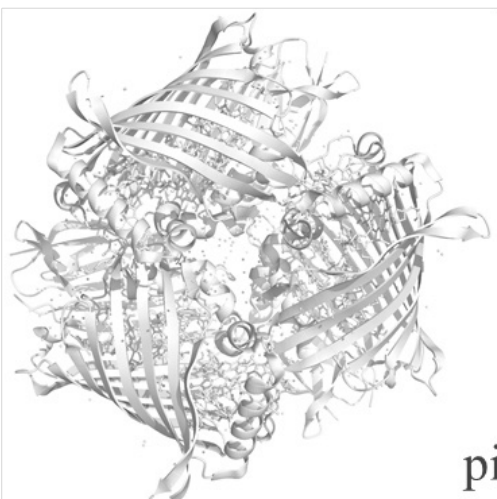
$\gamma$  = characteristic frequency

$\lambda$  = reorganization energy

$T$  = temperature

$$\rho(\text{time}) = f[\text{time}; \text{simulation parameters}]$$

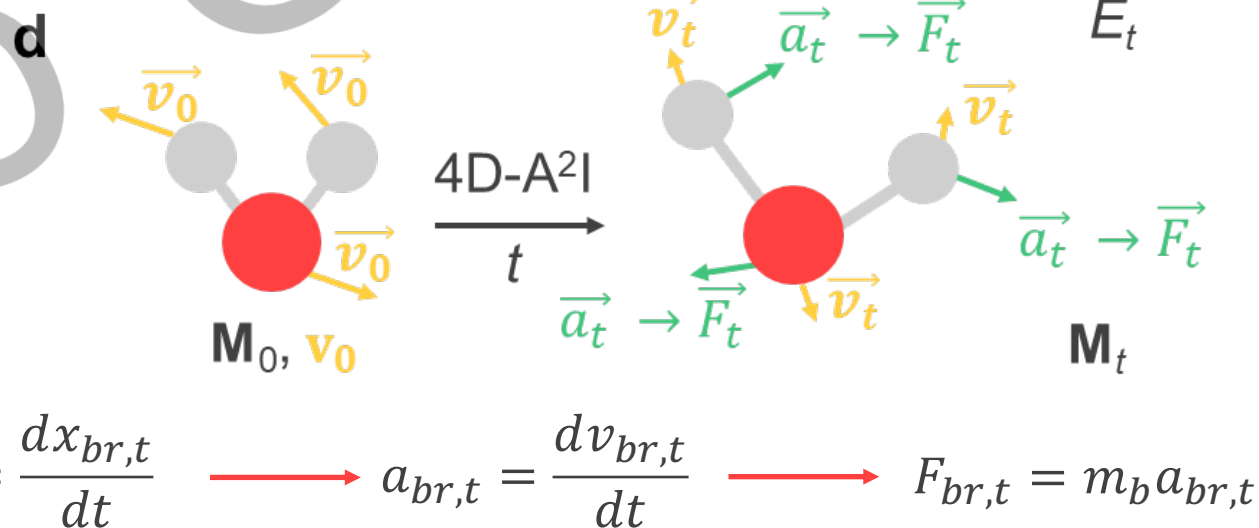
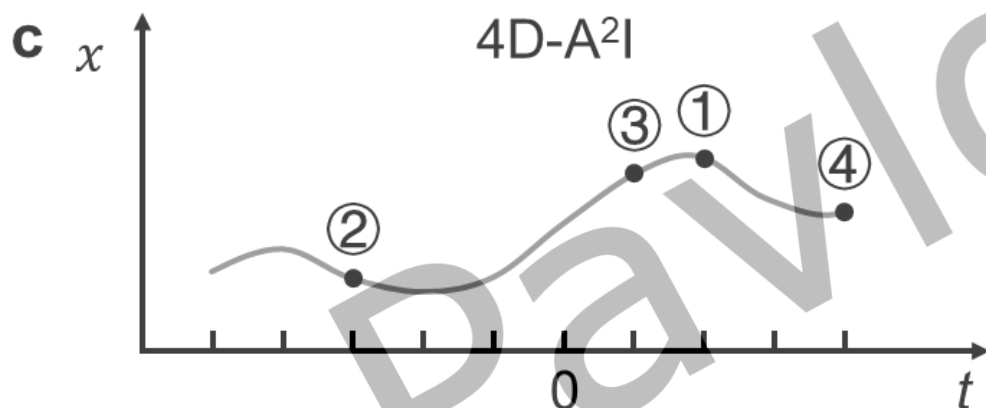
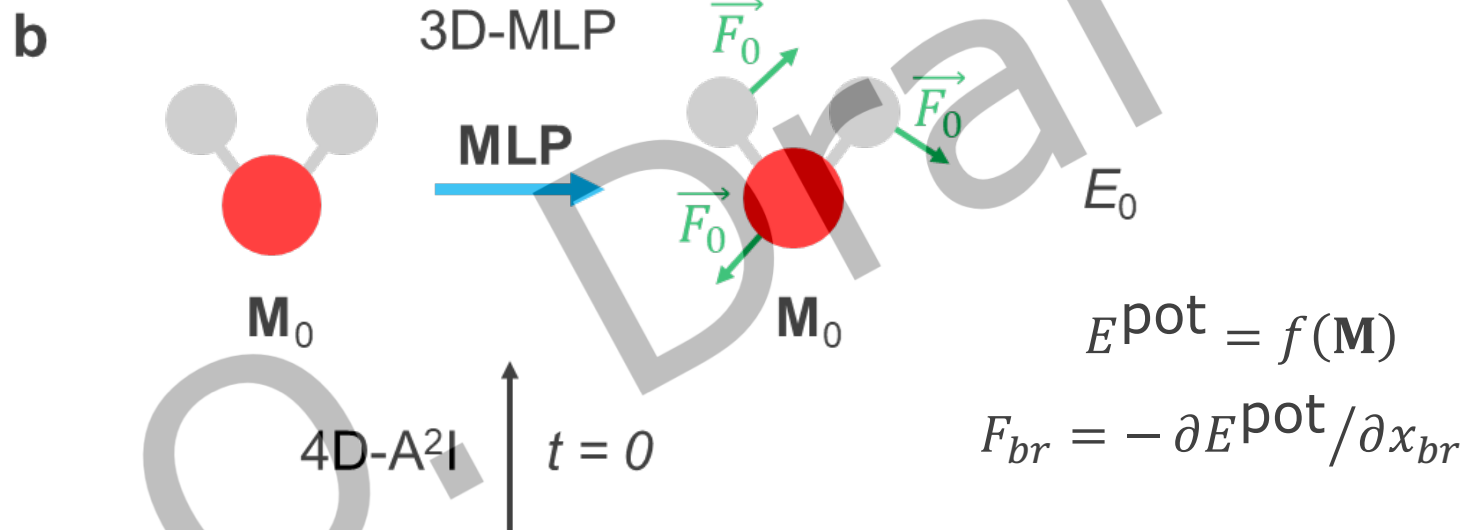
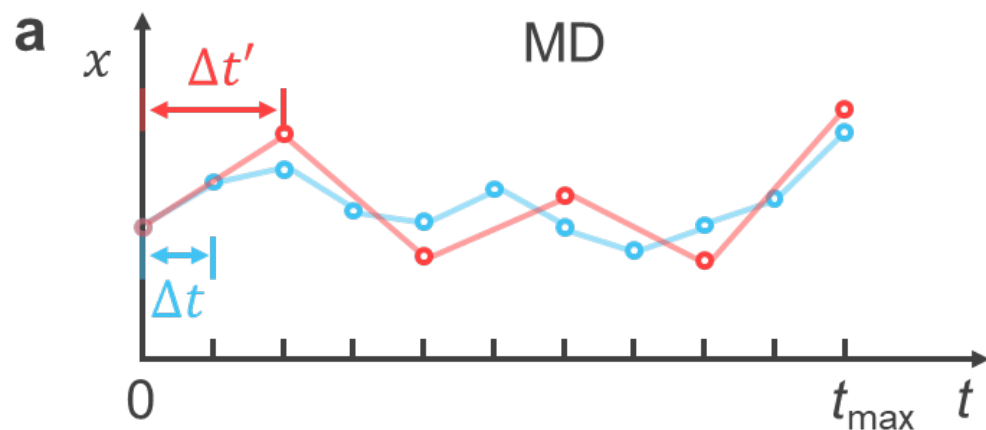
PDB code: 3ENI



