



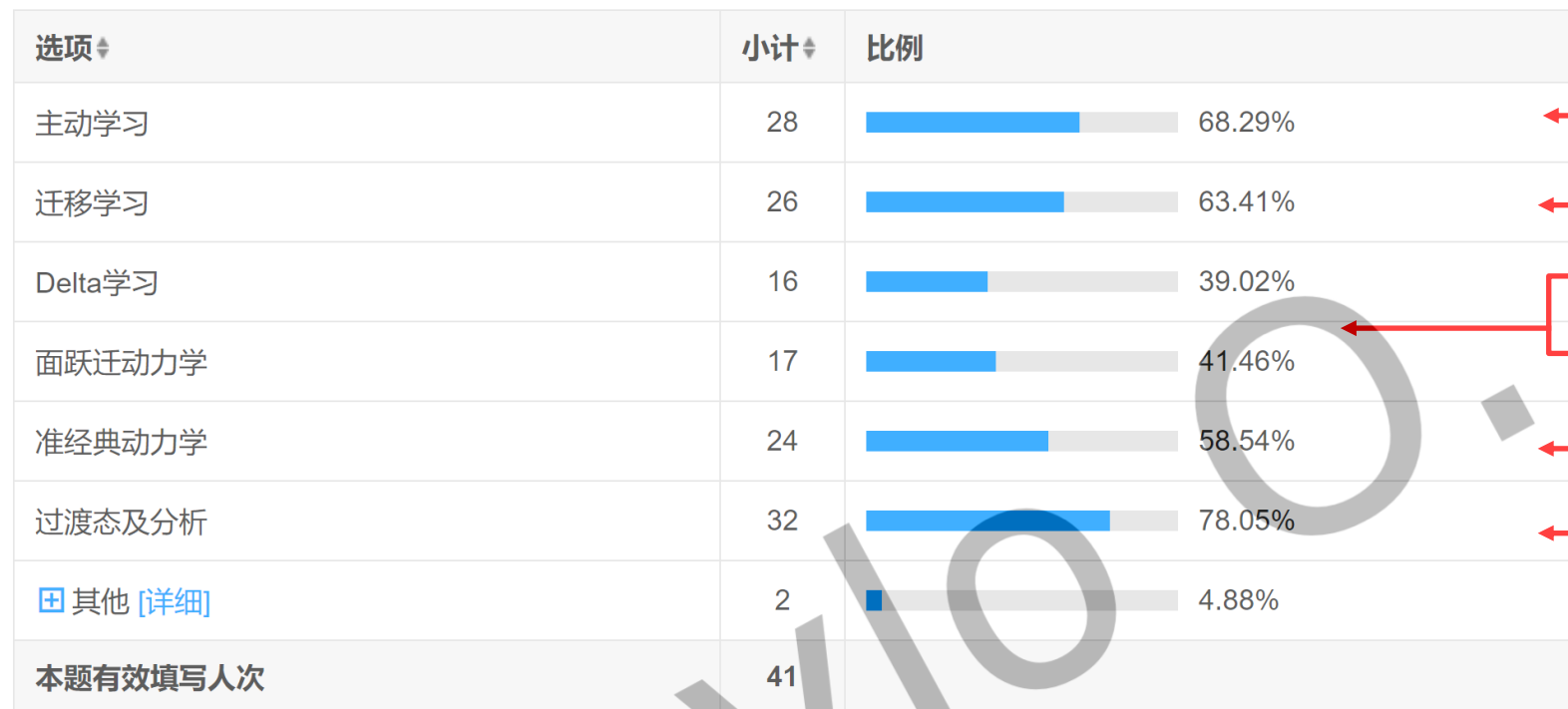
Advanced course on ML

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5 July 2024

第3题：您对机器学习的哪些课程比较感兴趣（可多选） [多选题]



4. Active learning (Python)

1. Transfer learning (Python)

5-6 – remaining time

2. MD (quasi-classical)

3. TS

序号	提交答卷时间	答案文本	查看答卷
42	7月4日 14:02	LLM, feature engineering	查看答卷
44	7月4日 14:52	没有ml基础不太懂，所以想学	查看答卷

