

# Machine learning and chemistry

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the Kulik group at MIT, [hjkgrp.mit.edu/](http://hjkgrp.mit.edu/)

under the supervision of Professor Heather J. Kulik

for the most recent version and demos: [github.com/jpjanet/ML-chem-workshop](https://github.com/jpjanet/ML-chem-workshop)  
this revision: 795e5c5440fb08de2cf673d2742e989915a0e4e0 on branch master





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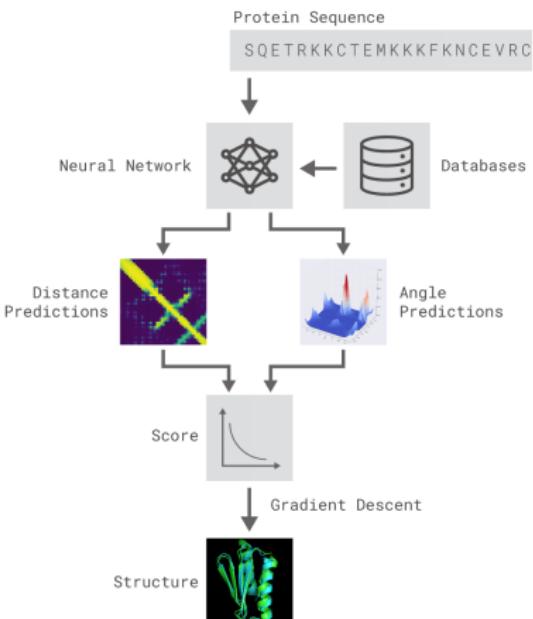
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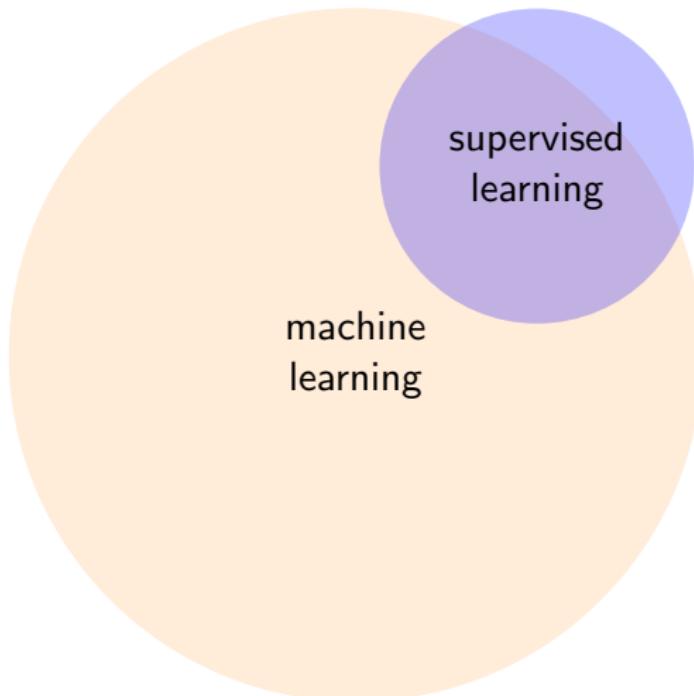
This is probably a bit strong, but all scientists generate data as a product. ML provides new, powerful ways to exploit their that information.

# Types of machine learning

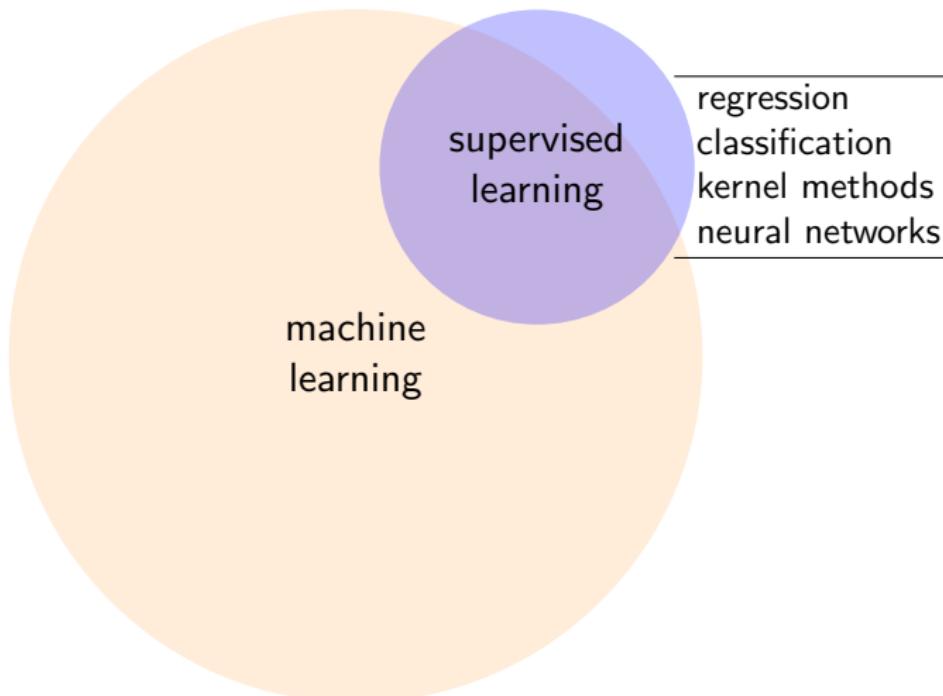
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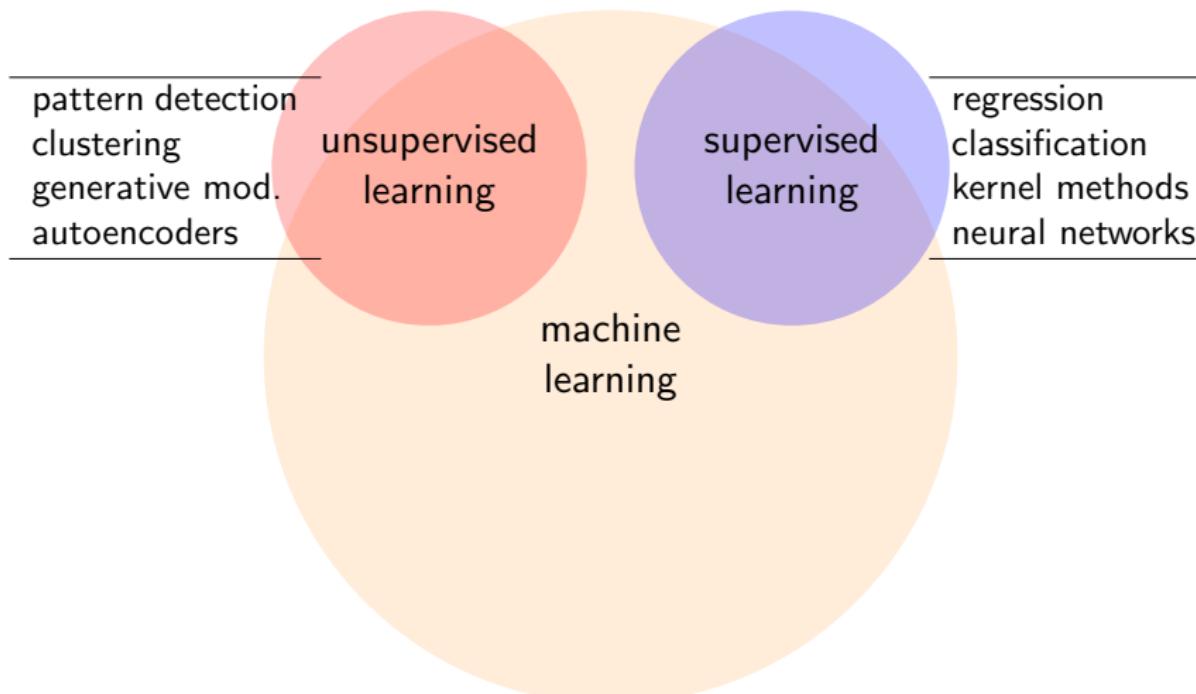
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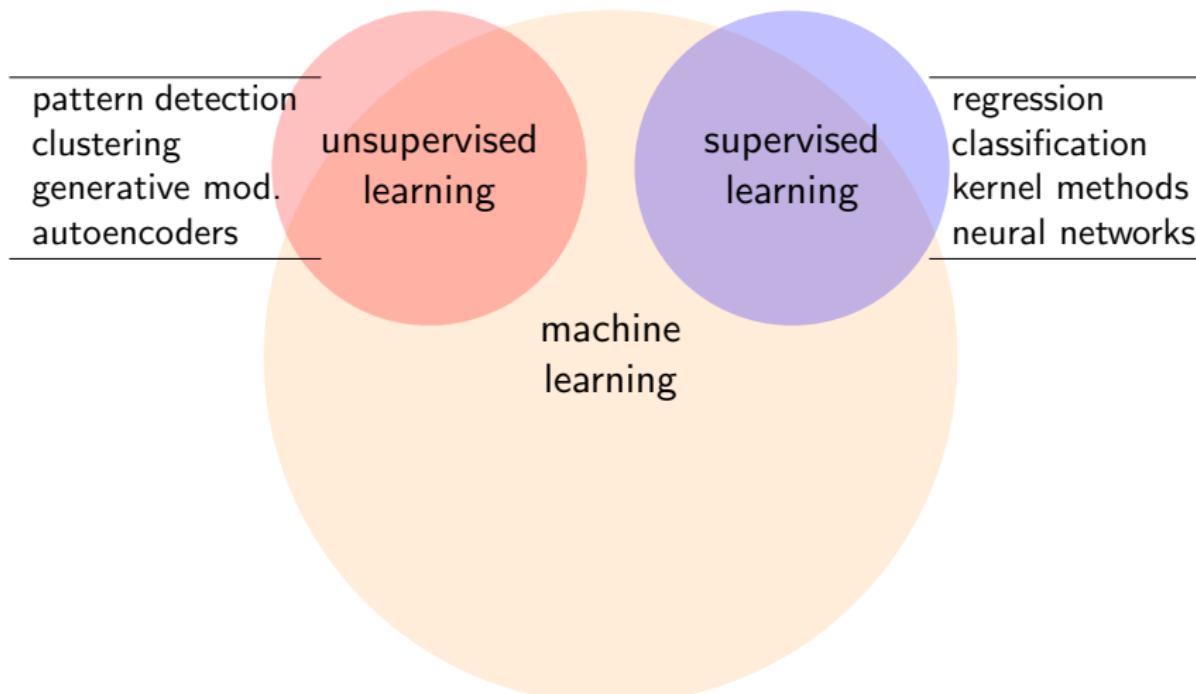
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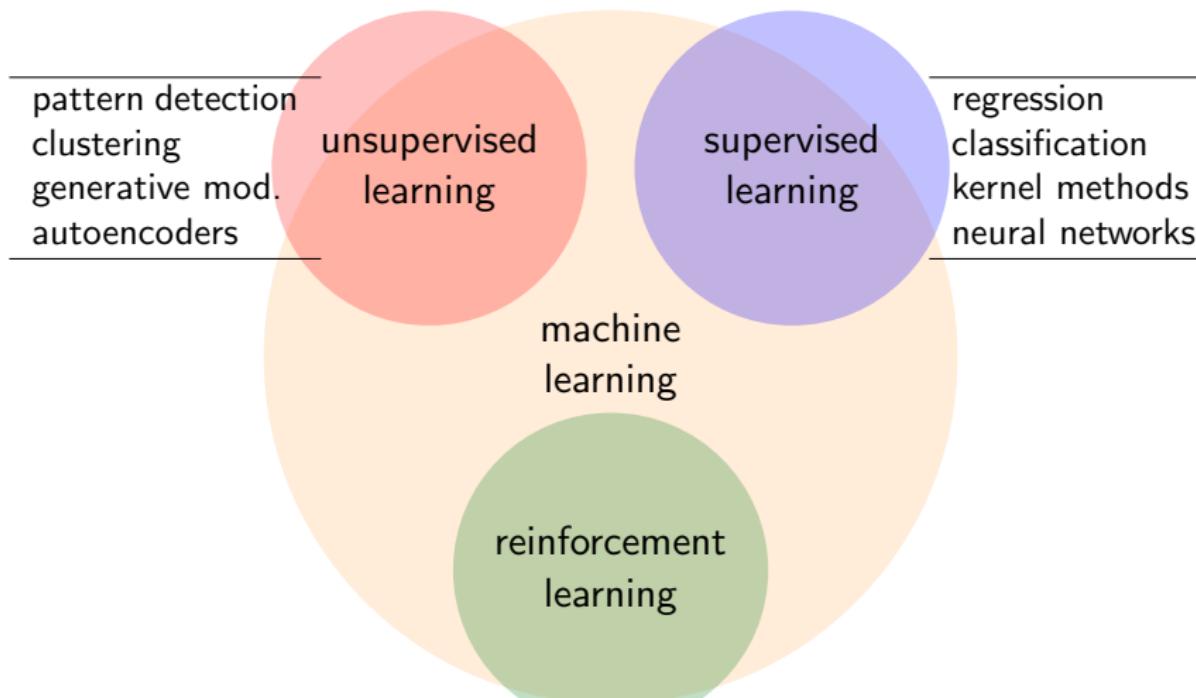
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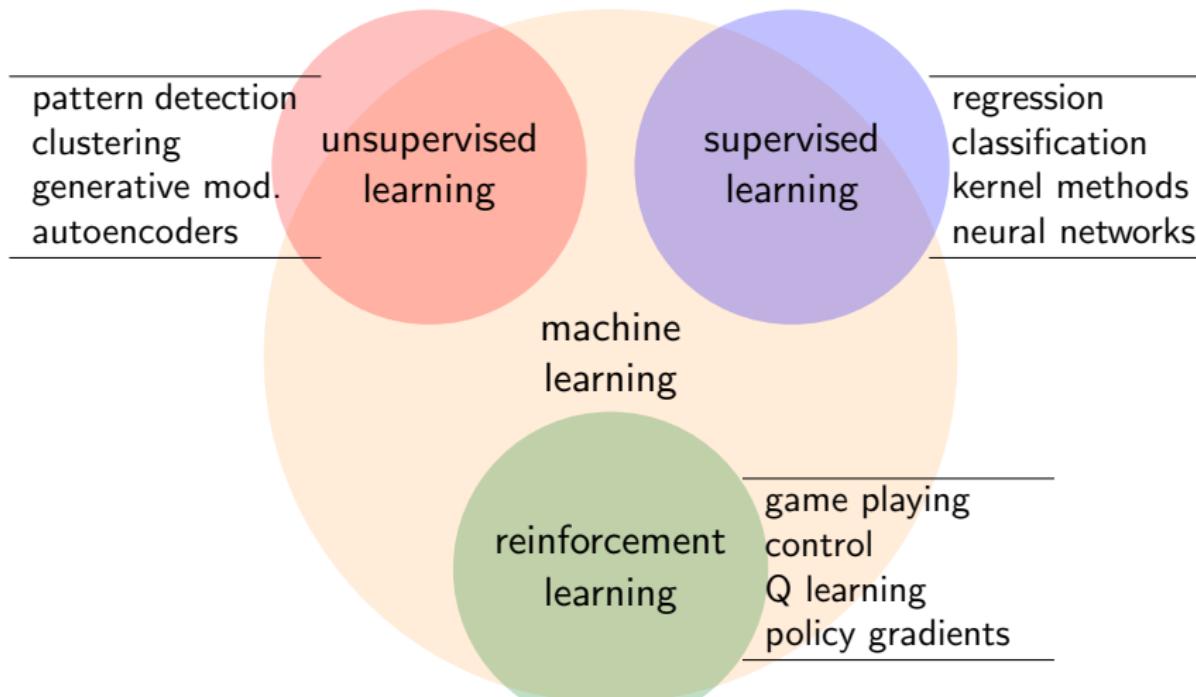
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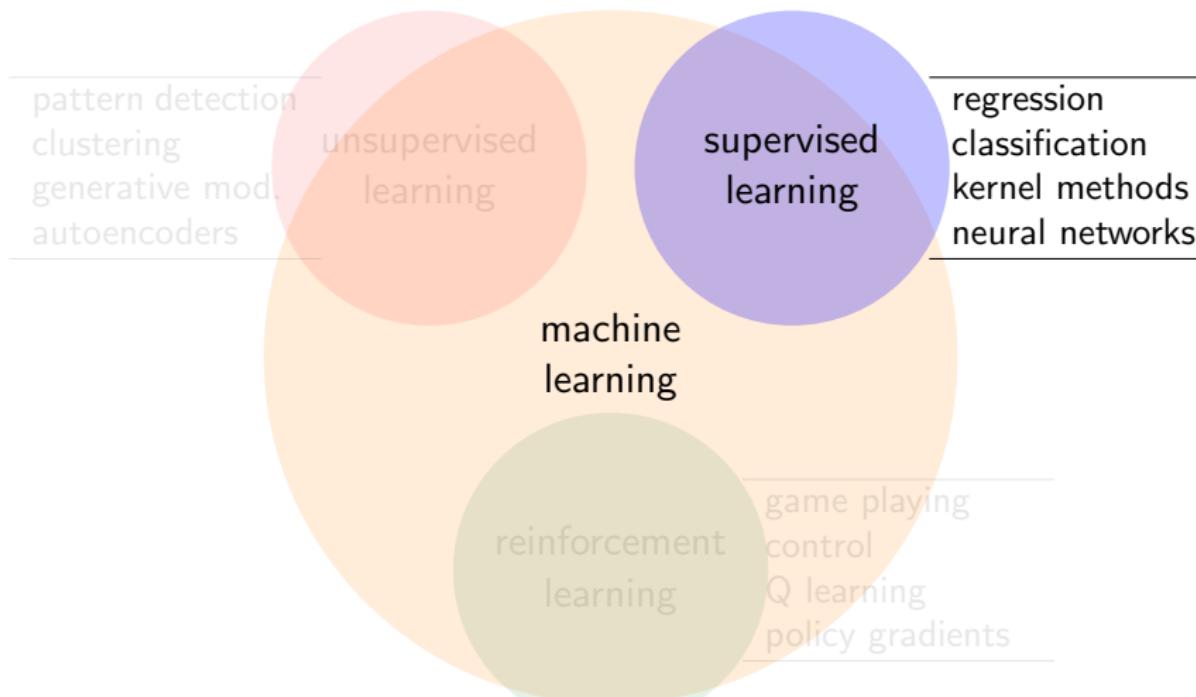
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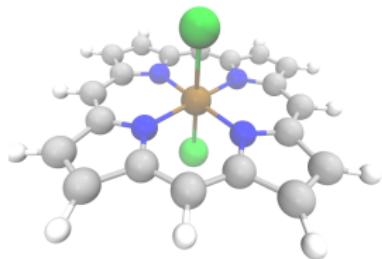
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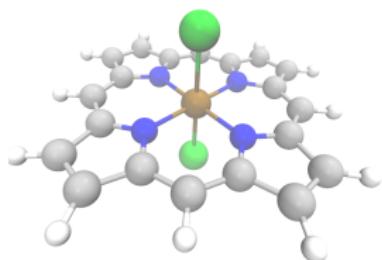
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## Why ML in chemistry?