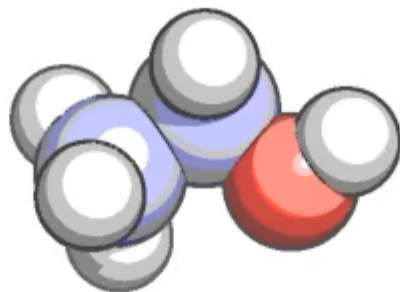
**Dots: Reference**  
**Line: AI-QD**

## 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



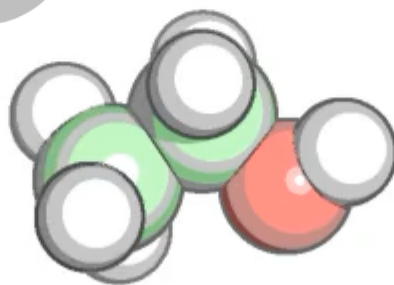
**Ref**



Very slow

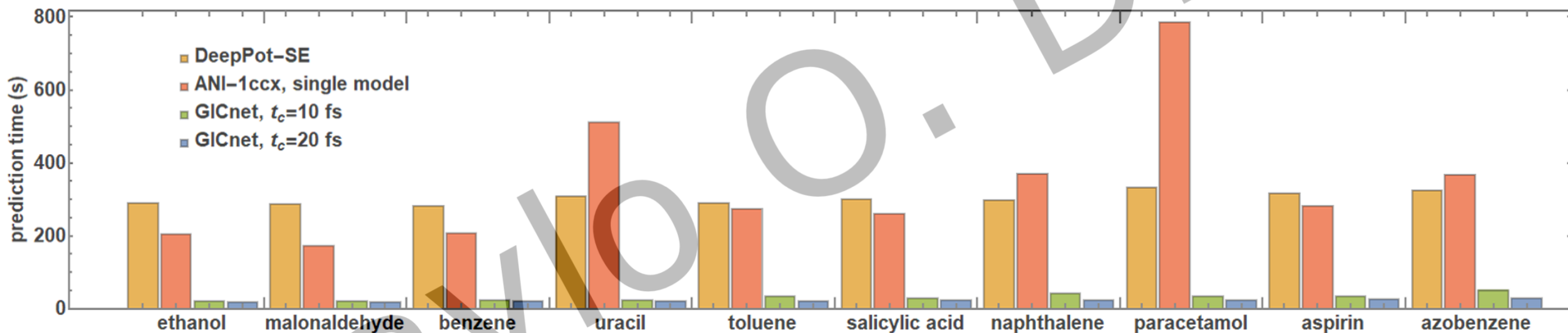
**Stable dynamics for > 1 ns**

**GICnet**



Very fast, e.g., 1 ps  
trajectory (time step 0.05 fs)  
within 1 minute

**0.0 fs**



# Can we do even better?

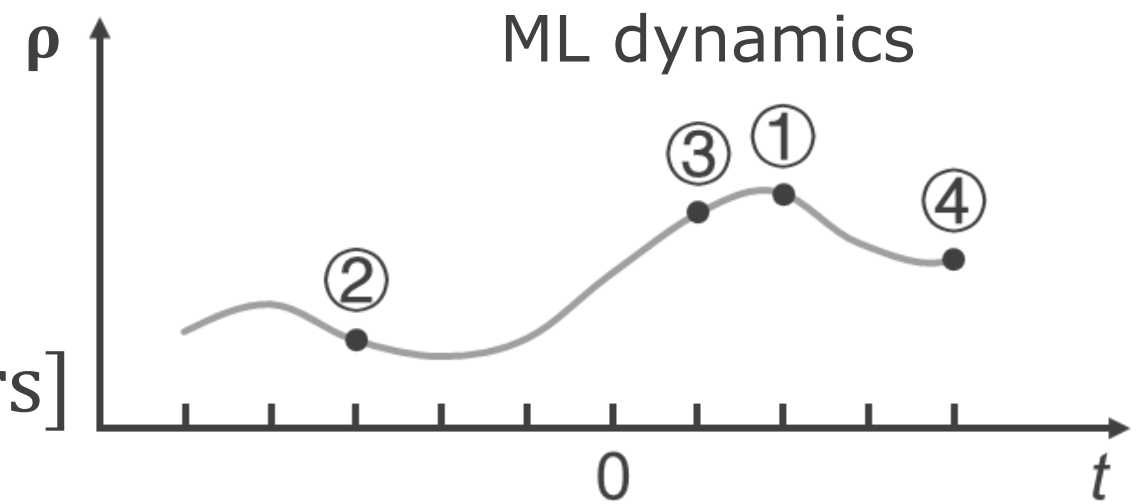
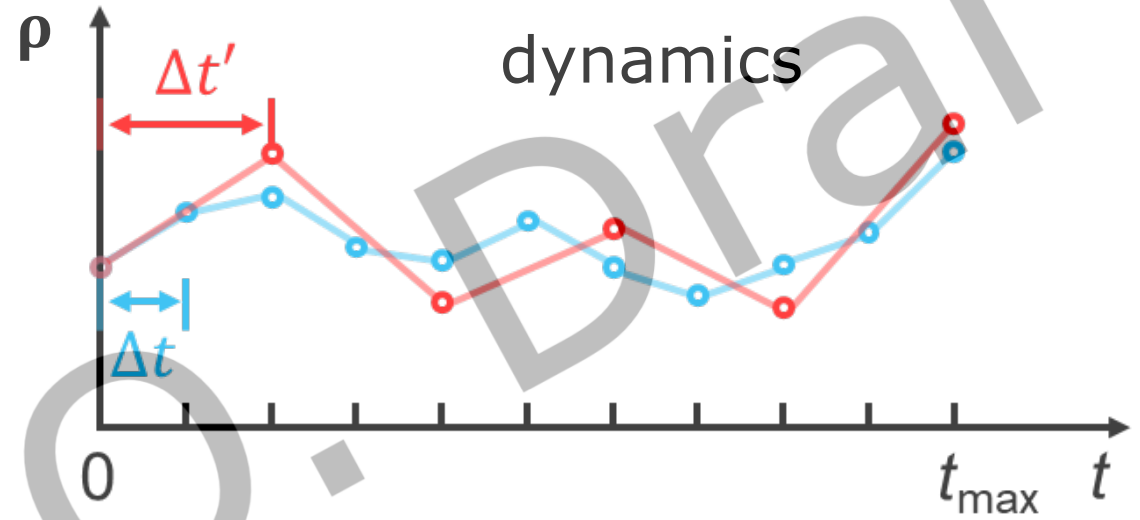
~~$$\rho(t) = f[\rho(t - \Delta t)]$$~~