Slides added to ML basics



8. Charge-shift bonding in propellane

MACHINE LEARNING

1. Machine learning basics

1.1. Slides

1.2. Power of ML

1.3. Training your first ML model

1.4. ML algorithms

2. ML for PES

3. Universal machine learning models

4. Spectroscopy

5. Transfer learning

6. Molecular dynamics

7. Transition state search and analysis

8. Active learning

9. Delta-learning

10. Surface-hopping nonadiabatic dynamics

1. Machine learning basics

1.1. Slides

↓ Slides:

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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

79

80

81

Linear regression

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 Φ :

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

In previous examples we new 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 in 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:

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

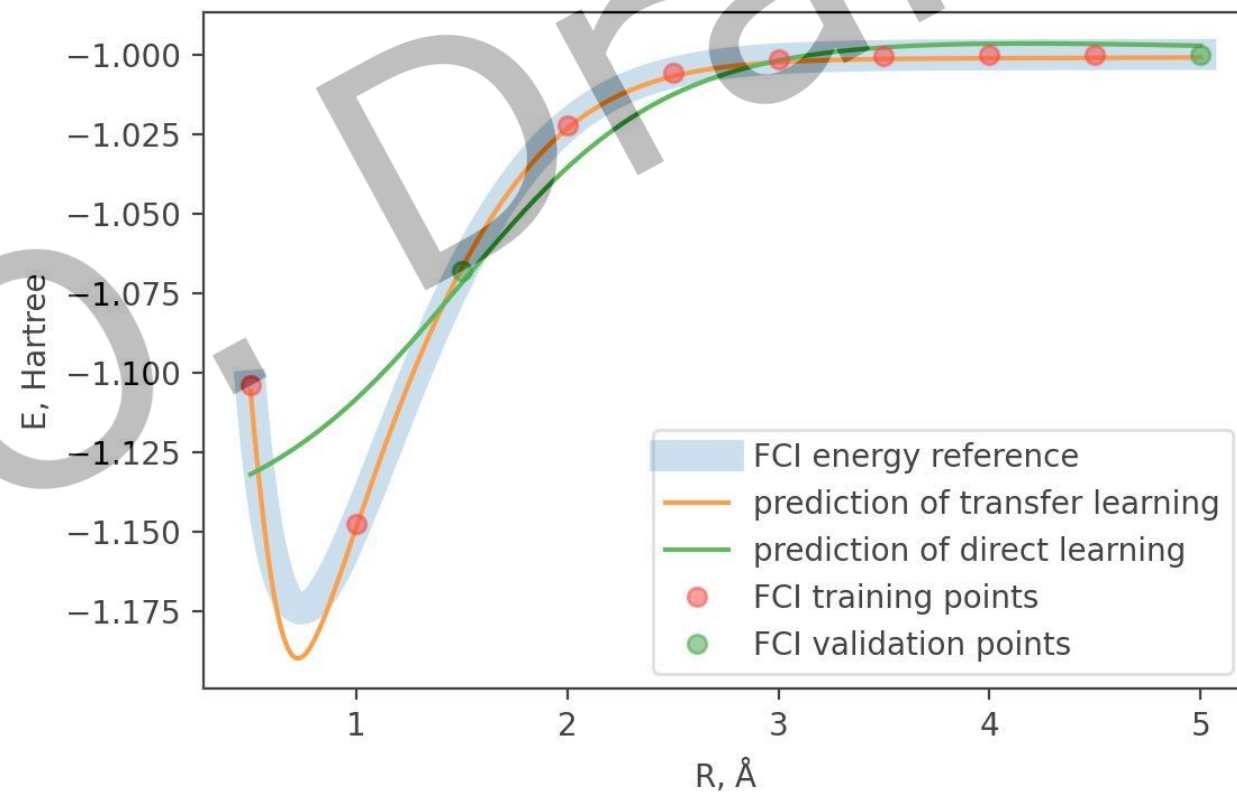
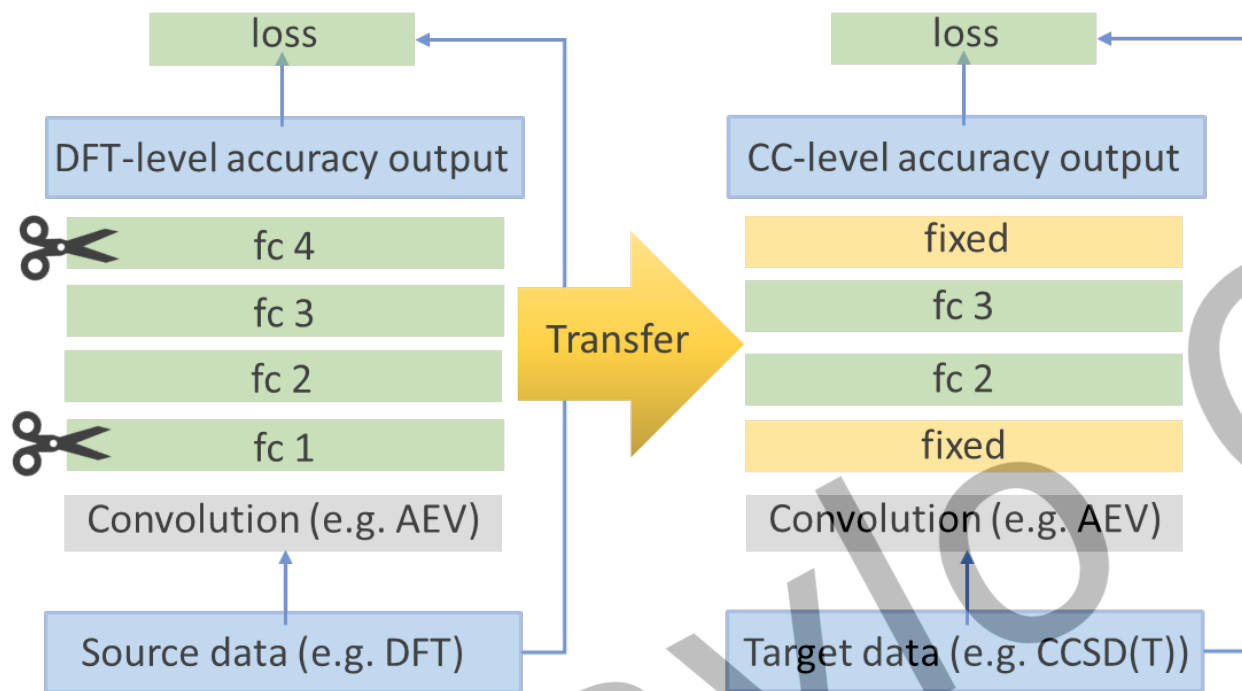