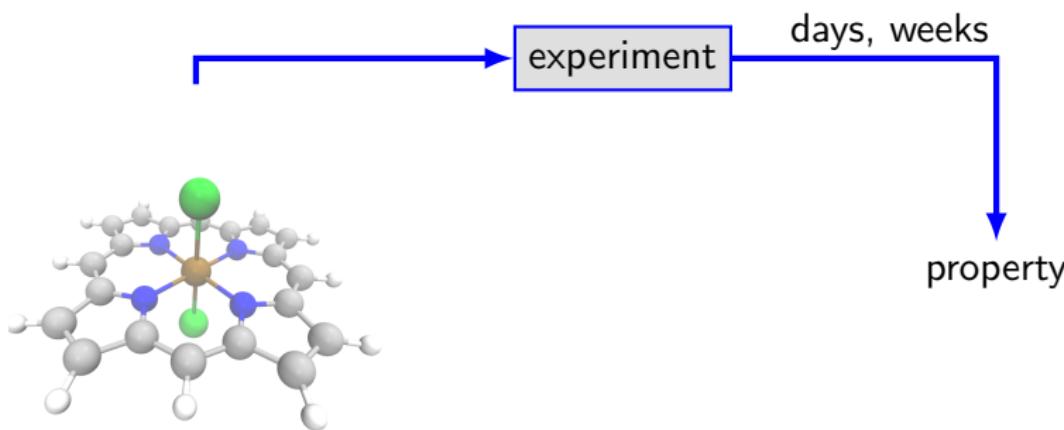


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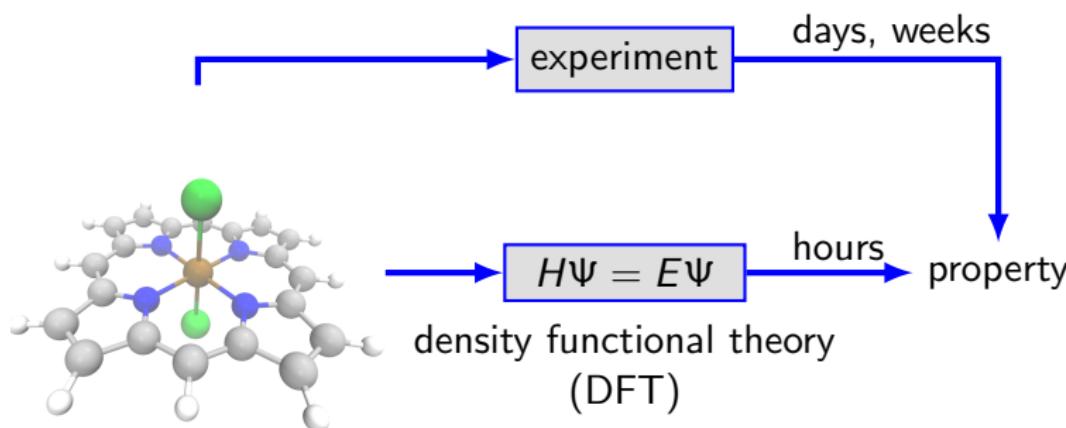


property

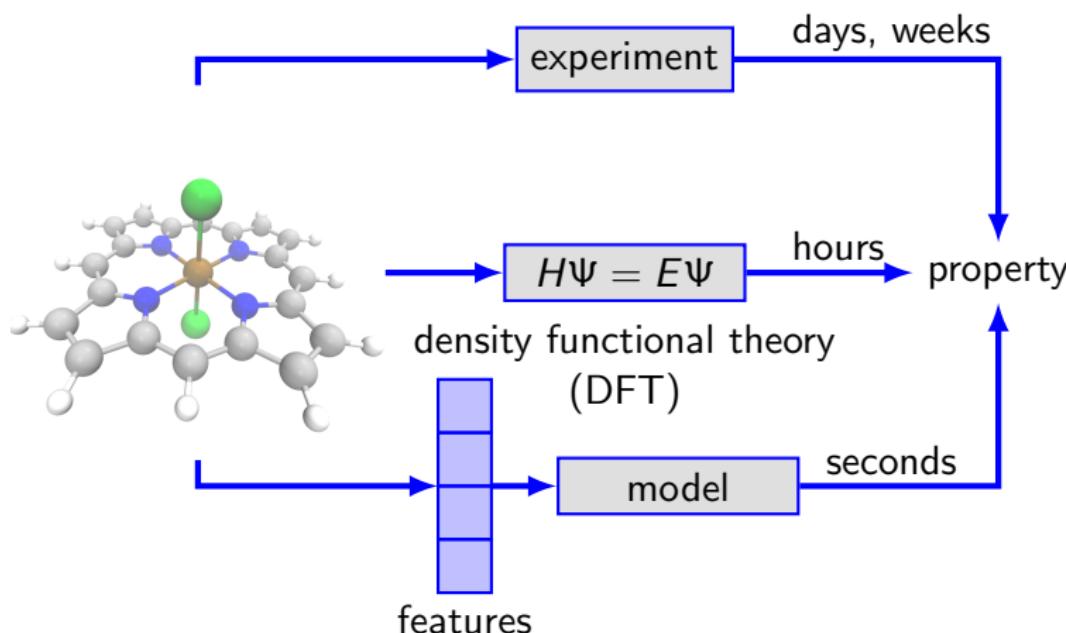
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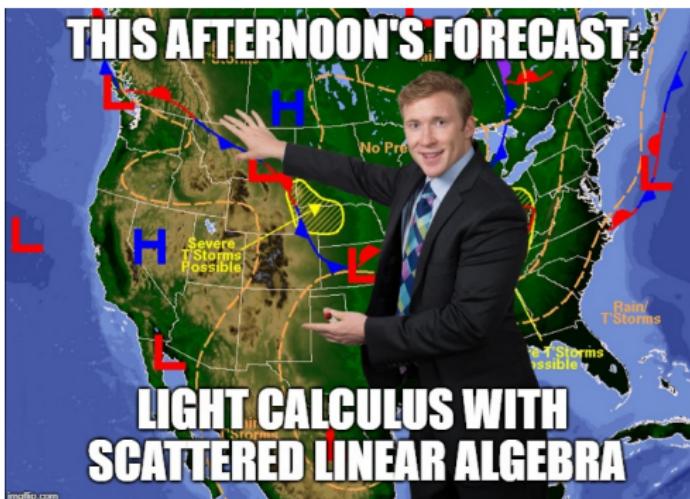
Please ask questions throughout!

# Disclaimer

Warning: this talk contains some *light* mathematics.

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Some useful notation:

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$X$	training data, as rows
$x^*$	one new molecule/systems
$y, \hat{y}$	property(energy?), predicted value
$\mathcal{L} = \ y - \hat{y}\ _2^2$	loss function
$W, w$	model parameters
$\hat{y} = f(x, W)$	our model

---

# The goal of statistical learning

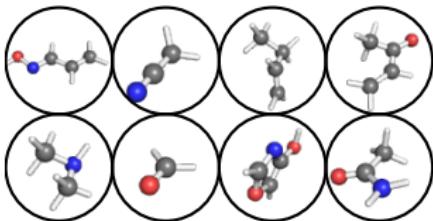
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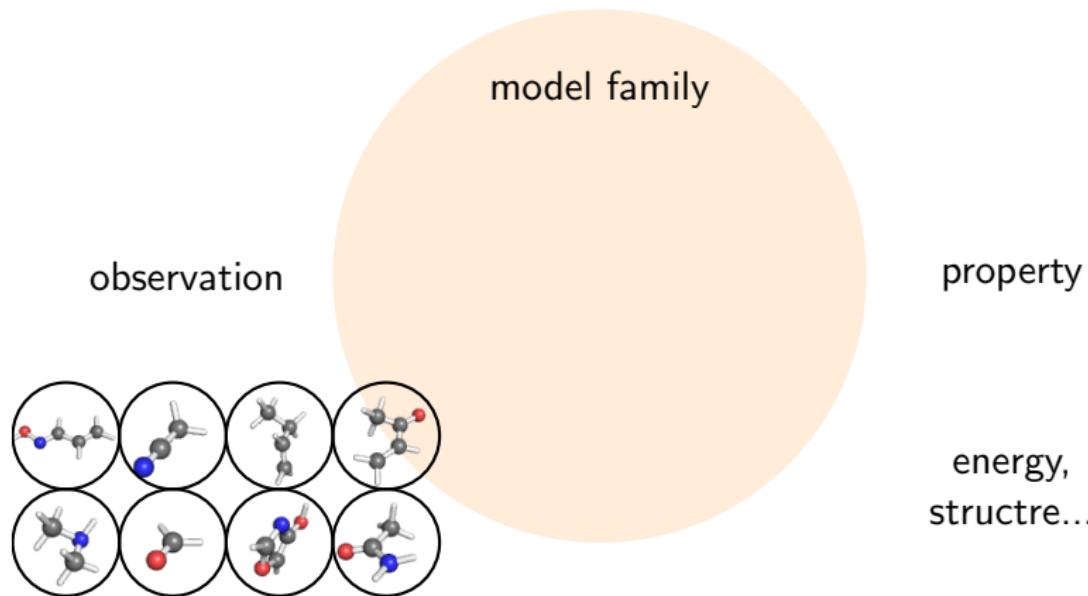
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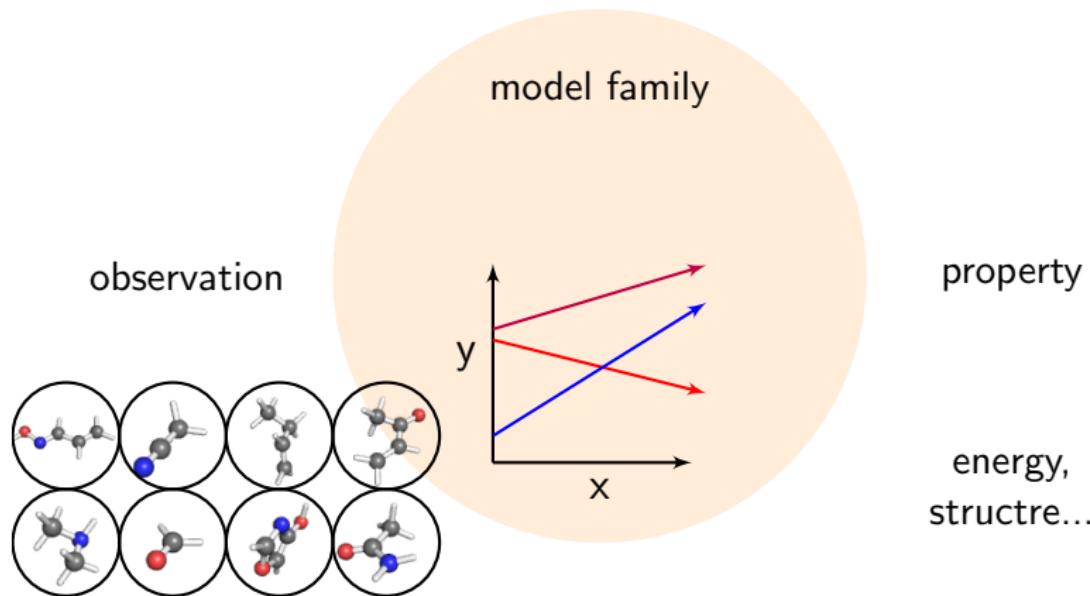
property

energy,  
structre...