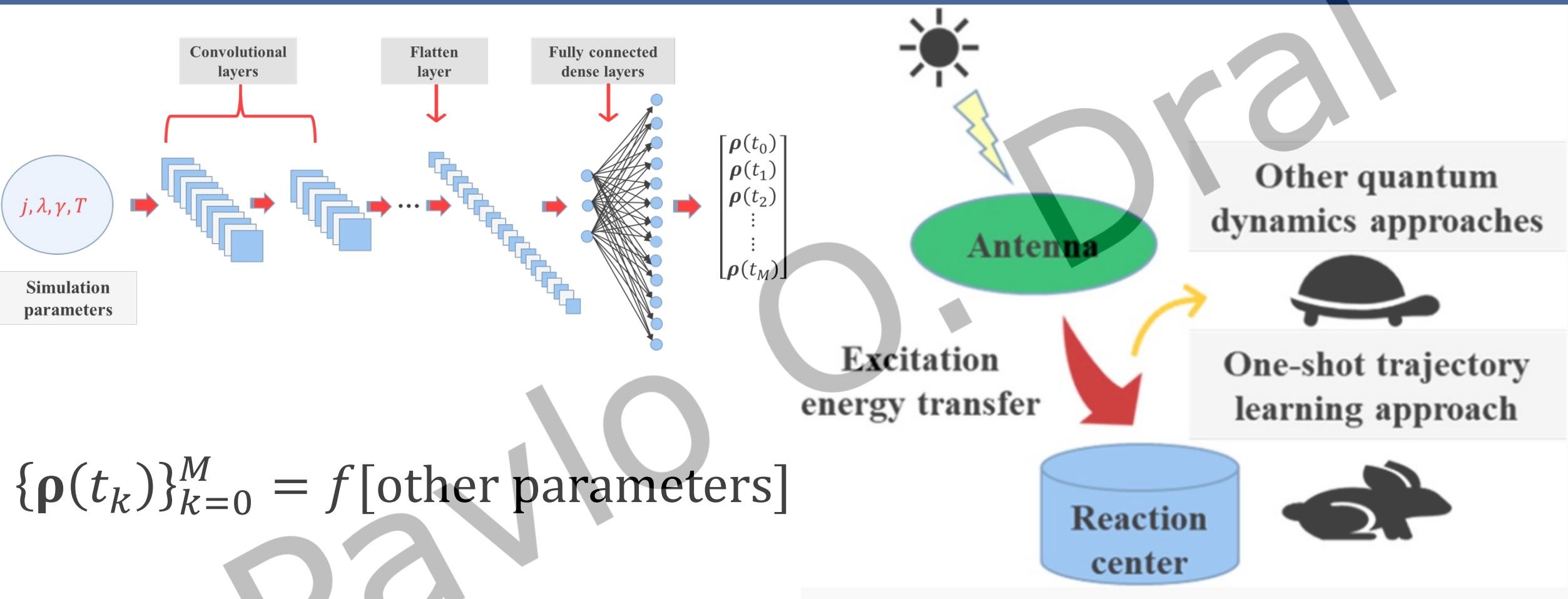
dynamics propagation is

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

~~$$\rho(t) = f[t; \text{other parameters}]$$~~

$$\{\rho(t_k)\}_{k=0}^M = f[\text{other parameters}]$$





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

TA的视频

36

最新发布

最多播放

最多收藏

播放全部 更多 >

Now available in MLatom 3.10 **NEW**

Three days of active learning for surface hopping

MS-ANI, 5950 points, S<sub>2</sub>  
MS-ANI, 5950 points, S<sub>1</sub>  
CASSCF, S<sub>2</sub>  
CASSCF, S<sub>1</sub>

Time (fs)

39:05

面跳跃动力学的主动学习

98 21小时前

Directly! MD

ICnet

Molecular Trajectory

F. Ge, L. Zhang, Y.-F. Hou, Y. Chen, A. Ullah, P. O. J. Phys. Chem. Lett. 2023, 14, 7732

01:48

直接学习分子动力学轨迹!

2923 7-17

molecular\_database molecule atom

database properties molecular properties atomic properties

database methods molecular methods atomic methods

molecular\_trajectory trajectory

01:45

MLatom@XACS中的Python数据对象详解

179 8-14

100 accurate trajectories in 1-2 h online!

03:31

在线模拟两可反应!

571 7-10

B3LYP wB97X-V DSD-BLYP

DENS24

00:55

选DFT的泛函? 不如全都要!

797 8-7

Active learning in MLatom 3.7.0i!

Somersault (4.6%)

01:20

主动学习: 数据高效的机器学习势构建

549 7-3

You asked and we delivered

MLatom.com

Periodic boundary condition in MLatom 3

00:41

终于! MLatom@XACS支持周期性边界条件啦!

559 7-31

CCSD(T) UAQM auto-selection

Computational timecost Accuracy

UAQM self-improvement

CCSD(T) UAQM self-improvement

Computational timecost Accuracy

02:30

UAQM: 通用且可更新的AI模型让你火力全开!

1122 6-27

Scattering Activity (A<sub>2</sub>/A<sub>1</sub>)

Frequency (cm<sup>-1</sup>)

00:35

分子拉曼光谱模拟现在上线啦!

2435 7-24

Intensity (arbitrary unit)

Frequency (cm<sup>-1</sup>)

00:24

分子红外光谱模拟现在上线啦!

3543 6-20





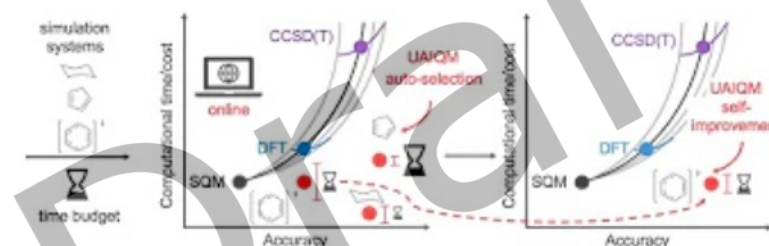
# XACS

@XACScloud · 184 subscribers · 41 videos

Online resources with AI, computational chemistry & chemical bonding analysis ...more

[xacscloud.com](http://xacscloud.com)