Transfer learning

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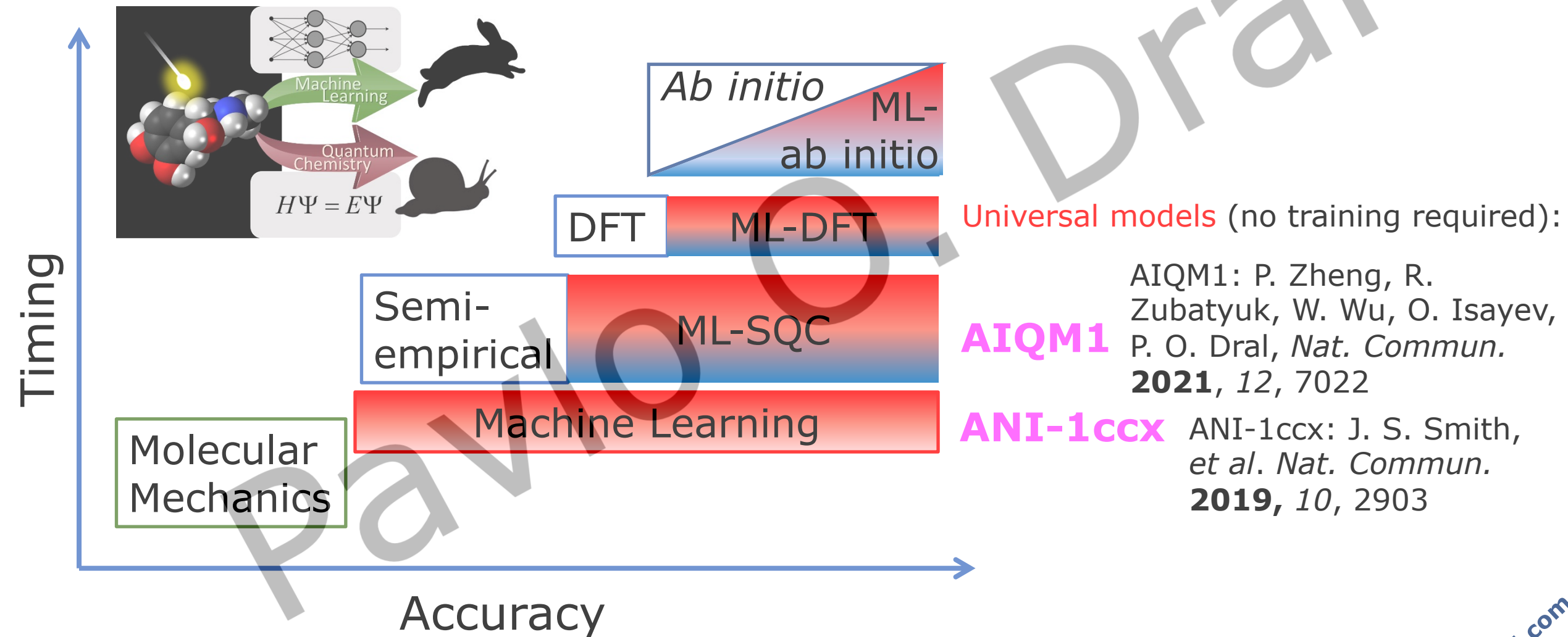
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Figures: Pavlo O. Dral, Tetiana Zubatiuk, Bao-Xin Xue, Learning from multiple quantum chemical methods: Δ -learning, transfer learning, co-kriging, and beyond. In *Quantum Chemistry in the Age of Machine Learning*, Pavlo O. Dral, Ed. Elsevier: **2023**.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