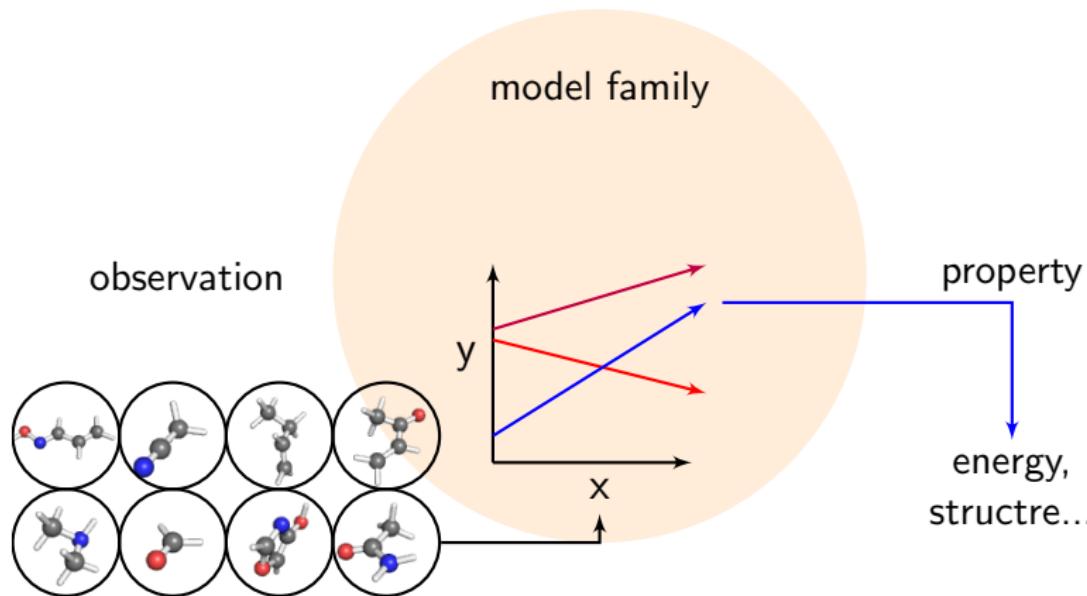
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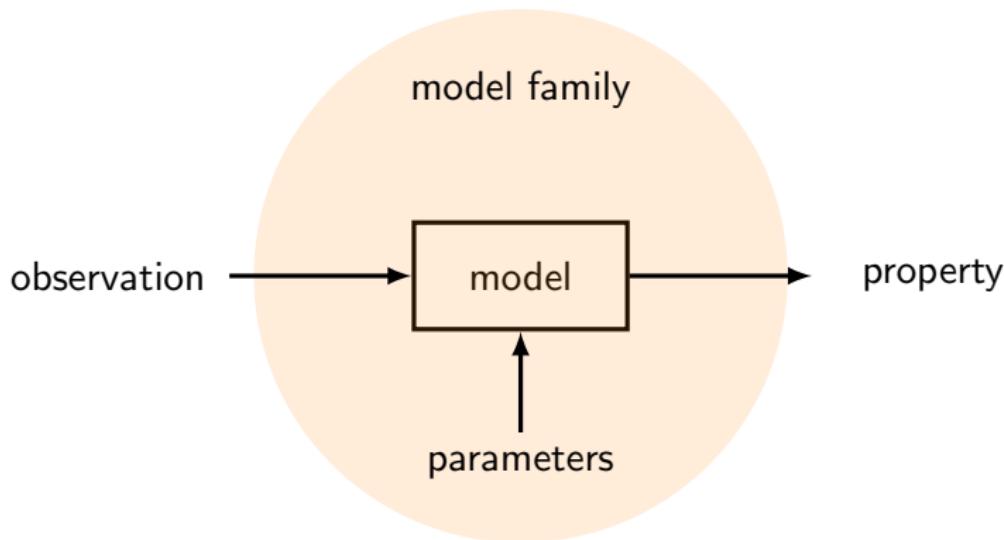
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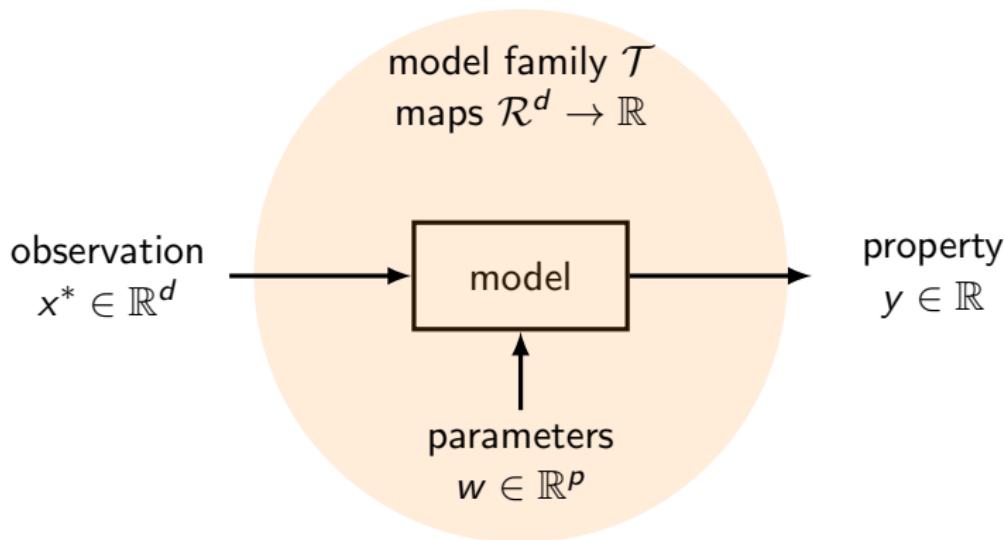
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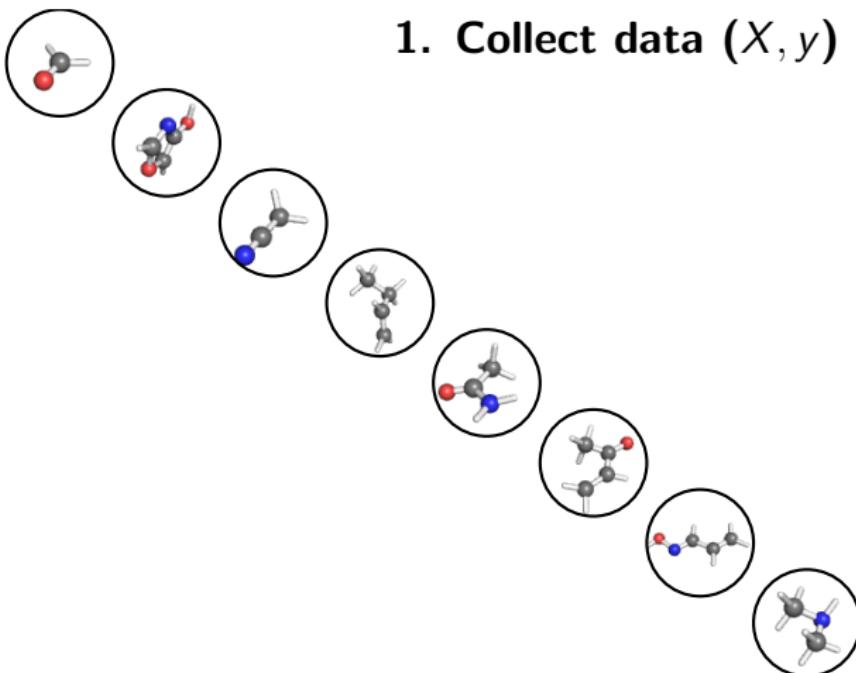


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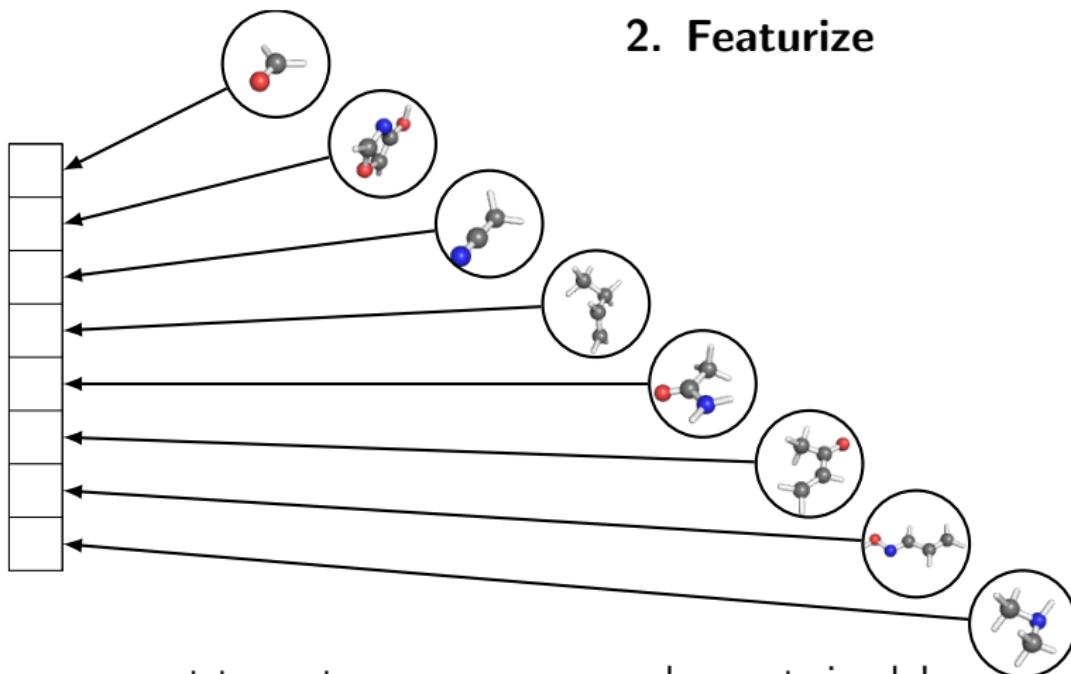
# Overview of supervised learning

1. Collect data ( $X, y$ )



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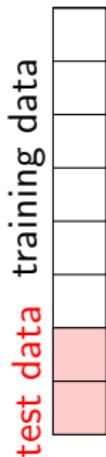
## 2. Featurize



convert to vectors, preprocess, scale – not simple!

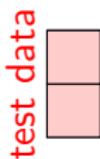
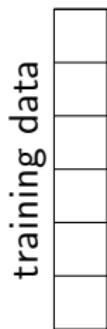
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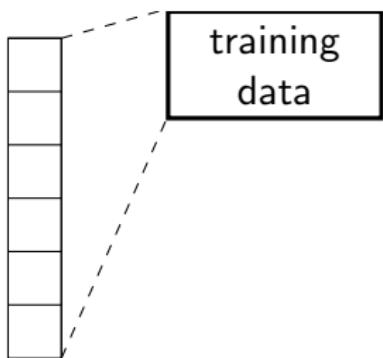


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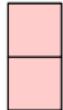
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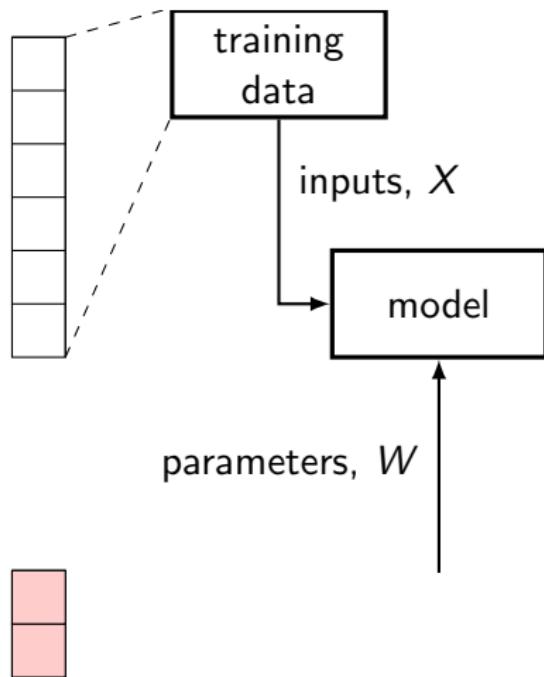
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## 4. Learning phase

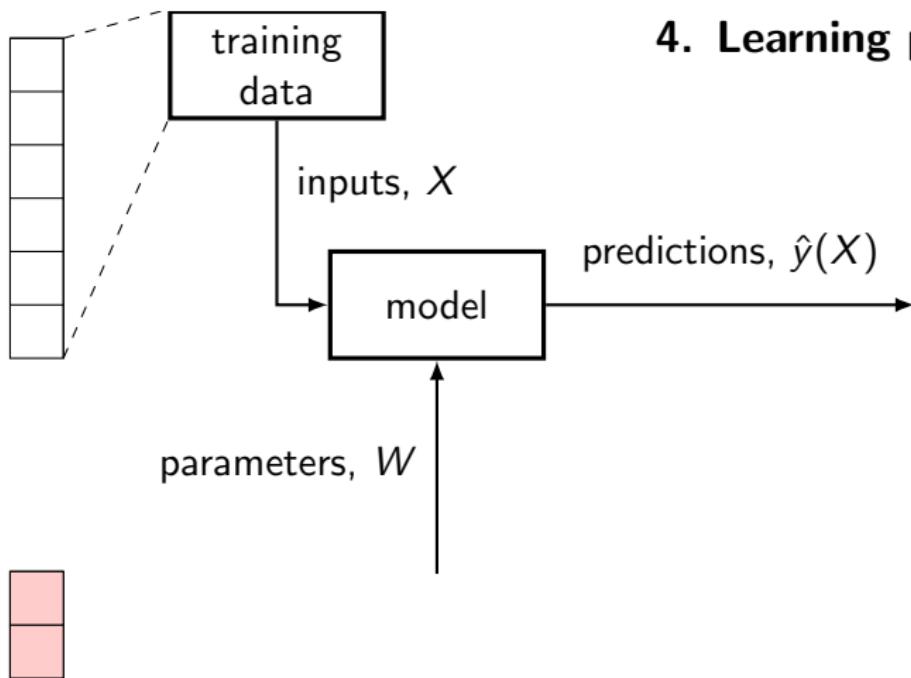


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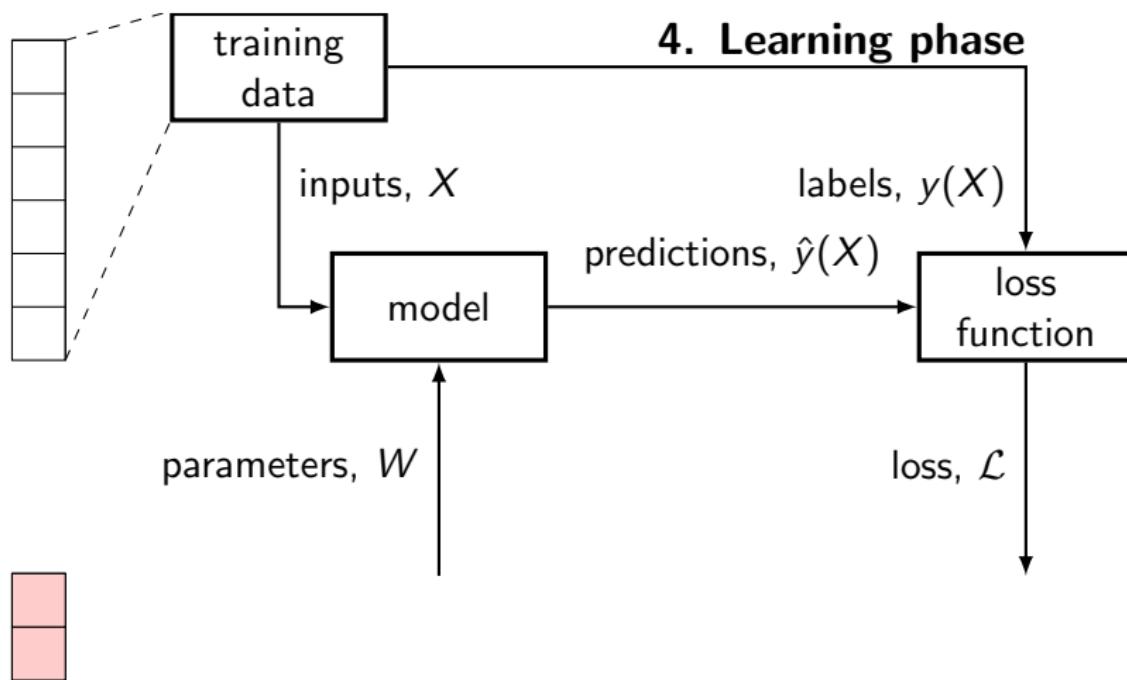
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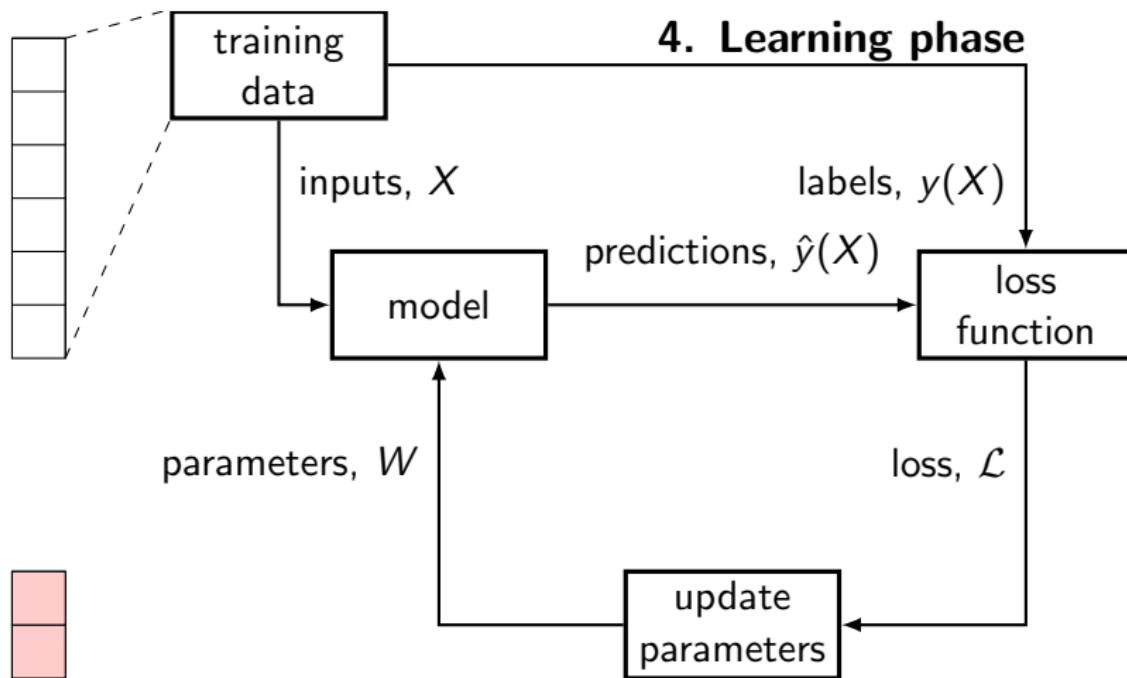