Subscribed



Preprint: <https://doi.org/10.26434/chemrxiv-2024-604wb>

2:30

## Supercharge your computational chemistry with the universal and...

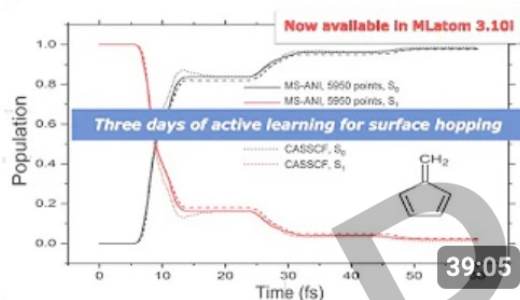
165 views · 1 month ago

Home Videos Shorts Live Playlists

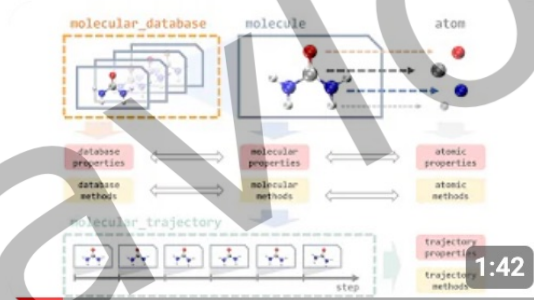
Latest

Popular

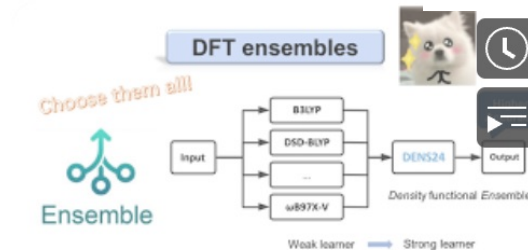
Oldest



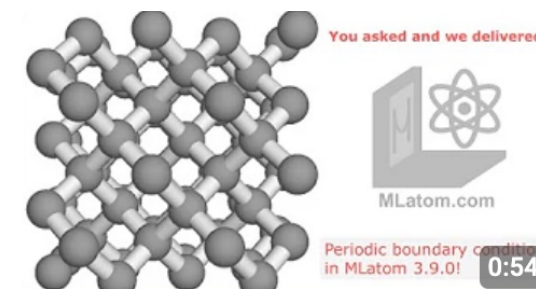
39:05



1:42



Y. Rui, Y. Chen, E. Ivanova, I. Grabowski, P. O. Dral. The best DFT functional is the ensemble of... <https://doi.org/10.26434/chemrxiv-2024-604wb>



Periodic boundary conditions in MLatom 3.9.0! 0:54

### Active learning for surface hopping dynamics

20 views · 16 hours ago

### Data in MLatom's Python API

53 views · 7 days ago

### Not sure which DFT functional to choose? Choose them all!

84 views · 2 weeks ago

### Finally! Periodic boundary conditions in MLatom

92 views · 3 weeks ago

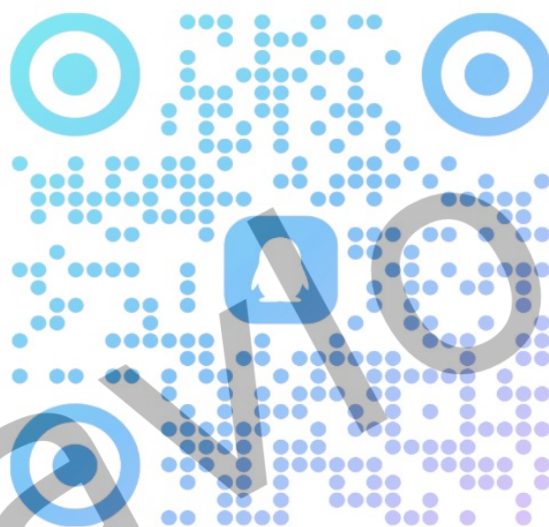
## Wechat

Follow  
XACS account



## QQ Group

Answering your  
questions in real time



## Slack

Answering your  
questions in real time



## Facebook

Join  
the XACS group



Our other channels:

Twitter/X: XACSprogram

Threads: XACSprogram

YouTube: XACScloud

bilibili: XACS团队

Please check the link  
[and register on XACS cloud – optional]



**before lunch break**



Questionary



# Tutorial: ML potentials I

**Pavlo O. Dral**  
Xiamen University, P.R. China

Visiting Professor in  
Nicolaus Copernicus University, Poland

10 September 2024

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

Please check the link  
[and register on XACS cloud – optional]