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



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

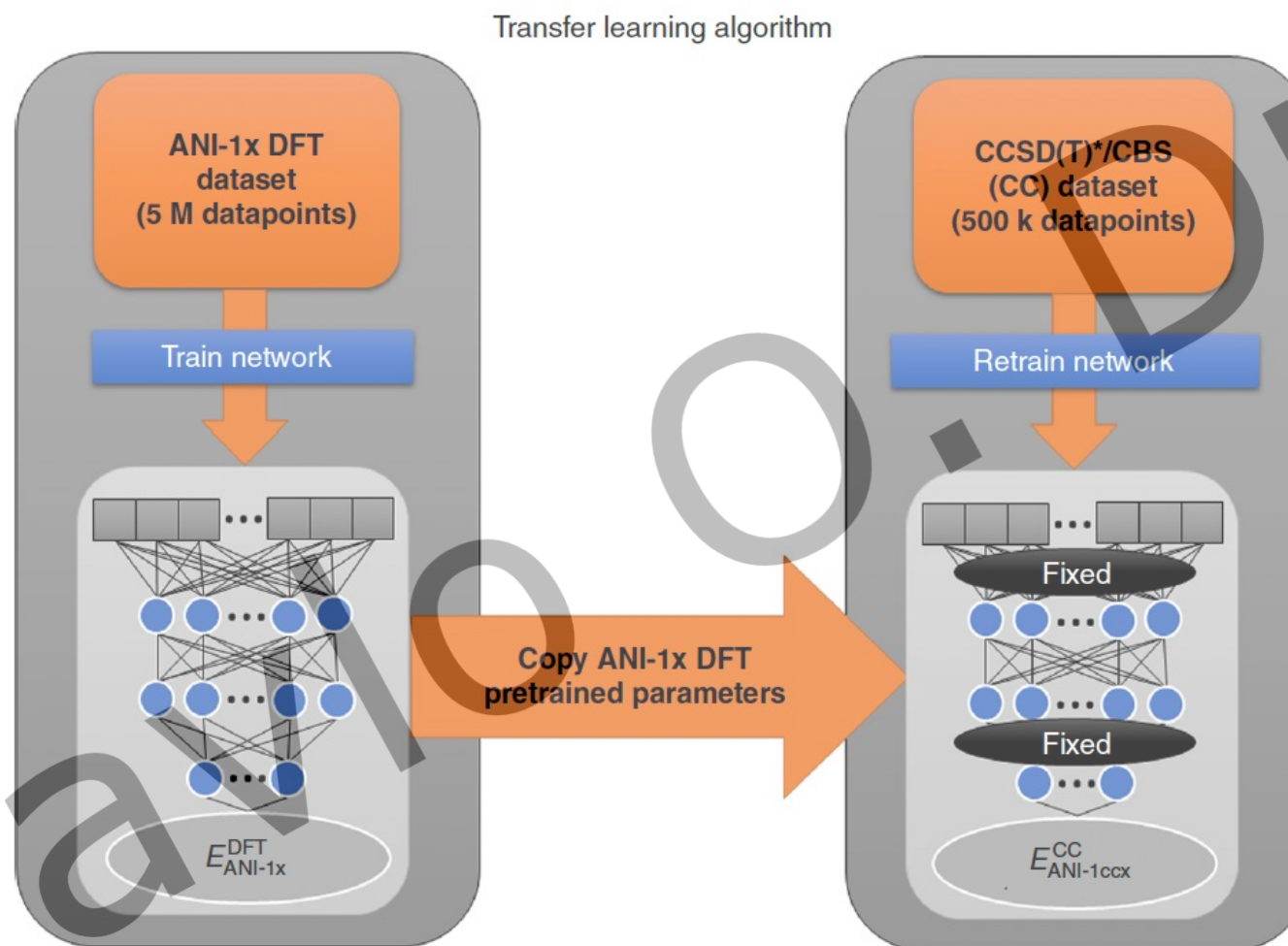
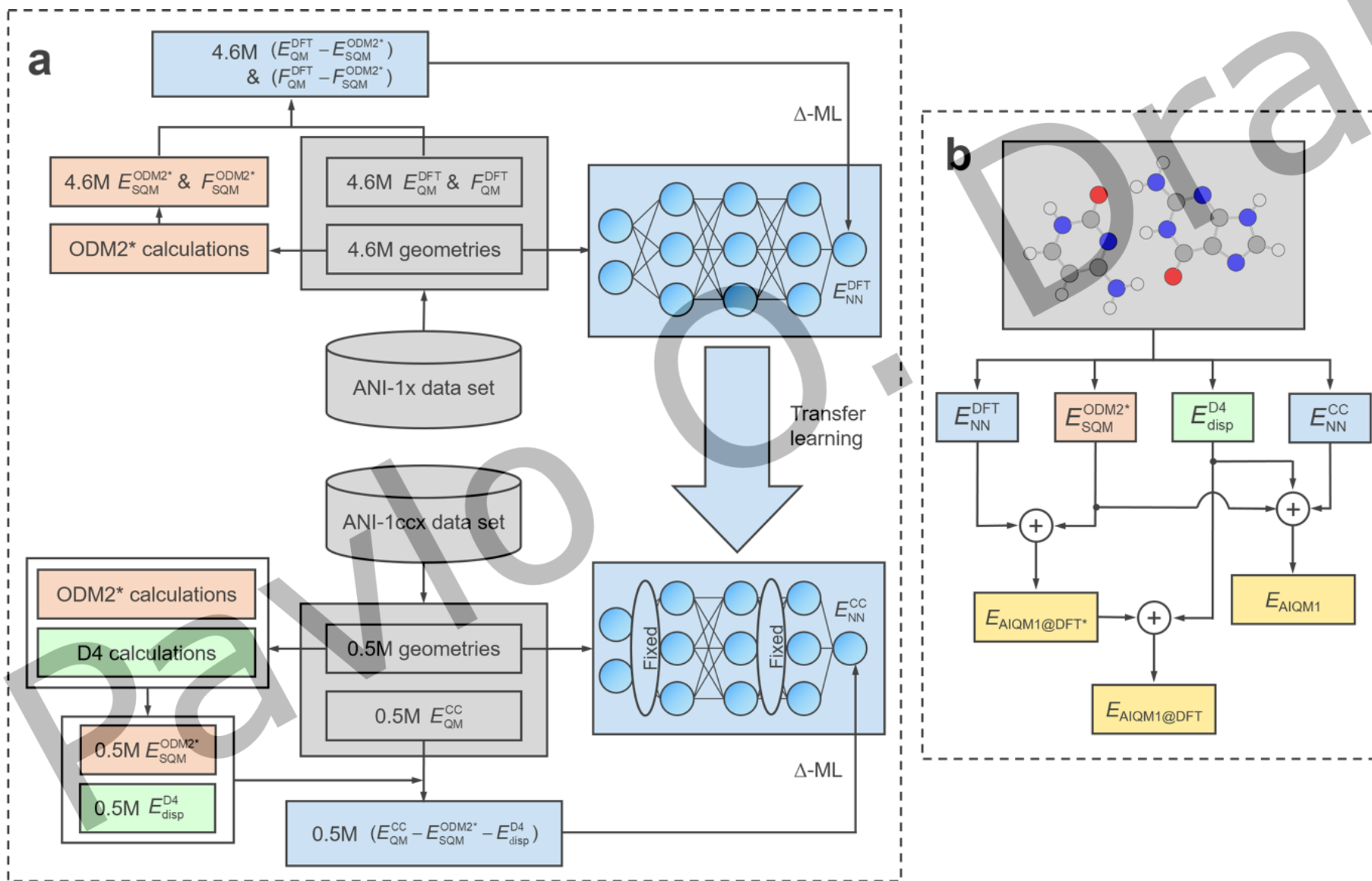