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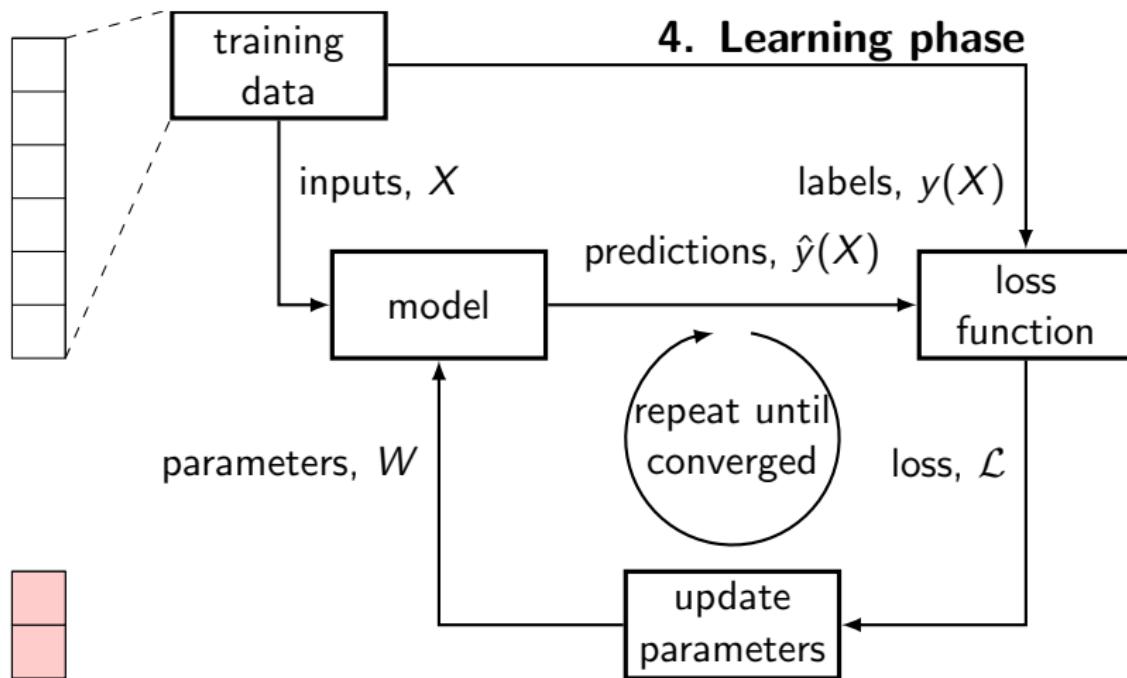
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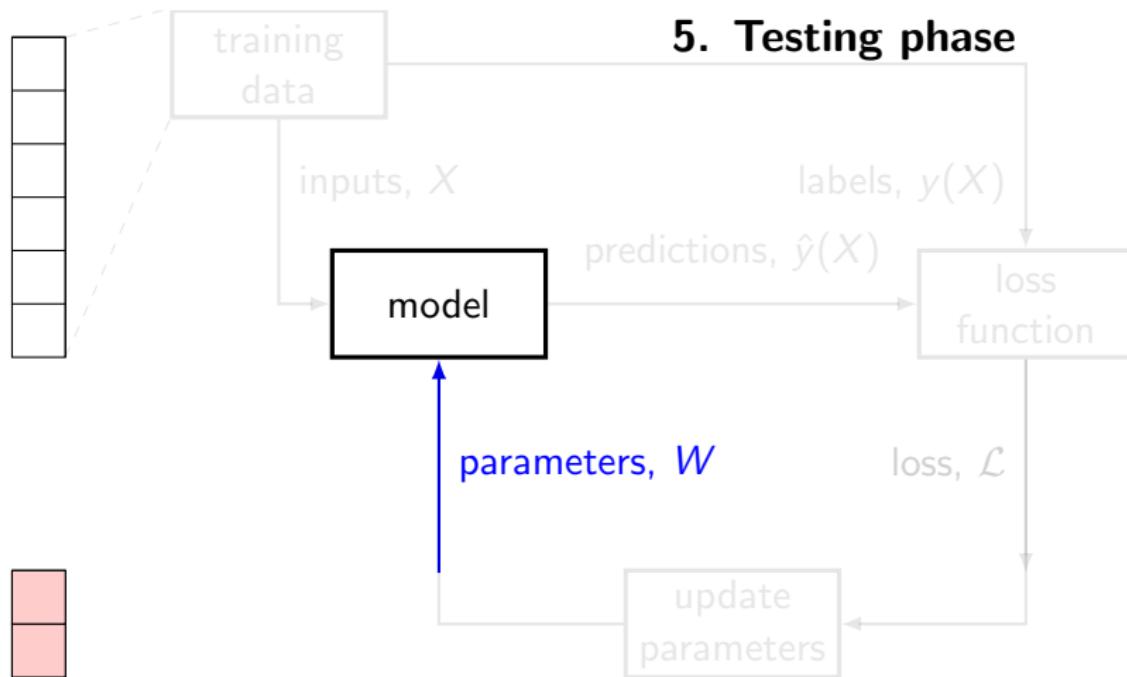
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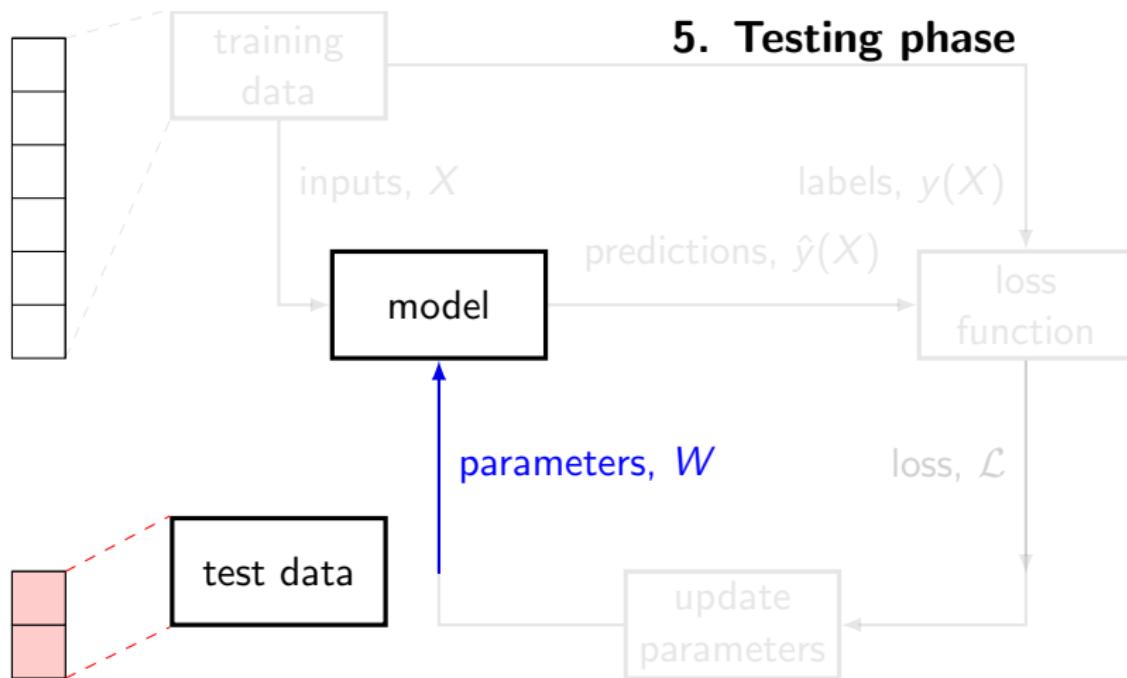
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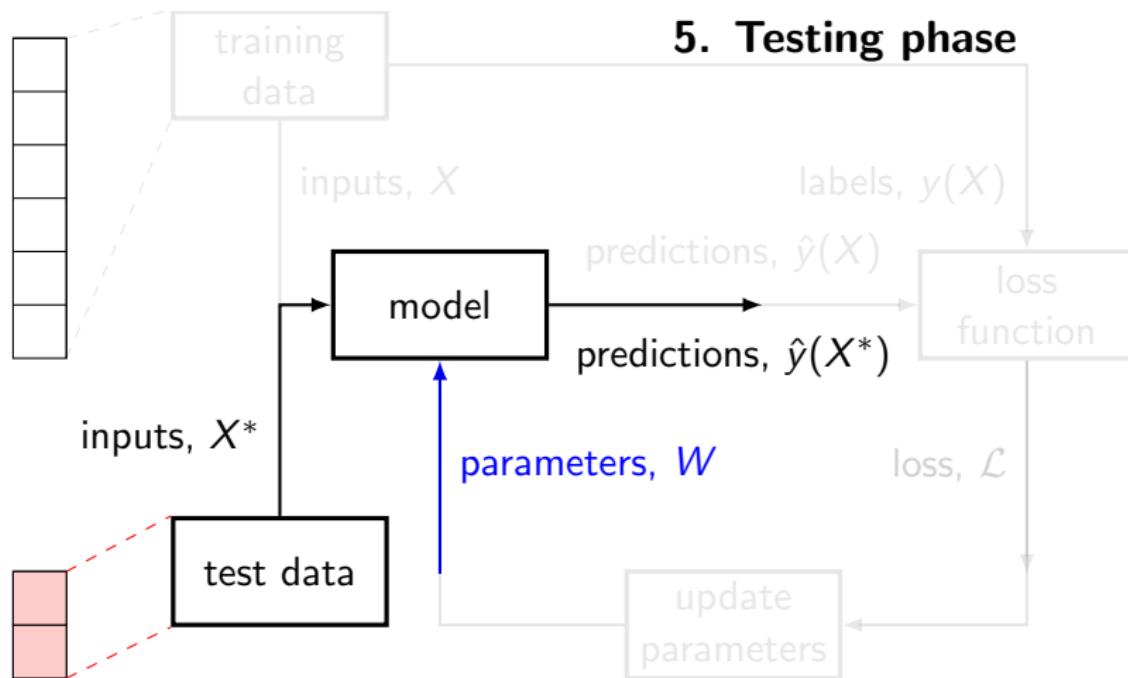
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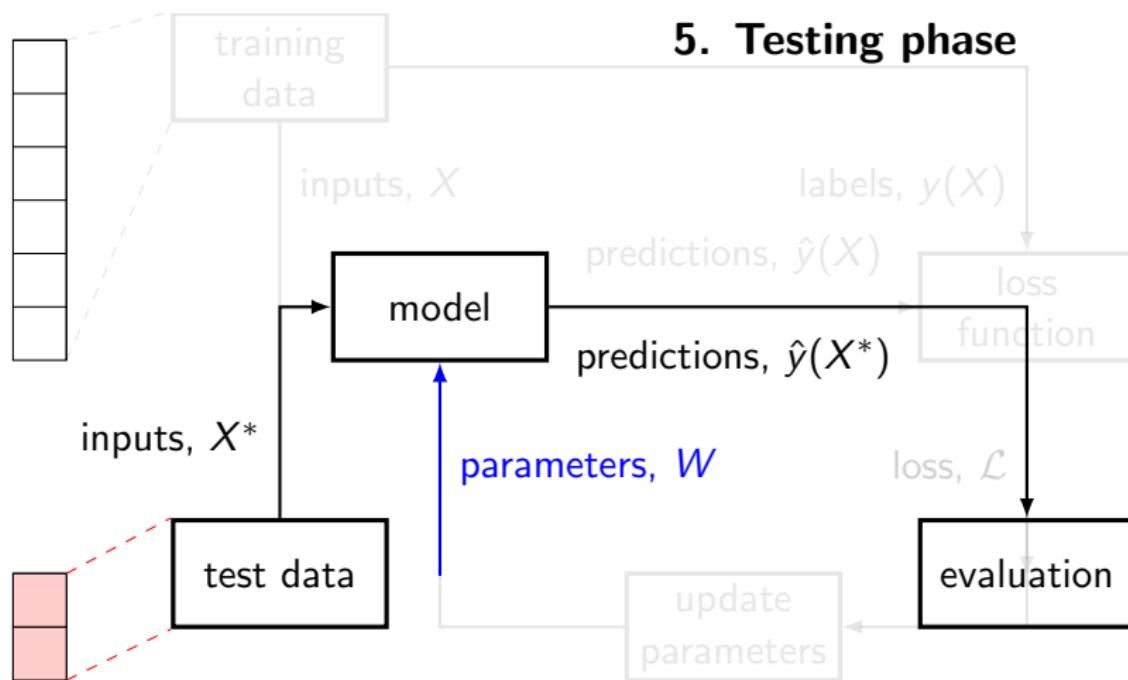
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## Risk and generalization - I

Our training data defines the *empirical risk*

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We want  $\mathcal{T}$  to be large/complicated enough to have low approximation error, **but no more complicated**.

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With limited data, we are often better off searching for a model in a simpler family models of that 'learn' more robustly and quickly as opposed to very complicated models with lots of parameters.

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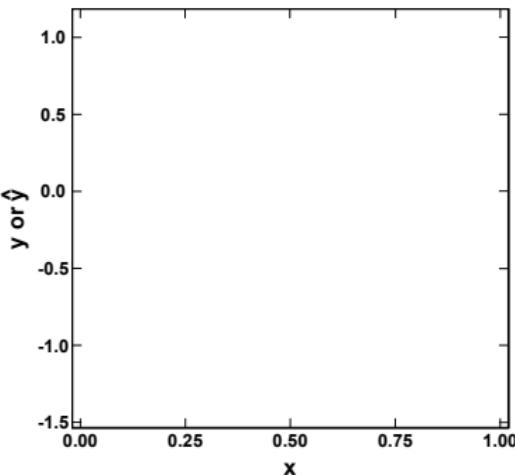
Conversely, a simple model will stop improving with more data past a certain point – where the approximation error dominates.

# Risk and generalization - IV

Let us use **polynomials** to estimate:

$$y(x) = \sin(2\pi x)$$

Note that  $f^* \notin \mathcal{T}$ !



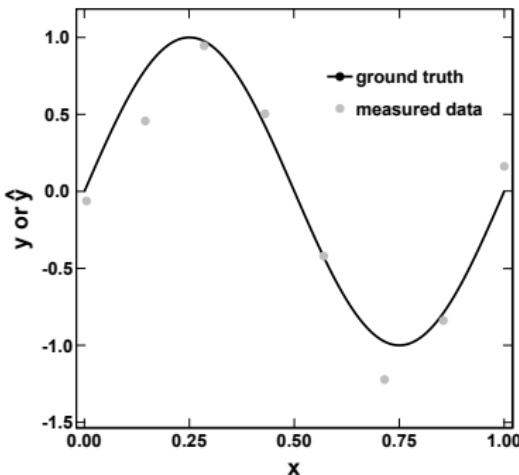
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Assume 8 measurements with noise  $\mathcal{N}(0, 0.2)$



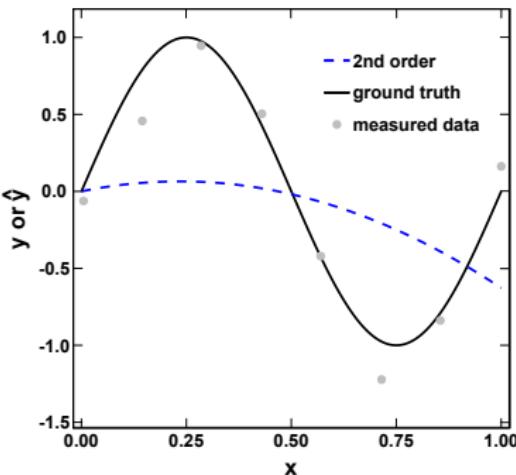
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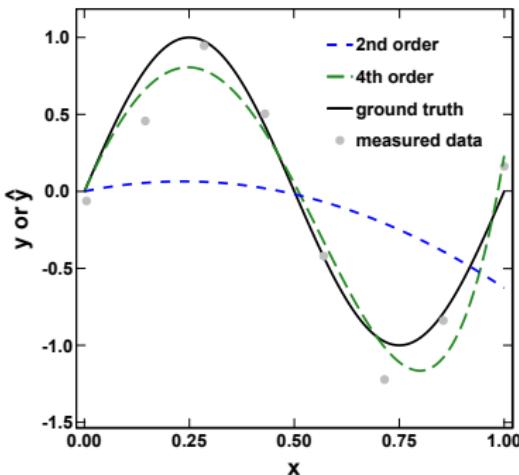
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Start with degree 2...What happens when we increase the order?