**before lunch break**



Questionnaire

# Machine Learning



# Simulations



**MLatom**

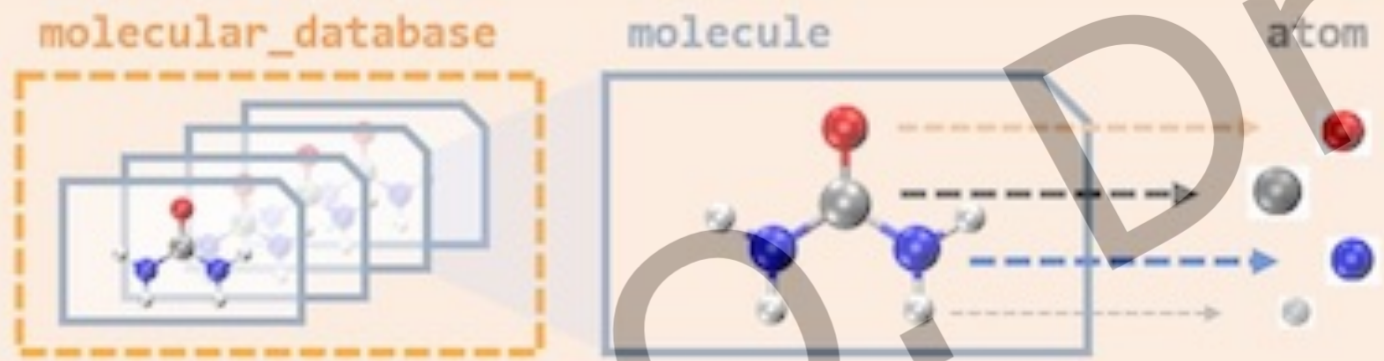


# Quantum Chemistry

# Data

MLatom@GitHub  
≠  
MLatom@XACS

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



➤ Python API **V3**

```
import mlatom as ml
```

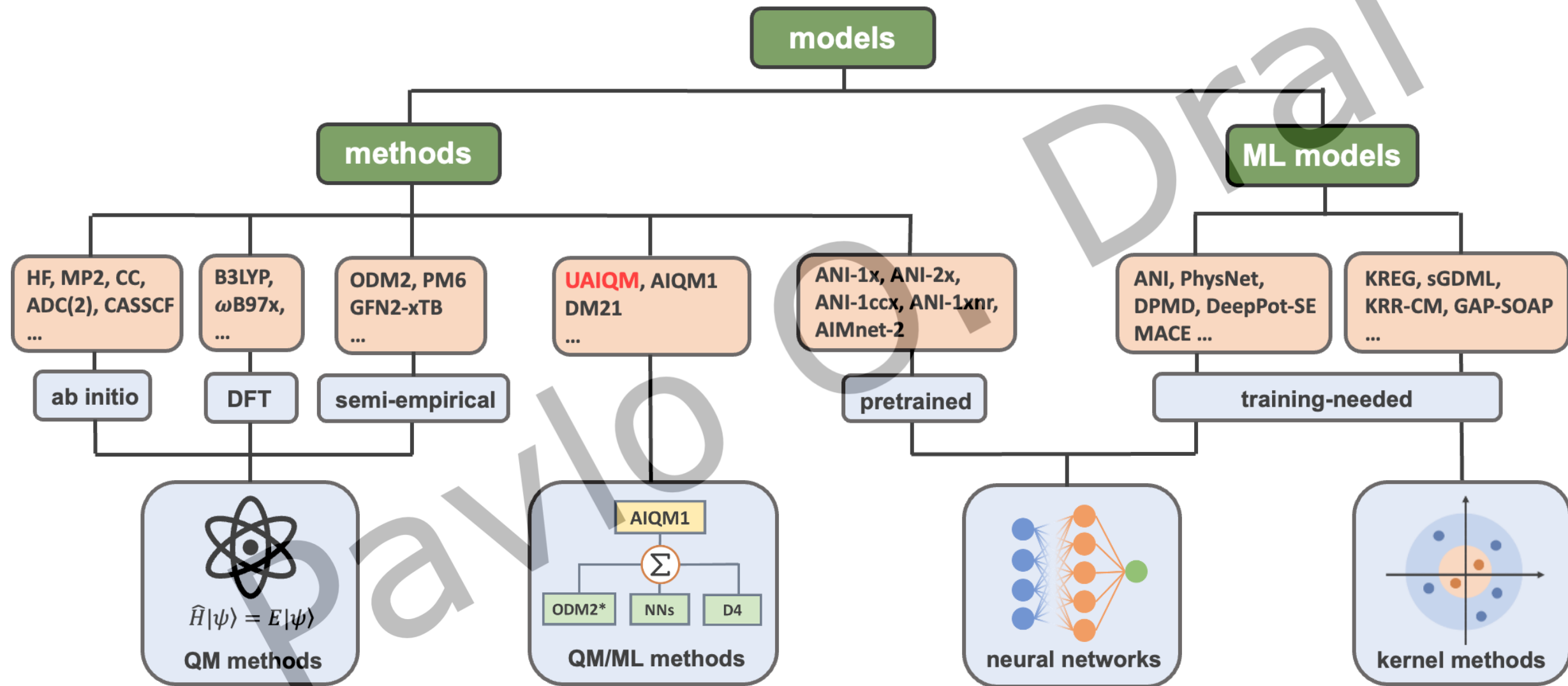


Input file

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

➤ Command line

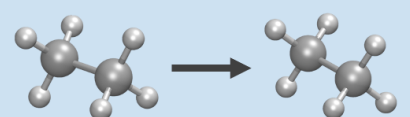
```
> $mlatom ANI-1ccx geomopt xyzfile=init.xyz optxyz=opt.xyz
```



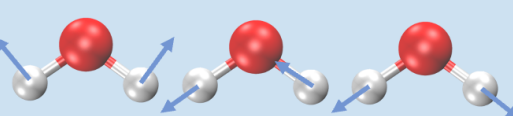


Single point calculations Energies, forces, Hessian matrix...

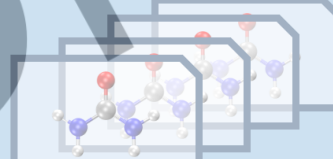
Geometry optimizations



Frequency calculations



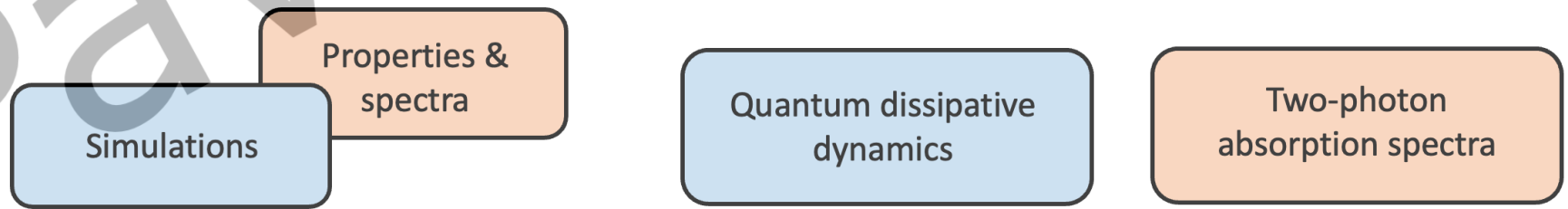
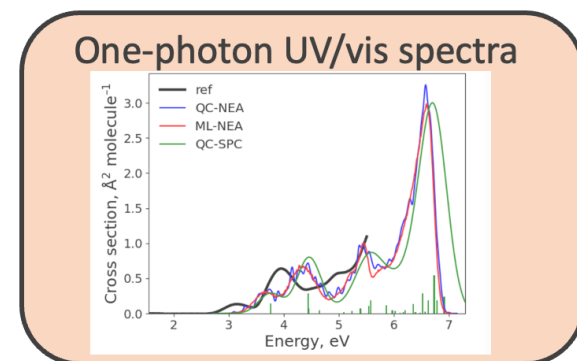
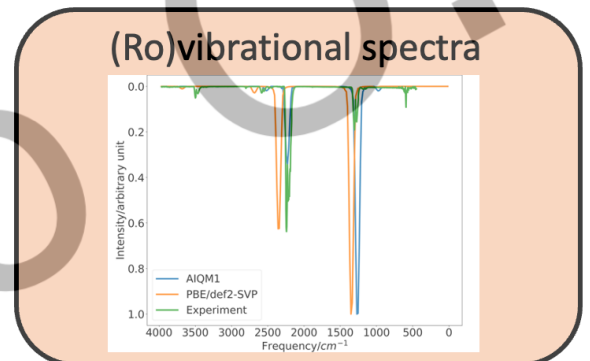
Molecular dynamics

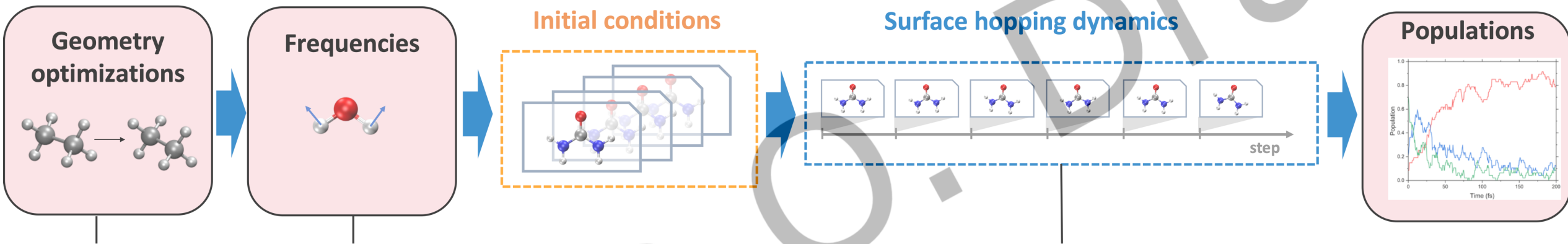


+PIMD coming soon...

Thermochemistry calculations

Heat of formation:

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




**Single point calculations**

$$E, \frac{\partial E}{\partial \mathbf{x}}, \frac{\partial^2 E}{\partial \mathbf{x}^2}$$

```
aiqm1=mlatom.models.methods(method='AIQM1')
geomopt=mlatom.optimize_geometry(model=aiqm1, ...)
...
mlatom.namd.surface_hopping_md(model=model, ...)
```

**Models**

- HF, MP2, CC  
ADC(2), CASSCF  
...
- B3LYP  
ωB97x  
...
- PM6, GFN2-xTB  
AM1, PM3, OMx  
...
- AIQM1  
AIQM1@DFT  
AIQM1@DFT\*
- ANI-1x, ANI-2x  
ANI-1ccx  
...
- ANI, PhysNet, DPMD  
DeepPot-SE, MACE  
...
- KREG, sGDML,  
KRR-CM, GAP-SOAP  
...