Fig. 4 Diagram of the transfer learning technique evaluated in this work. Transfer learning starts from a pretrained ANI-1x DFT model, then retrains to higher accuracy CCSD(T)* / CBS data with some parameters fixed during training



methods

6. Computing of diabatic states with VB theory

7. Menshutkin Reaction $\text{NH}_3 + \text{CH}_3\text{Cl} \rightarrow [\text{NH}_3\text{CH}_3]^+ + \text{Cl}^-$

8. Charge-shift bonding in propellane

MACHINE LEARNING

1. Machine learning basics

2. ML for PES

3. Universal machine learning models

4. Spectroscopy

5. Transfer learning

5.1. Jupyter notebook

5.2. Tutorial

5.3. Transfer learning in MLatom

5.4. ANI TL model

5.1. Jupyter notebook

- [📄 tutorial notebook](#)

Download this Jupyter notebook

You can upload it to the XACS cloud, launch Jupyter lab (click 'Connect' after launching), and open it in the Jupyter lab.

5.2. Tutorial

Transfer learning (TL) is an often-used technique in machine learning that helps you train better neural network models. TL transfers knowledge (parameters) of the pre-trained model from a different task, by fixing and changing some of its layers and there are several reasons that we might utilize this technique:

- **Better training performance**

The pre-trained models can give you a good starting point to kick off the training, which helps the convergence of the final model.

- **Less data hungry**

Training from a pre-trained model does not require data as much as training from scratch, since the pre-trained model has information from its training data. This can be critical, especially when the accessibility to the data for the new task is limited.

Cloud Computing

 Job Submitter

 Terminal

 File Manager

 Job Manager

 Jupyter Lab

Jupyter Lab



The server is running.