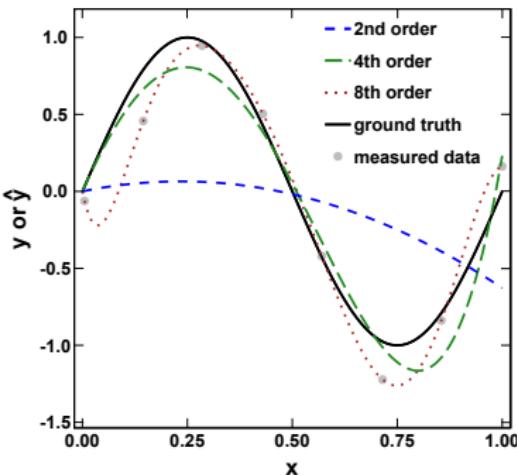


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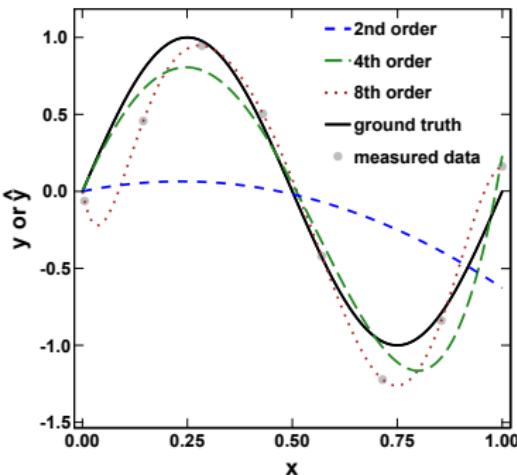
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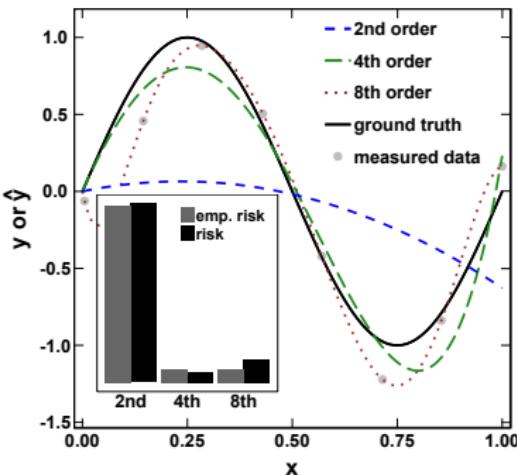
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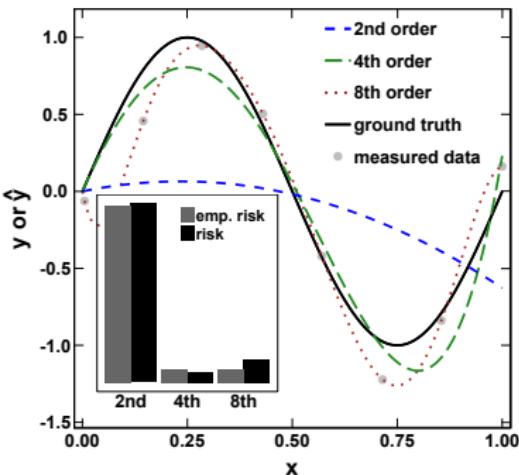


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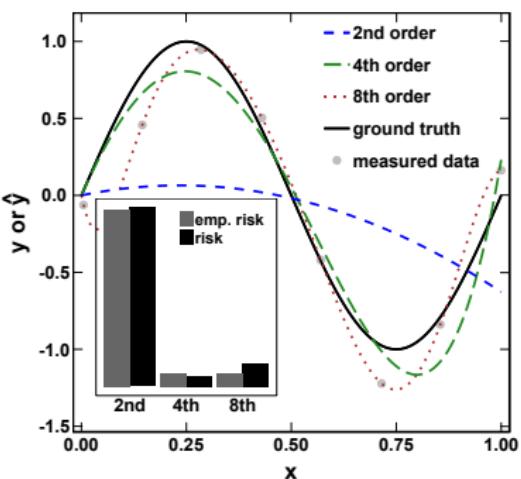
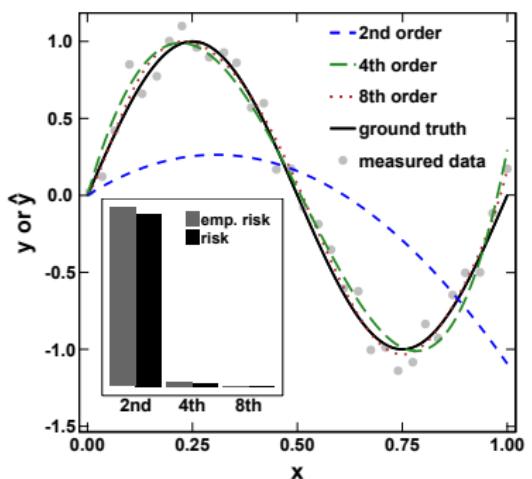
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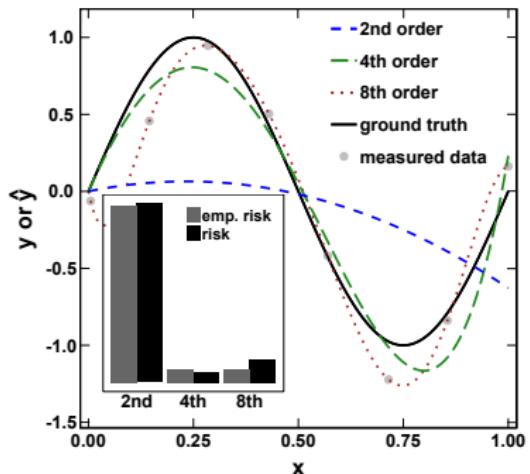
$$\begin{aligned}\mathcal{L}'(y, f(x, w)) &= \mathcal{L}(y, f(x, W)) + \lambda R(W) \\ (\text{Tiknohov/Ridge}) &= \frac{1}{n} \|y - f(x, W)\|_2^2 + \lambda \|W\|_2^2\end{aligned}$$

We choose to **penalize terms with large weights**, i.e. complicated functions.

- This makes empirical errors worse.
- This *can* improve generalization/excess risk.

# Controlling Complexity

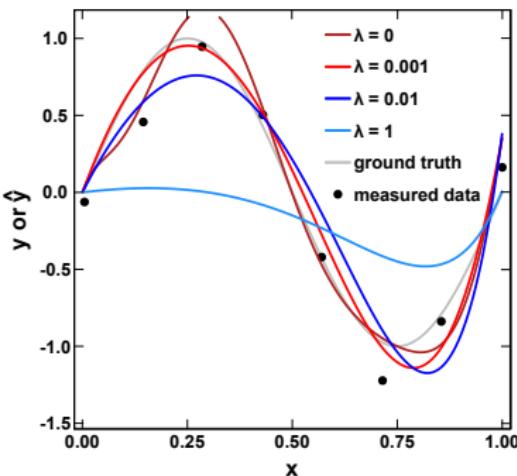
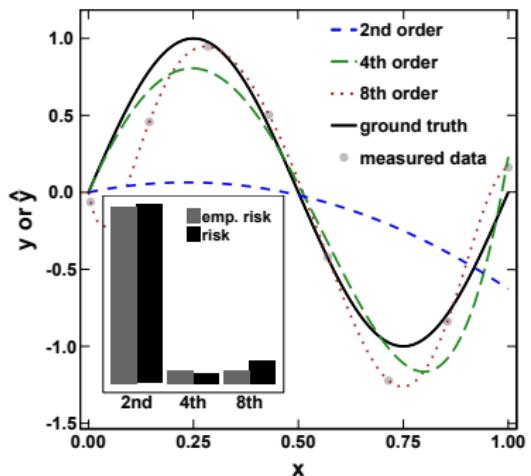
Let's return to our previous example:



Remember: larger  $\lambda$  = simpler, flatter model.

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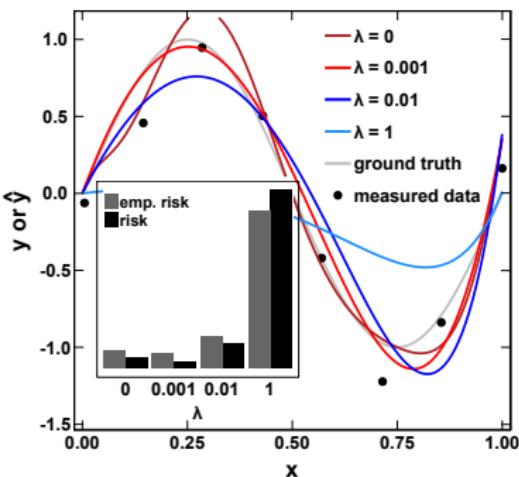
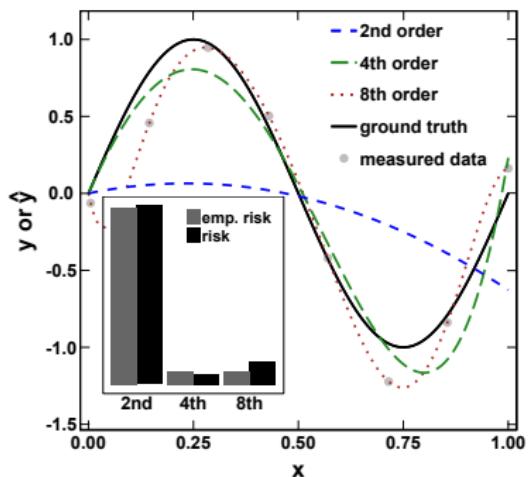
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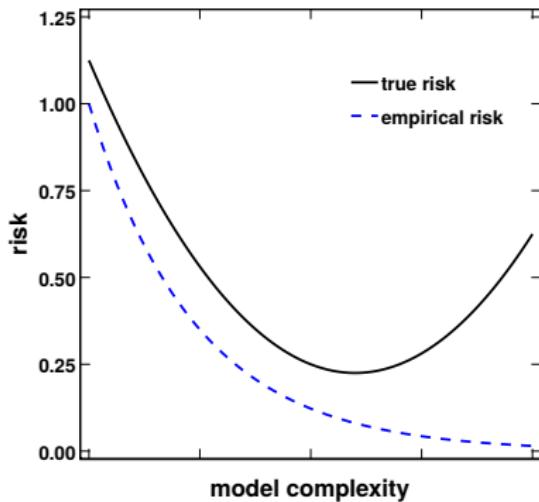
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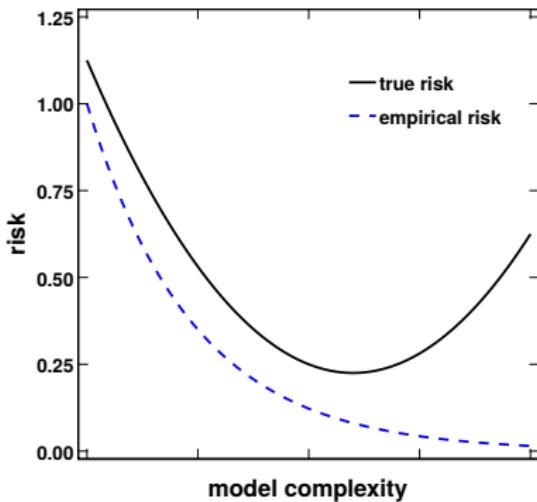


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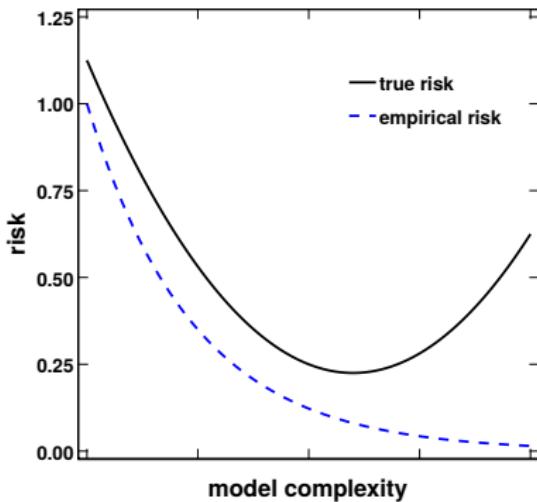


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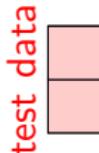
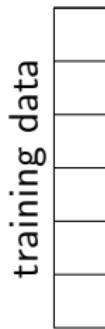
None of this helps us understand how complicated our model should be.

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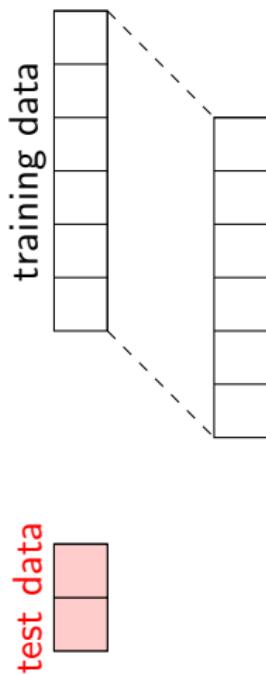


None of this helps us understand how complicated our model should be. Unfortunately, **errors on our training data cannot tell us the answer**