## Extras!

### MLQD

A Package for Quantum  
Dissipative Dynamics with  
Machine Learning  
by Arif Ullah, Anhui University

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

Semi-empirical quantum chemical programs:

Machine learning programs:

Dynamics and other atomistic simulation:

**Wechat**  
Follow  
XACS account



**Slack**  
Answering your  
questions in real time



Quantum chemical programs:



Gaussian

PySCF

Turbomole



ORCA

COLUMBUS



SPARROW

MNDO

PhysNet

GAP

TORCHANI



SGML

hyperopt

ASE



Not everything is available on the cloud...

- ⊕ 超参数优化
- ⊕ 测试使用
- ⊕ MACE势:
  - (p)KREG势
  - 基准测试
- Transfer learning
- AIQM1
- Quantum chemical methods
- 通用机器学习模型
- 更多教程

输入文件/命令行的使用手册

- 概览
- 模拟
- 学习
- 数据

PYTHON接口手册

- 概览
- Data
- Models
- Simulations

# Python API

对于KREG模型，我们可以使用简单的网格搜索优化

```

model = ml.models.kreg(model_file=f'kreg.npz')

sub, val = molDB.split(number_of_splits=2, fract:

model.hyperparameters['sigma'].minval = 2**-5 # ,
model.optimize_hyperparameters(subtraining_molec
optimization_alg
hyperparameters=
training_kwargs=
prediction_kwarg:

lmbd = model.hyperparameters['lambda'].value ; s
valloss = model.validation_loss
print('Optimized sigma:', sigma)
print('Optimized lambda:', lmbd)
print('Optimized validation loss:', valloss)
# Train the model with the optimized hyperparame:
model.train(molecular_database=molDB, property_t:
# Train the model with the optimized hyperparame:
model.train(molecular_database=molDB, property_t:

```

输出如下所示（它可能随子训练集和验证集的随机子

```

Optimized sigma: 0.10511205190671434
Optimized lambda: 2.910383045673381e-11
Optimized validation loss: 3.1550365181164988e-01