ID: 1727477

Expiration 2024-07-08 01:20:34

Connect

Stop

Upload Jupyter notebook there

Start Jupyter Lab

Pavlo

Dral

tl.ipynb

Code

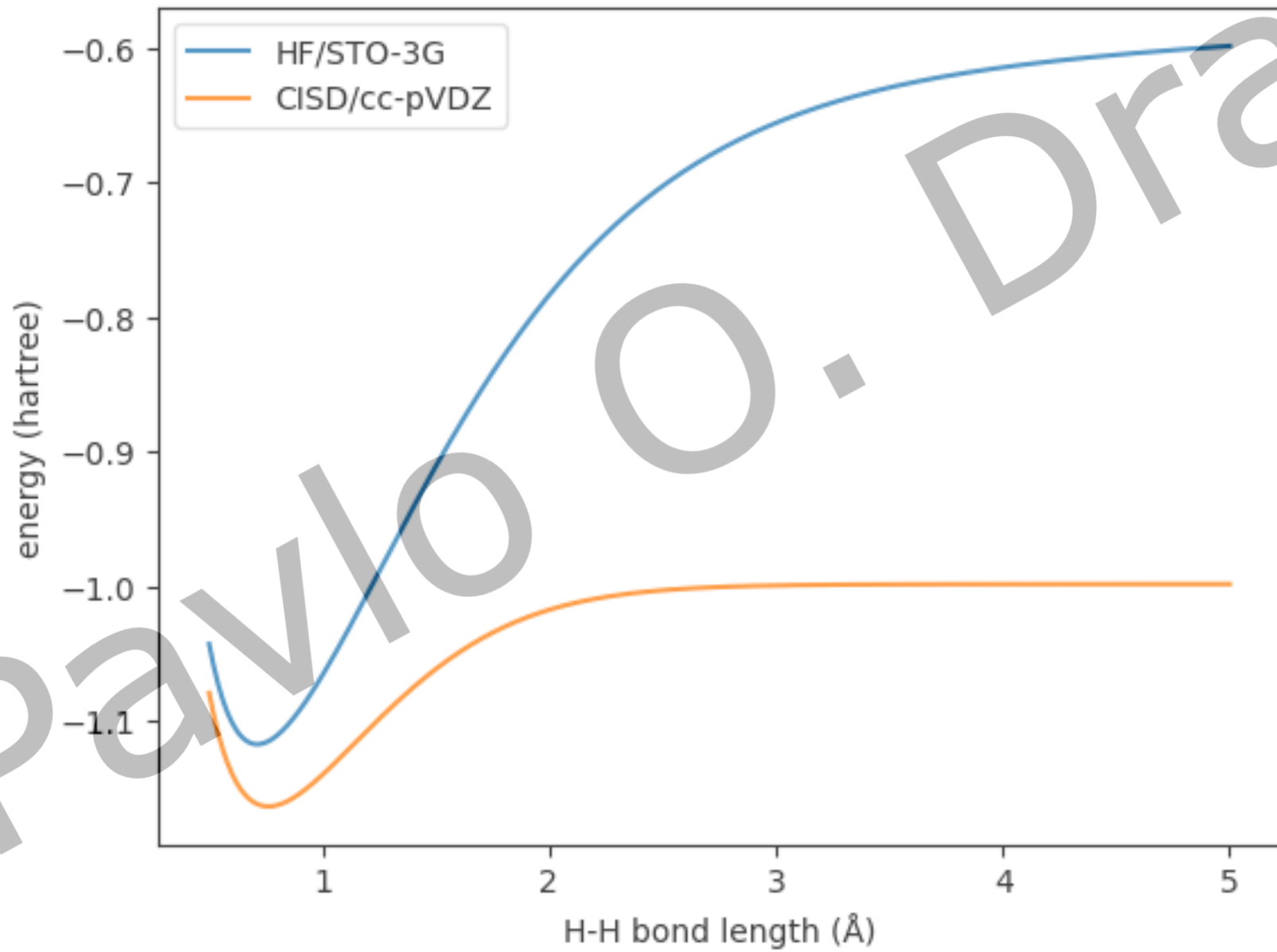
```
[1]: import mlatom as ml
import numpy as np
import matplotlib.pyplot as plt

[2]: # 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)

[3]: # calculate HF energies
hf = ml.models.methods(method='HF/STO-3G', program='PySCF')
hf.predict(molecular_database=molDB, calculate_energy=True)
molDB.add_scalar_properties(molDB.get_properties('energy'), 'HF_energy') # save HF energy with a new name

[4]: # calculate CISD energies
cisd = ml.models.methods(method='CISD/cc-pVDZ', program='PySCF')
cisd.predict(molecular_database=molDB, calculate_energy=True)
molDB.add_scalar_properties(molDB.get_properties('energy'), 'CISD_energy')

[5]: # plot the energies
plt.plot(xyz[:, 1, 2], molDB.get_properties('HF_energy'), label='HF/STO-3G')
plt.plot(xyz[:, 1, 2], molDB.get_properties('CISD_energy'), label='CISD/cc-pVDZ')
plt.legend()
plt.xlabel('H-H bond length (Å)')
plt.ylabel('energy (hartree)')
plt.show()
```




```
[6]: # train ANI model with HF energies
! rm ANI-HF.pt &> /dev/null
ani = ml.models.ani(model_file='ANI-HF.pt', verbose=False)
ani.train(molecular_database=molDB, property_to_learn='HF_energy')
# predict with trained ANI model
ani.predict(molecular_database=molDB, property_to_predict='ANI_HF_energy')
```

```
[7]: # show the model structure
ani.model
```

```
[7]: Sequential(
  (0): AEVComputer()
  (1): ANIModel(
    (H): Sequential(
      (0): Linear(in_features=48, out_features=160, bias=True)
      (1): CELU(alpha=0.1)
      (2): Linear(in_features=160, out_features=128, bias=True)
      (3): CELU(alpha=0.1)
      (4): Linear(in_features=128, out_features=96, bias=True)
      (5): CELU(alpha=0.1)