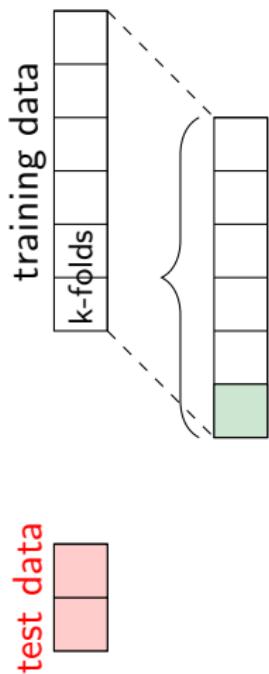
# (Cross)-validation



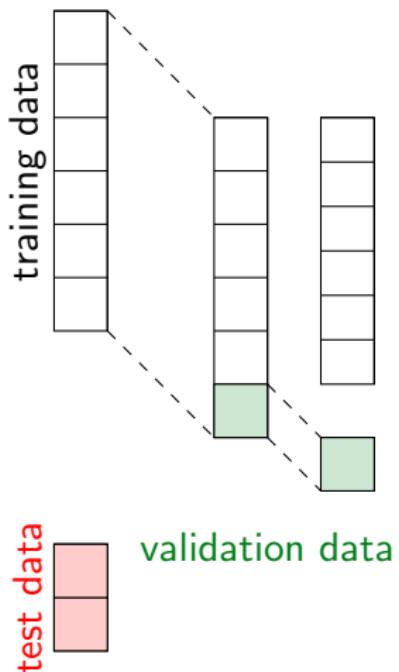
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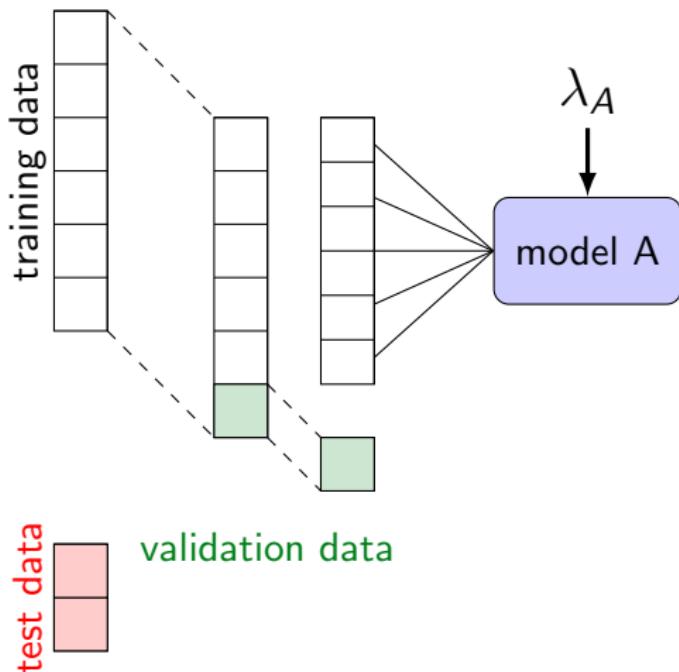
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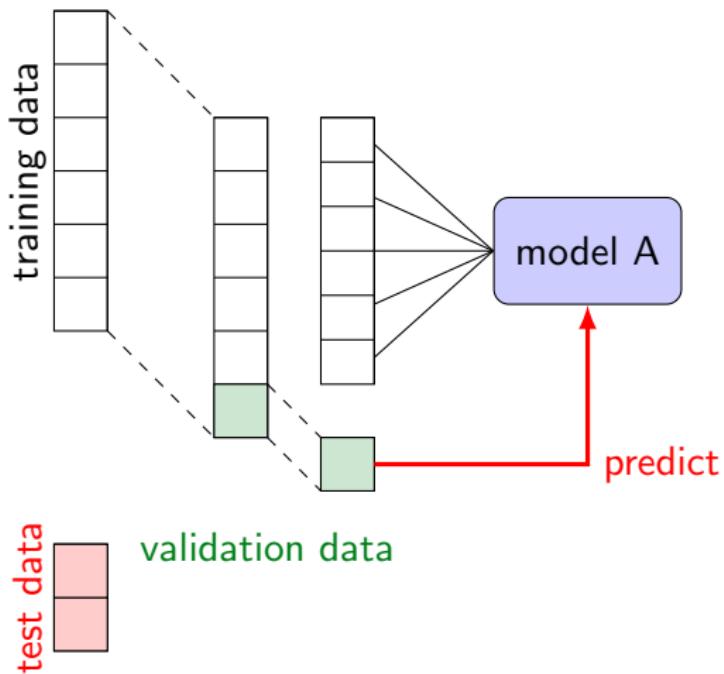
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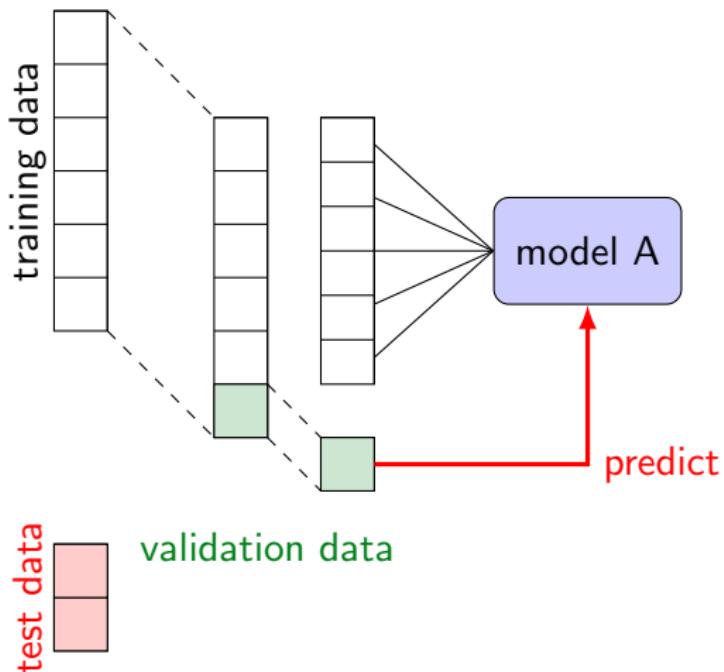
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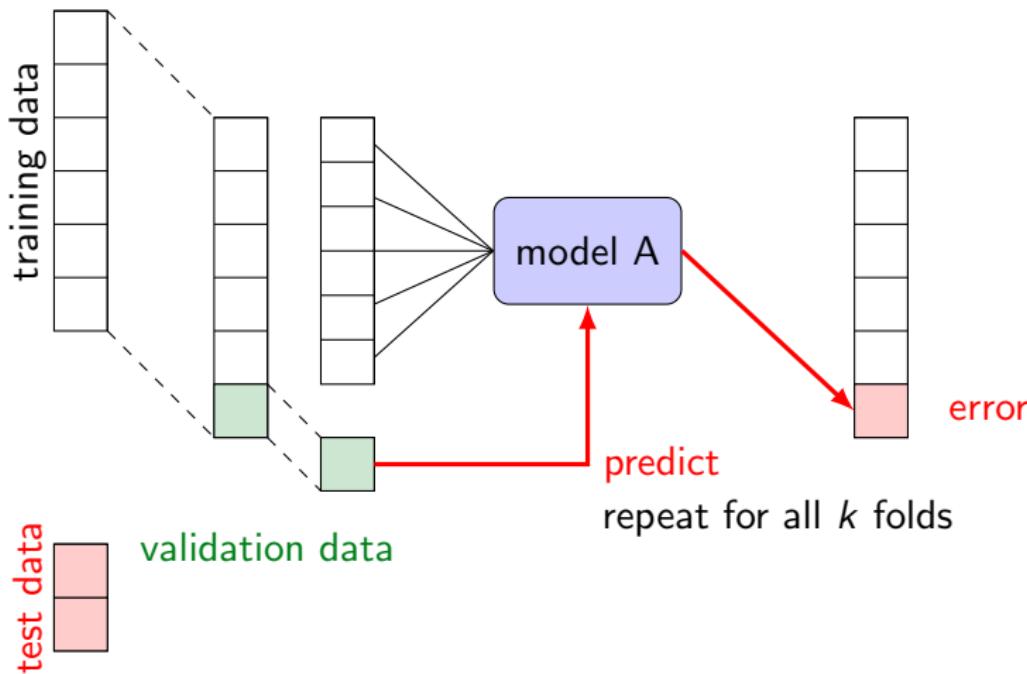
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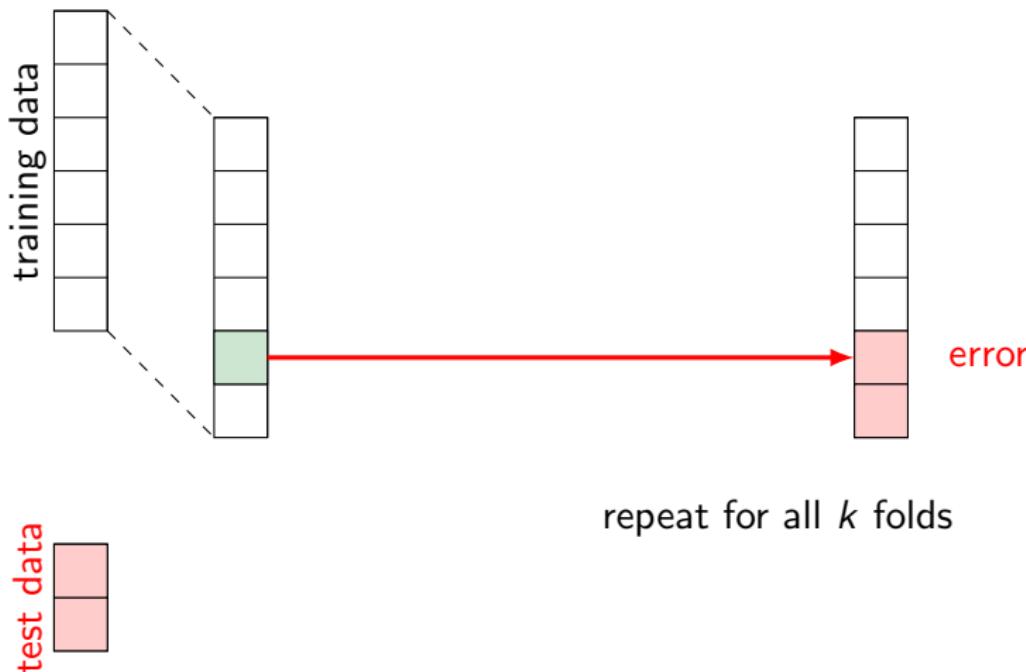
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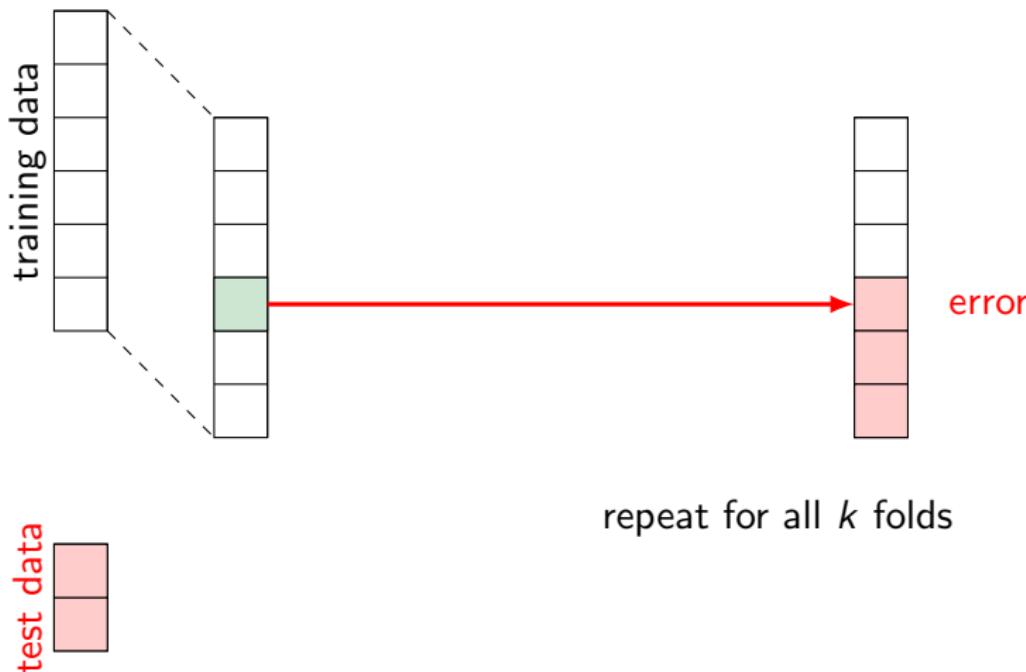
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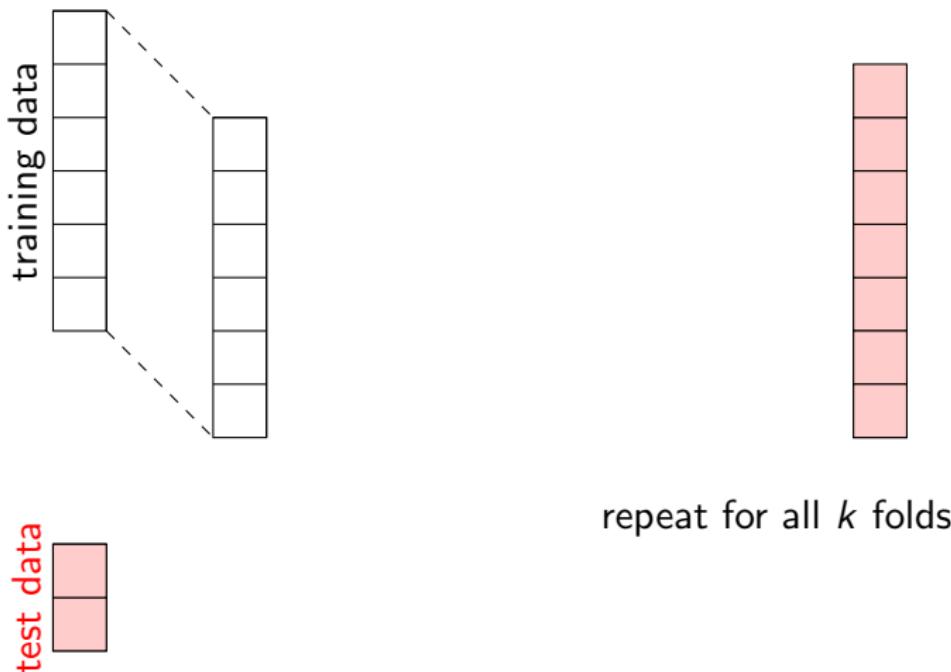
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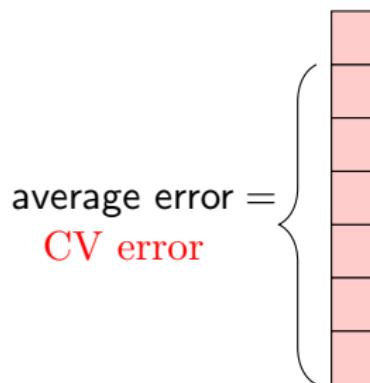
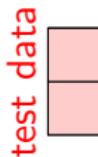
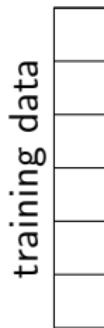
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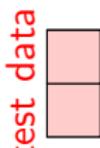
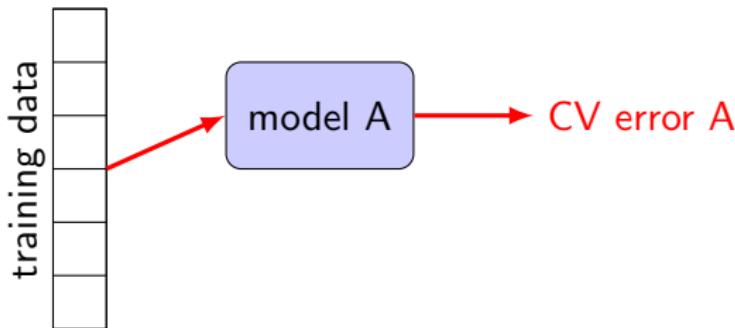
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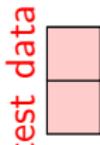
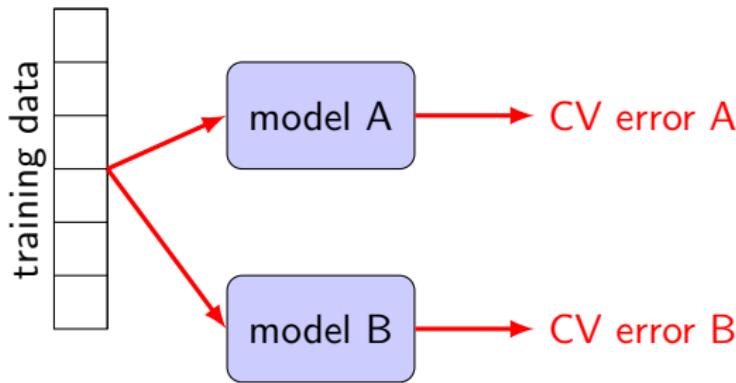


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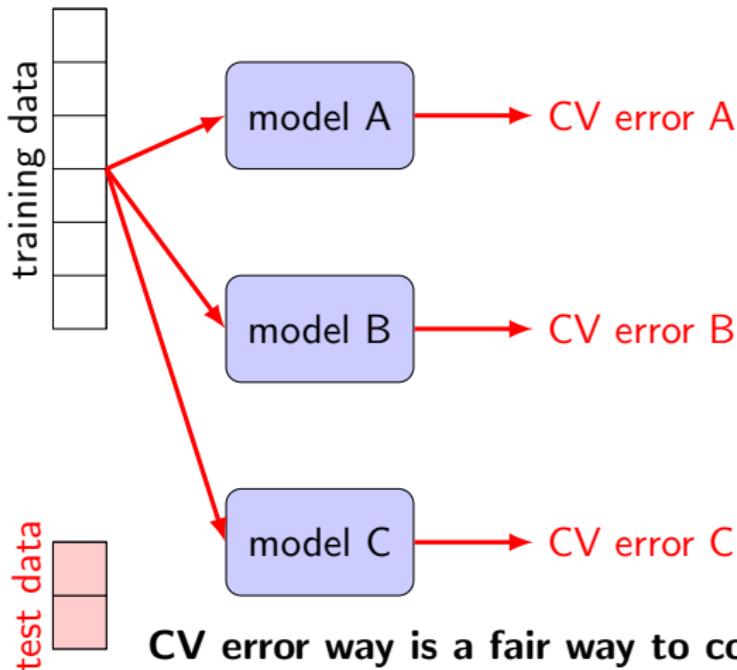
**CV error way is a fair way to compare models**

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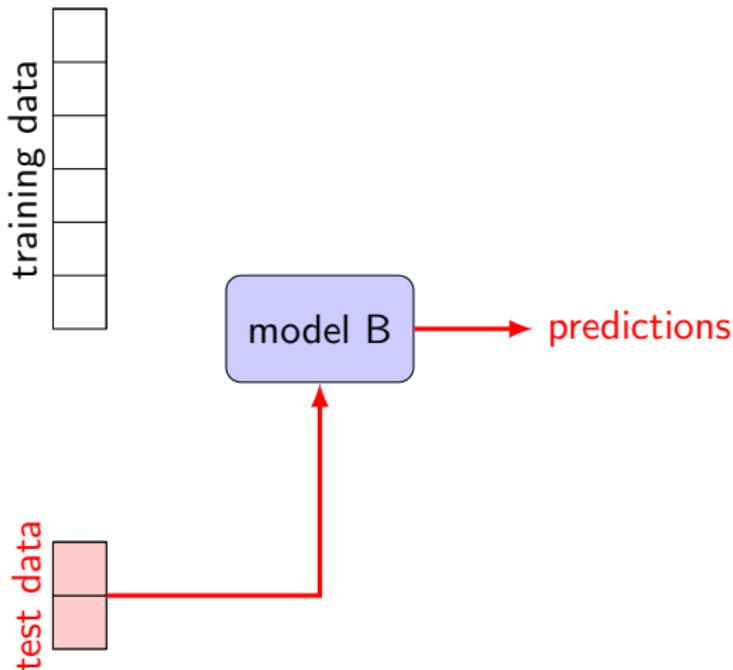


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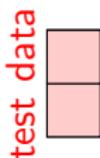
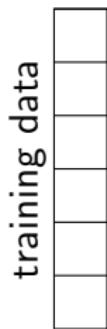
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Deep neural networks (might) need better theories.