```

其他参数也是可用的，例如SciPy(Nelder-Mead, BFGS, krylov, trust-exact)和hyperopt库(TPE)。

## Cloud Computing

- 🕒 Job Submitter
- 🖥️ Terminal
- 📁 File Manager
- 📋 Job Manager
- 🌐 Jupyter Lab

## Software

- 📄 Download

## Learning

- 📖 Courses
- 👤 Workshops

## Statistics

Total CPU time used  
4756h 1m

## Job Information

\* Job Name 2024-05-30\_0757

Job Location [🏠](#) > from\_job\_submitter

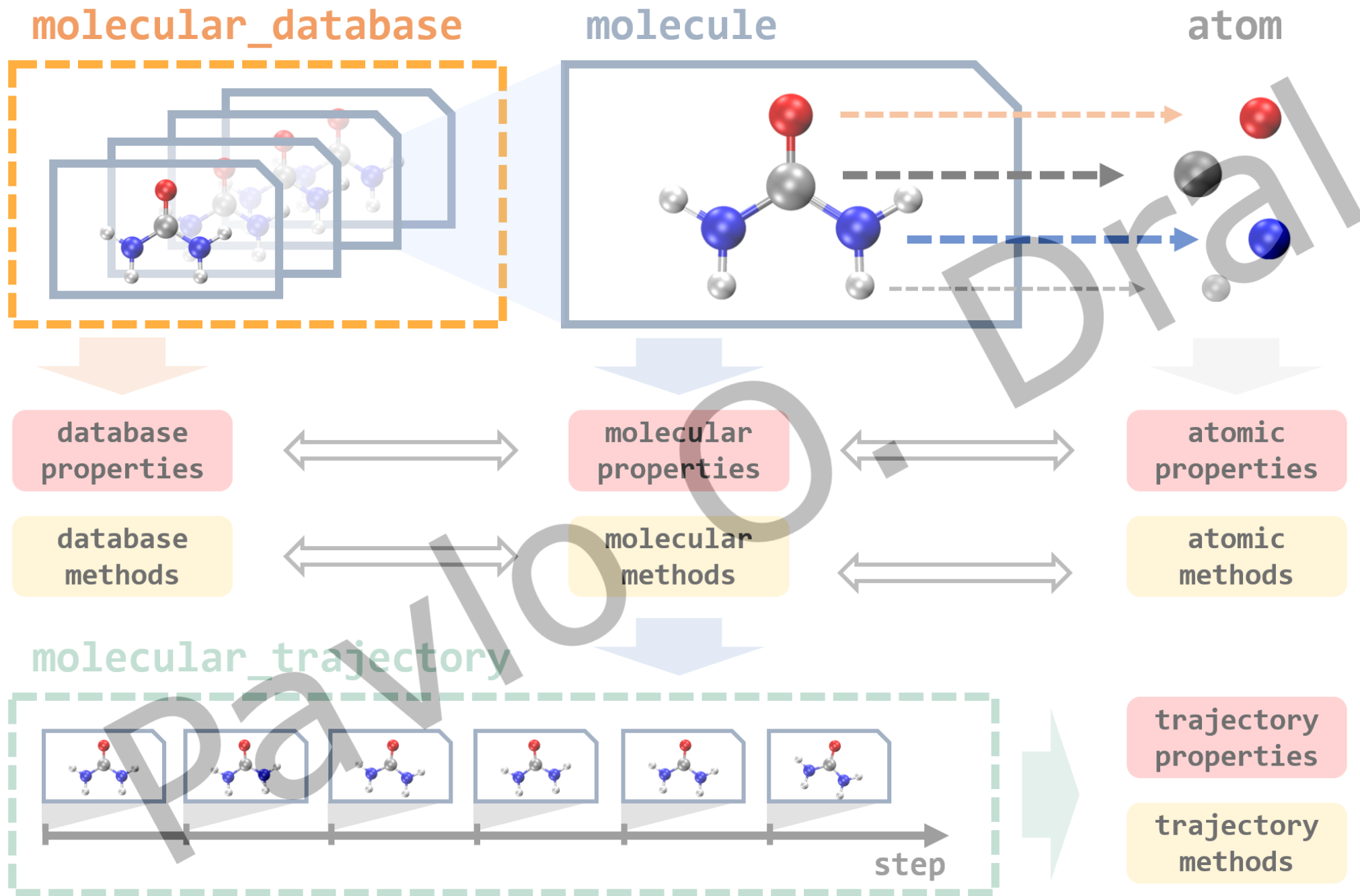
Job Type

- XACS (auto detect)
- Gaussian
- Mlatom\_d

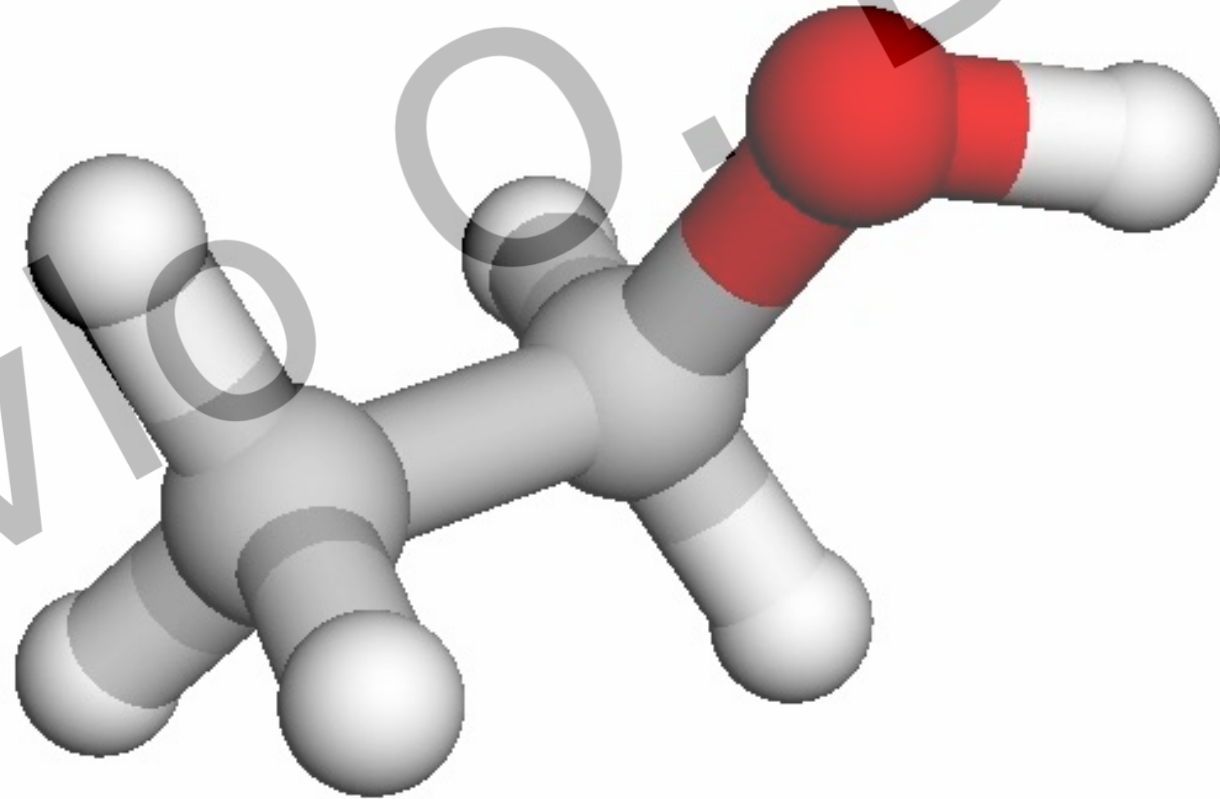
## Input File

📄 or edit XACS input file:

1

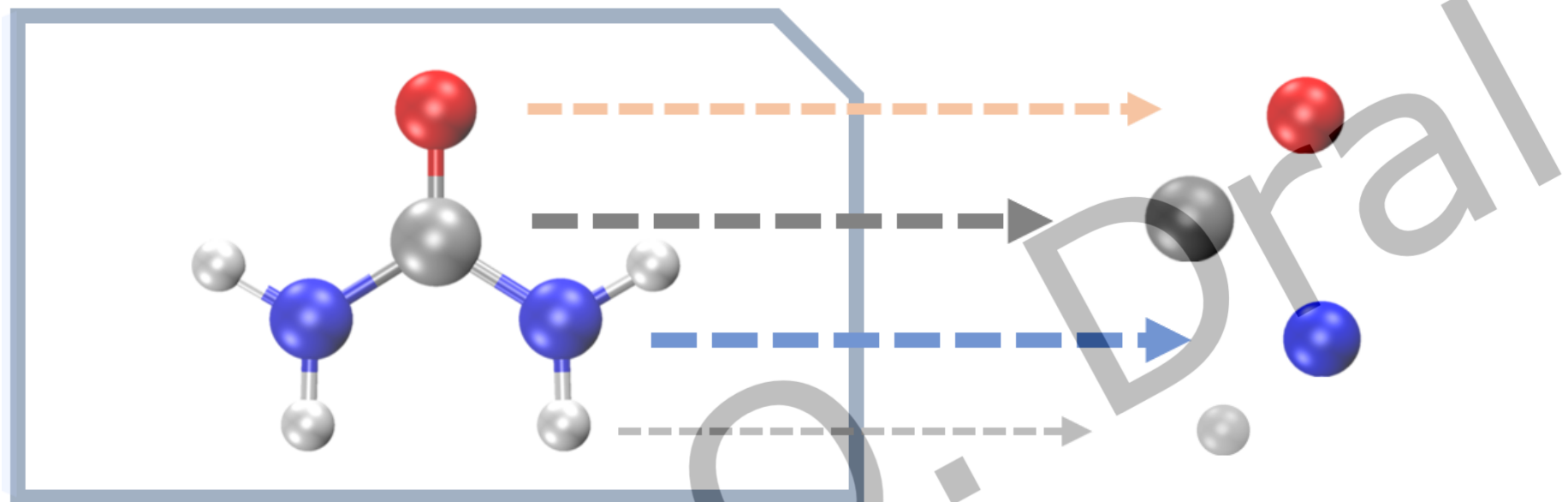


```
molvibr.view(normal_mode=normal_mode)
```



molecule

atom



```
mol = ml.molecule()
mol.read_from_xyz_string('''2
H 0 0 0
H 0 0 0.74
''')
# or
mol.read_from_xyz_file('h2.xyz')
# or
import numpy as np
xyz = np.array([[0, 0, 0], [0, 0, 0.7]])
mol.read_from_numpy(coordinates=xyz, species=np.array([1, 1]))
```

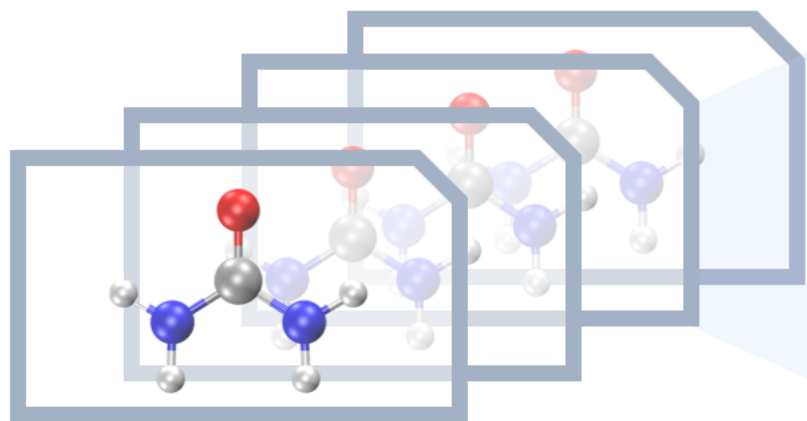
```
mol.dump(filename='h2.json')
mol2.load(filename='h2.json')
```

```
mol.charge = 1
mol.multiplicity = 2
```