      (6): Linear(in_features=96, out_features=1, bias=True)
    )
  )
)
```

```
[8]: # fix some of the layers
! cp ANI-HF.pt ANI-HF-TL.pt
ani = ml.models.ani(model_file='ANI-HF-TL.pt', verbose=False)
ani.fix_layers([[0, 6]])
# transfer leaning with every 40th of the data
step = 40
val = molDB[::step][::10]
sub = ml.molecular_database([mol for mol in molDB[::step] if mol not in val])
ani.energy_shifter.self_energies = None # let the model recalculate the self atomic energies
ani.train(molecular_database=sub, validation_molecular_database=val, property_to_learn='CISD_energy',
          hyperparameters={'learning_rate': 0.0001}, reset_optimizer=True)
# predict with TL model
ani.predict(molecular_database=molDB, property_to_predict='ANI_TL_energy')
```

```
[9]: # train ANI model with CISD energies directly
! rm ANI_CISD.pt &> /dev/null
ani_cisd = ml.models.ani(model_file='ANI_CISD.pt', verbose=False)
ani_cisd.train(molecular_database=sub, validation_molecular_database=val, property_to_learn='CISD_energy')
# predict with trained ANI model
ani_cisd.predict(molecular_database=molDB, property_to_predict='ANI_CISD_energy')
```

```
[10]: # plot the energies
plt.plot(xyz[:, 1, 2], molDB.get_properties('HF_energy'), label='HF/STO-3G')
plt.plot(xyz[:, 1, 2], molDB.get_properties('CISD_energy'), label='CISD/cc-pVDZ')
plt.plot(xyz[:, 1, 2], molDB.get_properties('ANI_HF_energy'), label='ANI-HF')
plt.plot(xyz[:, 1, 2], molDB.get_properties('ANI_TL_energy'), label='ANI-TL')
plt.plot(xyz[:, 1, 2], molDB.get_properties('ANI_CISD_energy'), label='ANI-CISD')
plt.plot(sub.xyz_coordinates[:, 1, 2], sub.get_properties('CISD_energy'), 'o', label='TL subtraining')
plt.plot(val.xyz_coordinates[:, 1, 2], val.get_properties('CISD_energy'), 'o', label='TL validation')
plt.legend()
plt.xlabel('H-H bond length (Å)')
plt.ylabel('energy (hartree)')
plt.show()
```