# Multiple linear regression



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We can write this in a matrix form as well:

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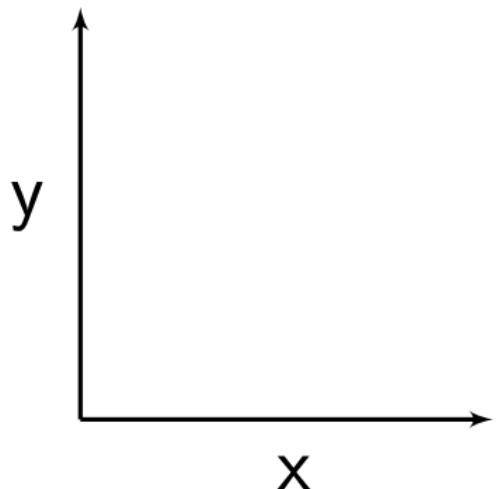
Notice how we handle the constant terms

## Multiple linear regression II

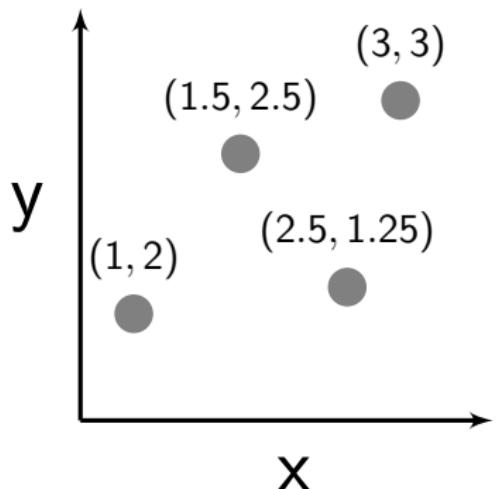
Let's solve our regularized least-squares problem:

$$\begin{aligned}
 w &= \arg \min_{w \in \mathbb{R}^p} \frac{1}{n} \|y_{data} - Xw\|_2^2 + \lambda \|w\|_2^2 \\
 &= \frac{1}{n} (y_{data} - Xw)^T (y_{data} - Xw) + \lambda w^T w \\
 \frac{\partial \mathcal{L}}{\partial w} &= -\frac{2}{n} X^T (y_{data} - Xw) + 2\lambda w = 0 \\
 \implies (\lambda I + X^T X)w &= X^T y_{data} \\
 w &= (\lambda I + X^T X)^{-1} X^T y_{data}
 \end{aligned}$$

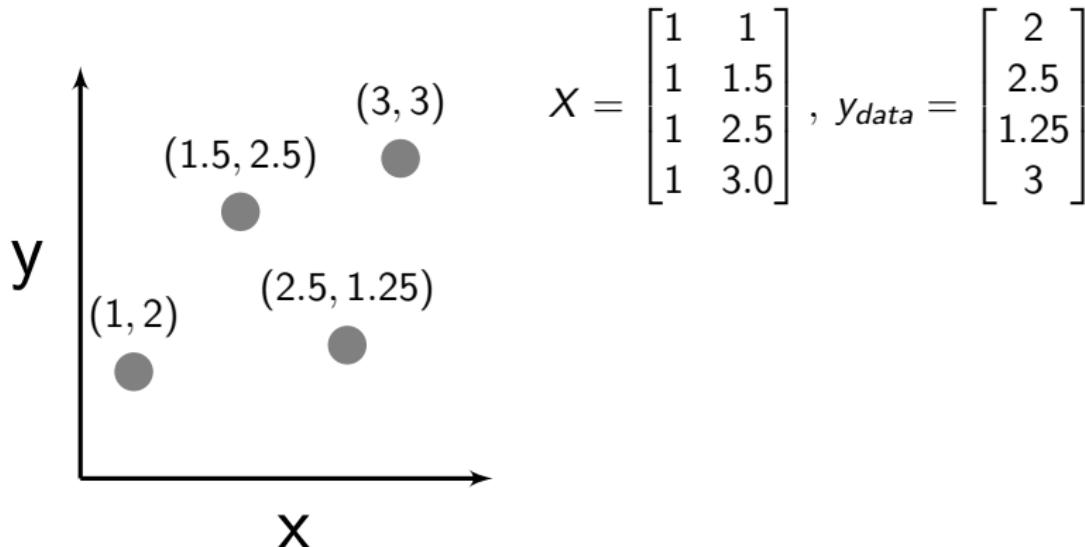
# Simple example in 1D



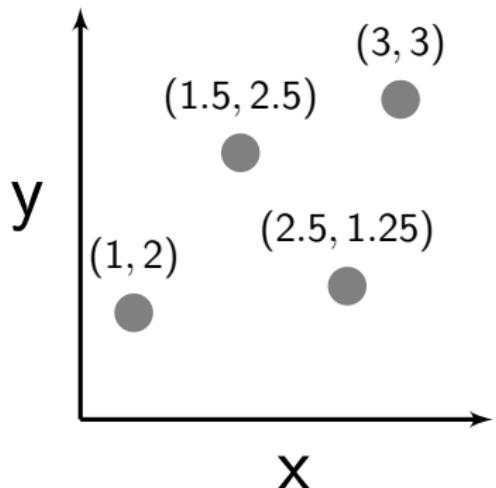
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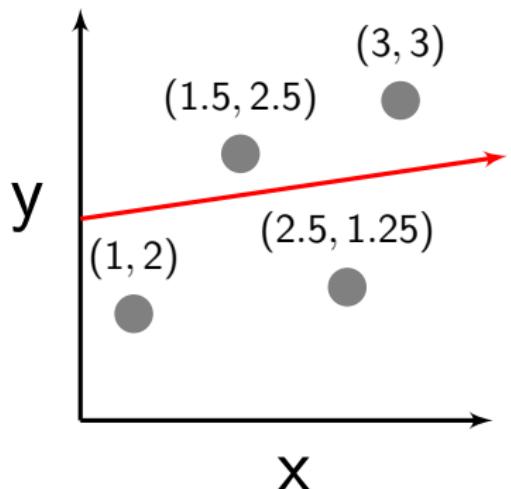
## Simple example in 1D



$$X = \begin{bmatrix} 1 & 1 \\ 1 & 1.5 \\ 1 & 2.5 \\ 1 & 3.0 \end{bmatrix}, y_{data} = \begin{bmatrix} 2 \\ 2.5 \\ 1.25 \\ 3 \end{bmatrix}$$

$$w = (X^T X + \lambda I)^{-1} X^T y_{data}$$

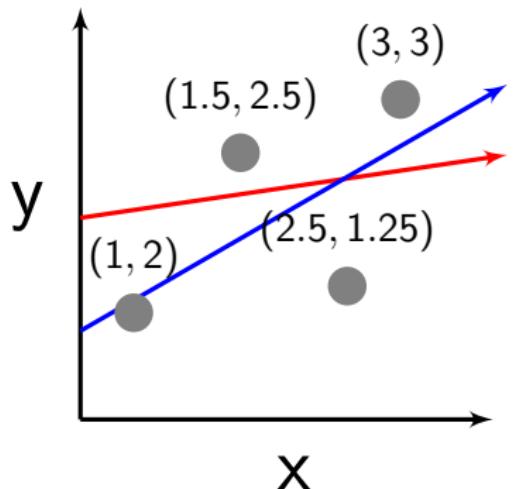
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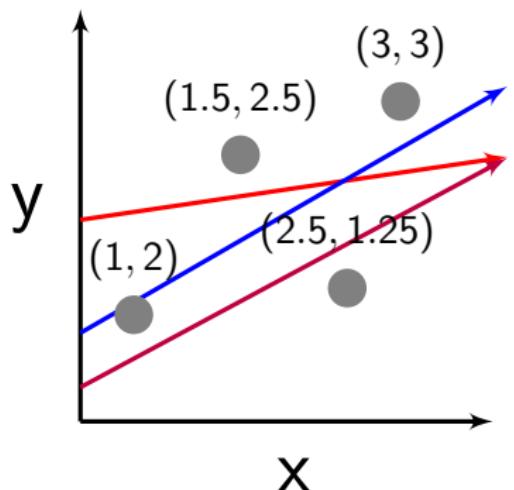
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$$\begin{aligned} w &= (X^T X + \lambda I)^{-1} X^T y_{data} \\ &= \begin{bmatrix} 0.82 \\ 0.58 \end{bmatrix} (\lambda = 1.0) \end{aligned}$$

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## The linear kernel

We can rewrite our result to express  $w = X^T a$  for  $a \in \mathbb{R}^n$  (shift of basis).

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The term  $k(x^*, x_i) = x^* x_i^T = \langle x^*, x_i \rangle$  is the **linear kernel**.

## The linear kernel II

The matrix  $K_{i,j} = \langle x_i, x_j \rangle$  is called the (linear) **kernel matrix**.

We can write the solution of the regression problem in this form – it is **exactly equivalent**:

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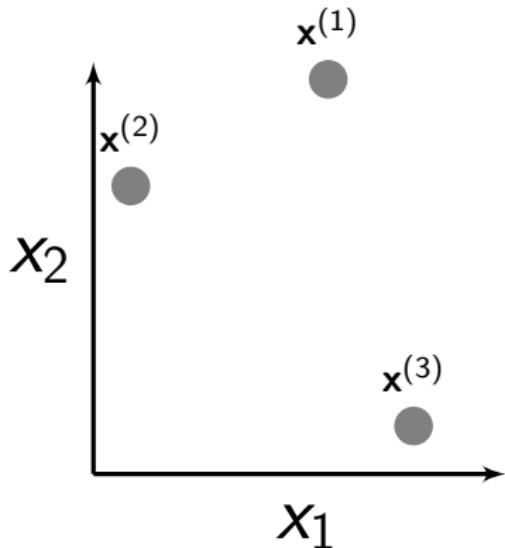
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The prediction at any new point is proportional to the inner product of each training point and the new point:

$$\hat{y}_{MLR}(x^*) = \sum_{i=1}^n k(x^*, x_i) a_i = \sum_{i=1}^n k \langle x^*, x_i \rangle a_i$$

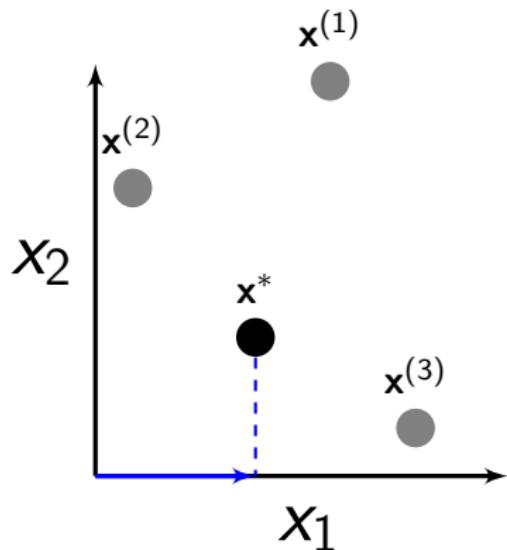
# Linear kernel geometric picture

$$y(x^*) = w_1 x_1^* + w_2 x_2^*$$



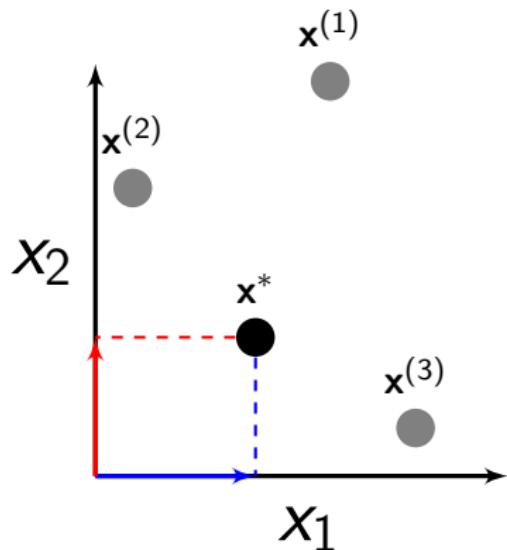
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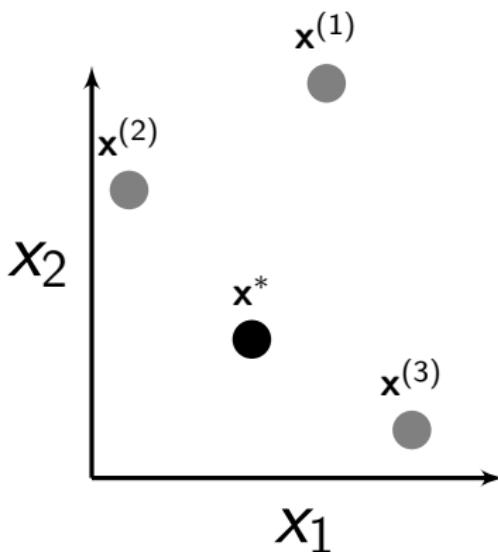
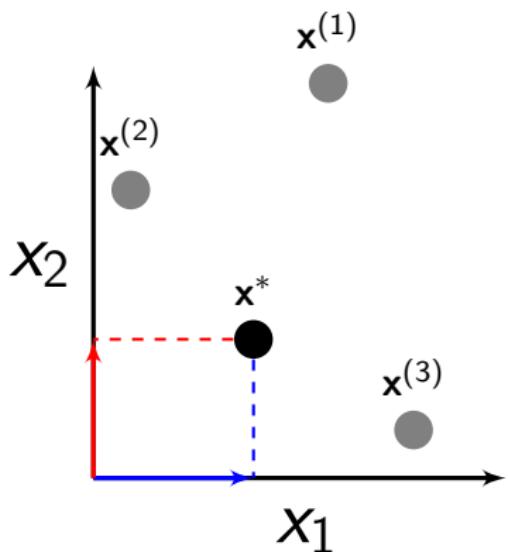
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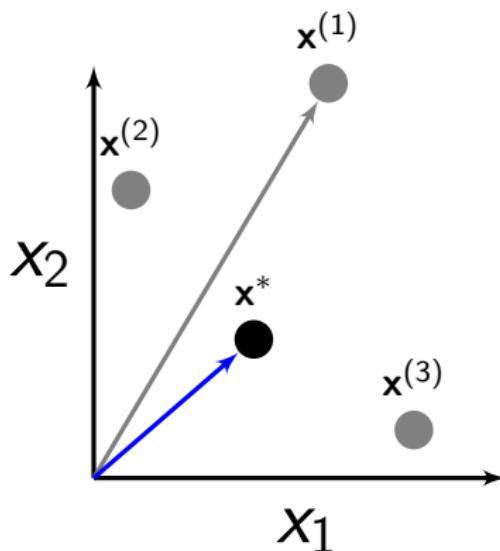
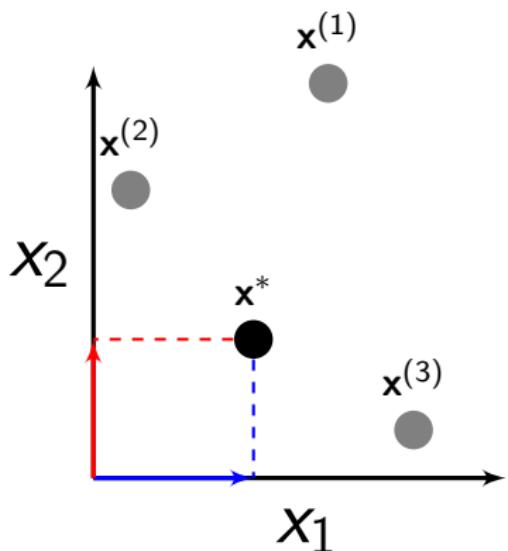
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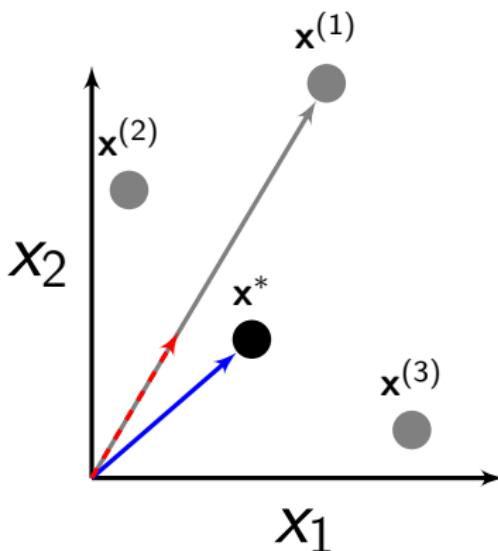
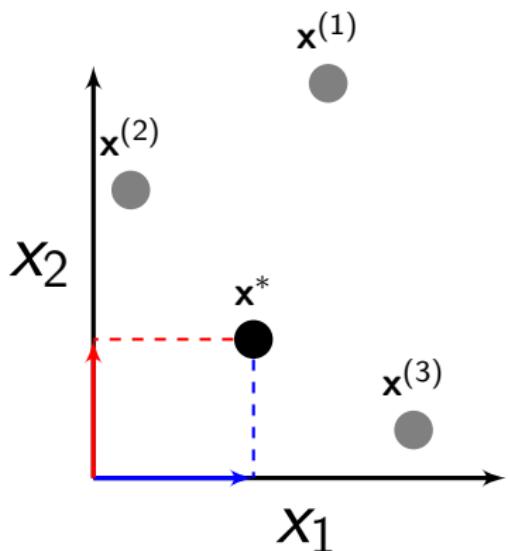
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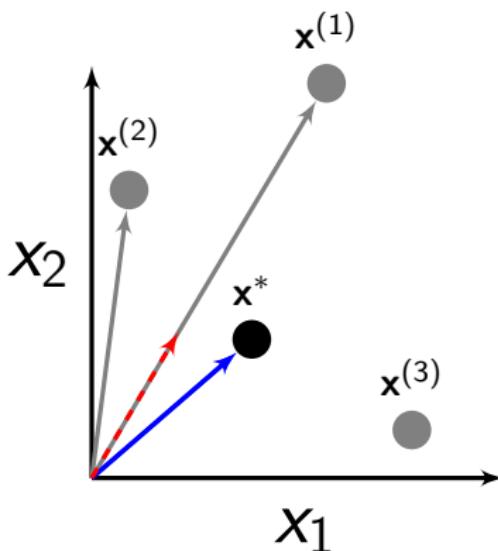
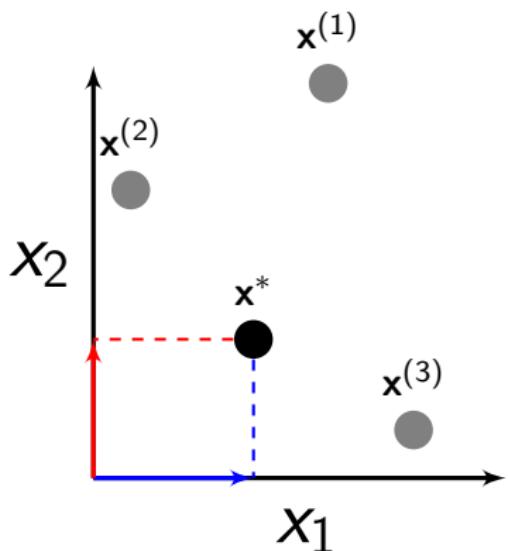
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# Linear kernel geometric picture