molecular\_database

molecule

atom



```
# prepare H2 geometries with bond lengths ranging from 0.5 to 5.0 Å
xyz = np.zeros((451, 2, 3))
xyz[:, 1, 2] = np.arange(0.5, 5.01, 0.01)
z = np.ones((451, 2)).astype(int)
molDB = ml.molecular_database.from_numpy(coordinates=xyz, species=z)

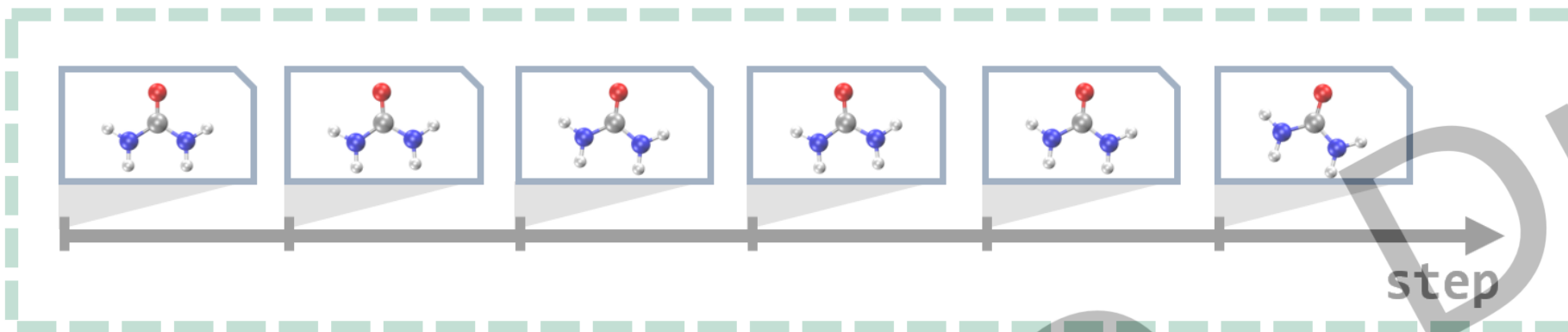
molDB.dump(filename='h2.json')
molDB2 = ml.molecular_database.load(filename='h2.json')
```

```
print(len(molDB))           # 451
print(len(molDB[:100:4]))  # 25
```

```
# let's change the splitting to 9:1 instead of default 8:2
train, test = molDB.split(fraction_of_points_in_splits=[0.9, 0.1])
```



## molecular\_trajectory



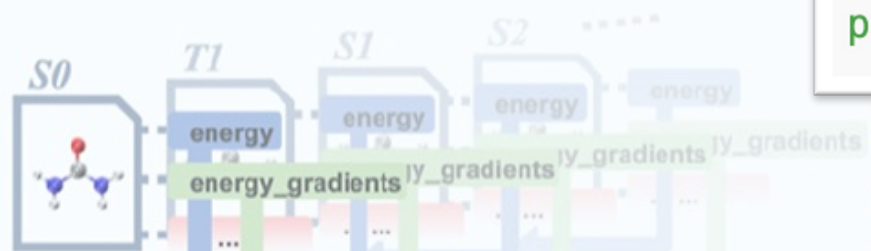
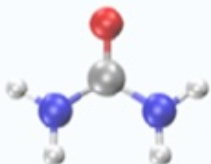
trajectory  
properties

trajectory  
methods

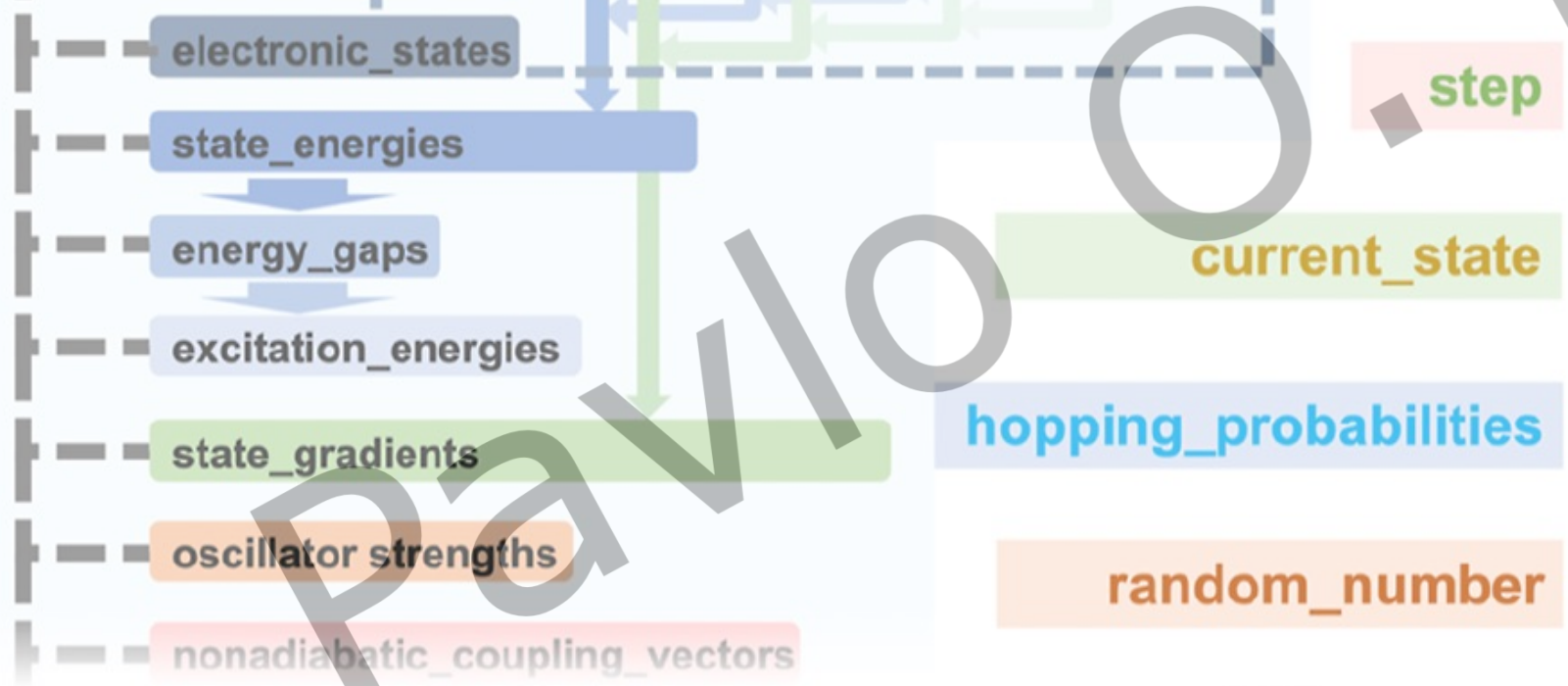
```
traj = ml.data.molecular_trajectory()  
...  
print(f'Info for MD trajectory step {traj.steps[5].step}')  
print(f'Molecule: {traj.steps[5].molecule}')  
print(f'Time: {traj.steps[5].time}')  
print(f'Kinetic energy: {traj.steps[5].molecule.kinetic_energy}')
```

# trajectory\_step

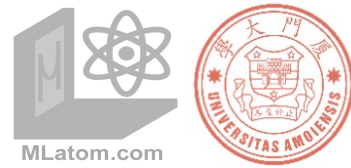
## molecule



```
# let's get S1 state
S1 = mol.electronic_states[1]
print(f'Energy of S1 state is {S1.energy}')
print(f'Its forces are')
print(-S1.energy_gradients)
print(f'checking multiplicity: {S1.multiplicity}')
```



**Subscribe to updates and submit your papers to the AI/ML journals**



Ukraine: Click here to read IOP Publishing's statement



### Machine Learning: Science and Technology

Home > Journals > Machine Learning: Science and Technology > Editorial board

Follow:

CALL FOR PAPERS

# Artificial Intelligence Chemistry

## Open for Submissions



Homepage

Author guidelines

About Machine Learning: Science and Technology

Editorial board

Track my article

shingsupport.iopscience.iop.org



## 3<sup>rd</sup> International Symposium on Machine Learning in Quantum Chemistry (SMLQC)

University of Tennessee, Knoxville

Fall 2025

Dates TBD