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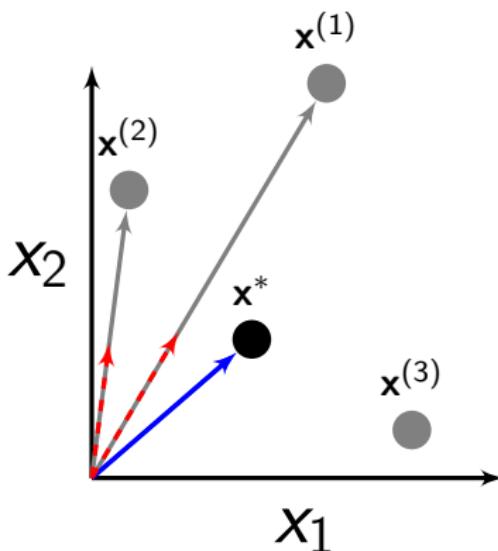
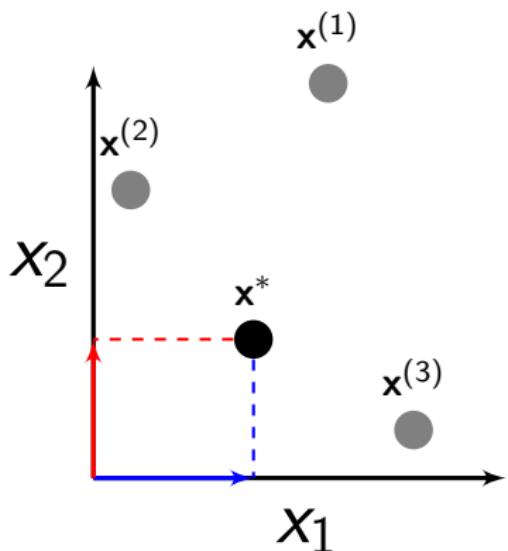
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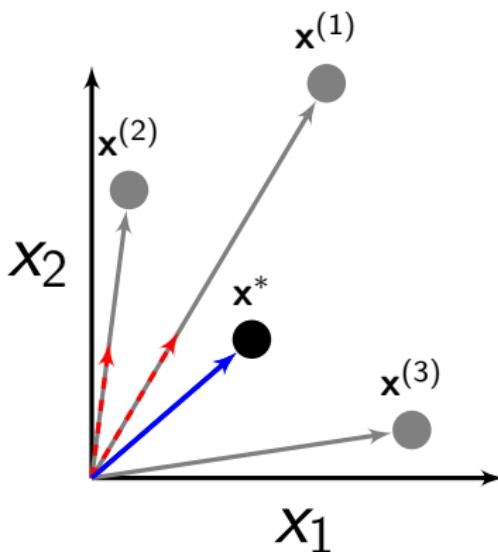
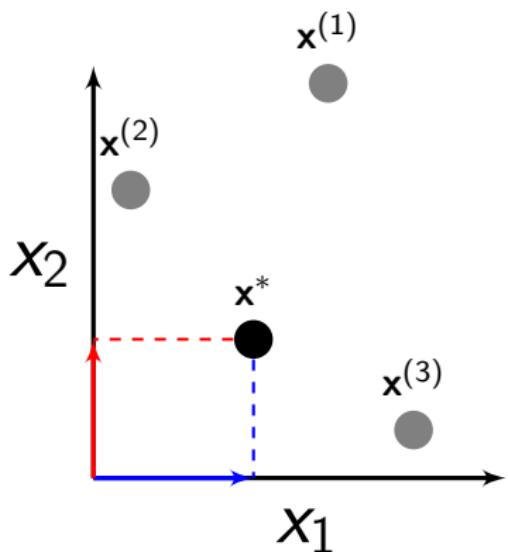
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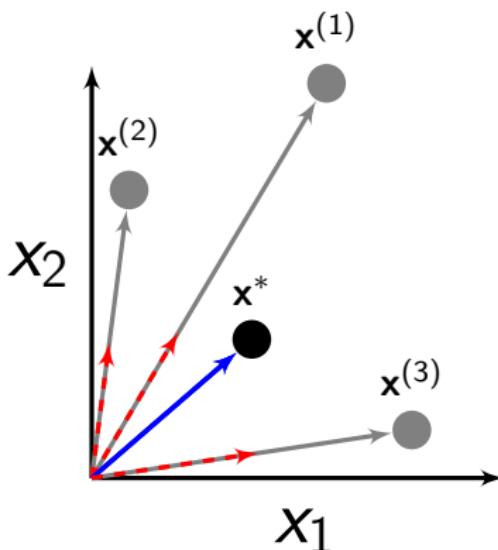
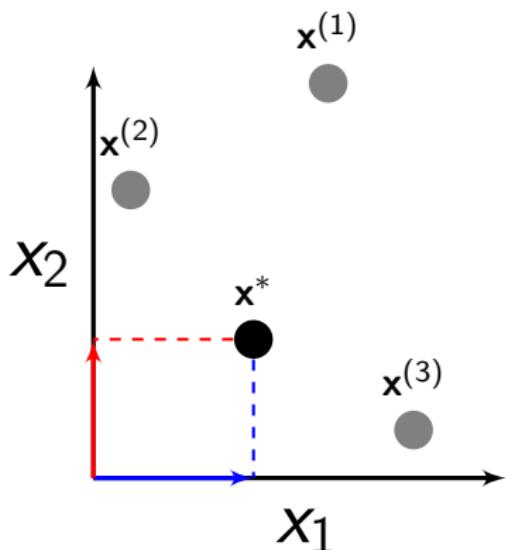
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Notice that this is *linear* in  $w$  for a 'lifted' feature space,  $\varphi(X)$ :

$$\begin{aligned} y_{QUAD}(x) &= \varphi(X)w \\ &= \begin{bmatrix} 1 & \sqrt{2}x_1^{(1)} & \sqrt{2}x_2^{(1)} & \sqrt{2}x_1^{(1)}x_2^{(1)} & (x_1^{(1)})^2 & (x_2^{(1)})^2 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & \sqrt{2}x_1^{(n)} & \sqrt{2}x_2^{(n)} & \sqrt{2}x_1^{(n)}x_2^{(n)} & (x_1^{(n)})^2 & (x_2^{(n)})^2 \end{bmatrix} \begin{bmatrix} w_1 \\ \vdots \\ w_6 \end{bmatrix} \end{aligned}$$

except the dimension has increased from  $\mathbb{R}^{n \times 2} \rightarrow \mathbb{R}^{n \times 6}$

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by direct analogy to the previous slides, there is also a kernel form:

$$\hat{y}(x^*) = \sum_{i=1}^n k(x^*, x_i) a_i$$

$$k(x^*, x_i) = \langle \varphi(x_i), \varphi(x_j) \rangle$$

$$= \begin{bmatrix} 1 & \sqrt{2}x_1^{(i)} & \dots & (x_1^{(i)})^2 & (x_2^{(i)})^2 \end{bmatrix} \begin{bmatrix} 1 \\ \sqrt{2}x_1^{(j)} \\ \vdots \\ (x_1^{(j)})^2 \\ (x_2^{(j)})^2 \end{bmatrix}$$

# The “kernel trick”

Notice that all that is required is vector products, i.e.

$$K_{i,j} = \langle \varphi(x_i), \varphi(x_j) \rangle$$

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which can be computed entirely using vectors in  $\mathbb{R}^2$ , so we never have to allocate the (factorially large) feature space.

# Detailed example of nonlinear regression

jupyter notebook: [github.com/jpjanet/ML-chem-workshop/  
blob/master/notebooks/workshop\\_compare\\_models.ipynb](https://github.com/jpjanet/ML-chem-workshop/blob/master/notebooks/workshop_compare_models.ipynb)

# General kernels

Both kernel methods are the same except:

	linear	quadratic
$K_{ij}$	$(x^{(i)})^T x^{(j)}$	$((x^{(i)})^T x^{(j)} + 1)^2$

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From the perspective of similarity, we can imagine arbitrary functions to be our kernel, without ever needing to know what the underlying feature map  $\varphi$  is.

# The Gaussian kernel and KRR

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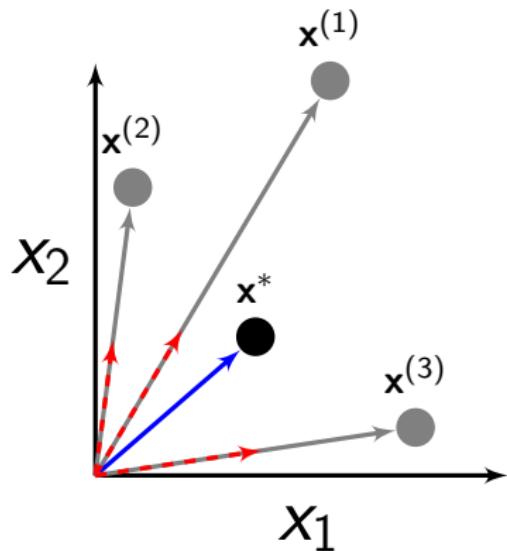
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Depends on  $\sigma$  to control non-locality.

# Similarity and Gaussian KRR

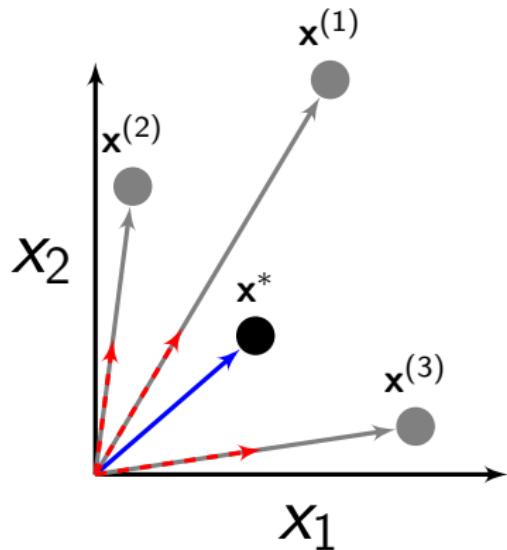
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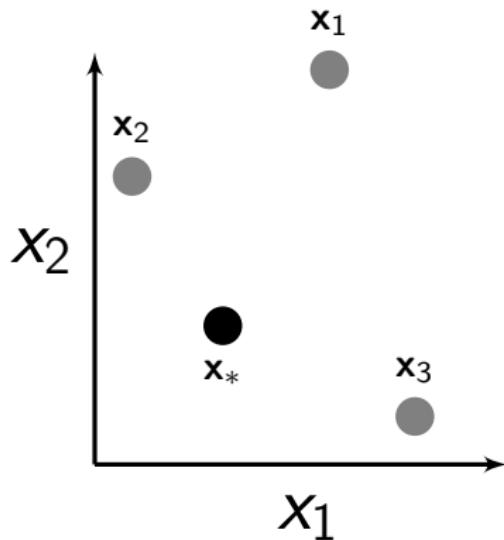
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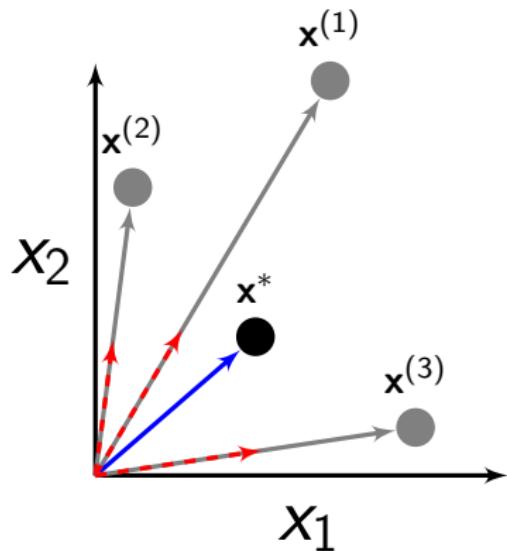
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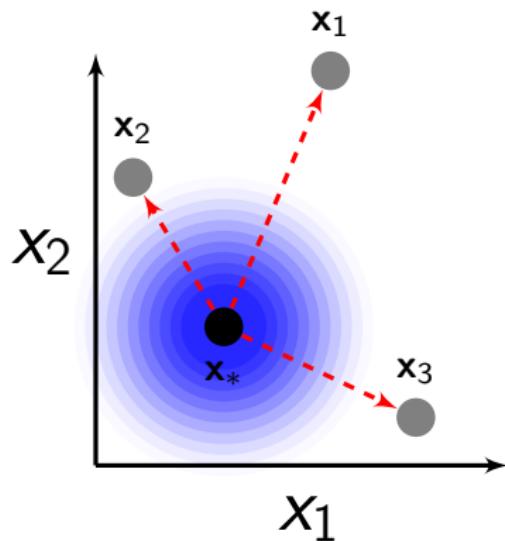
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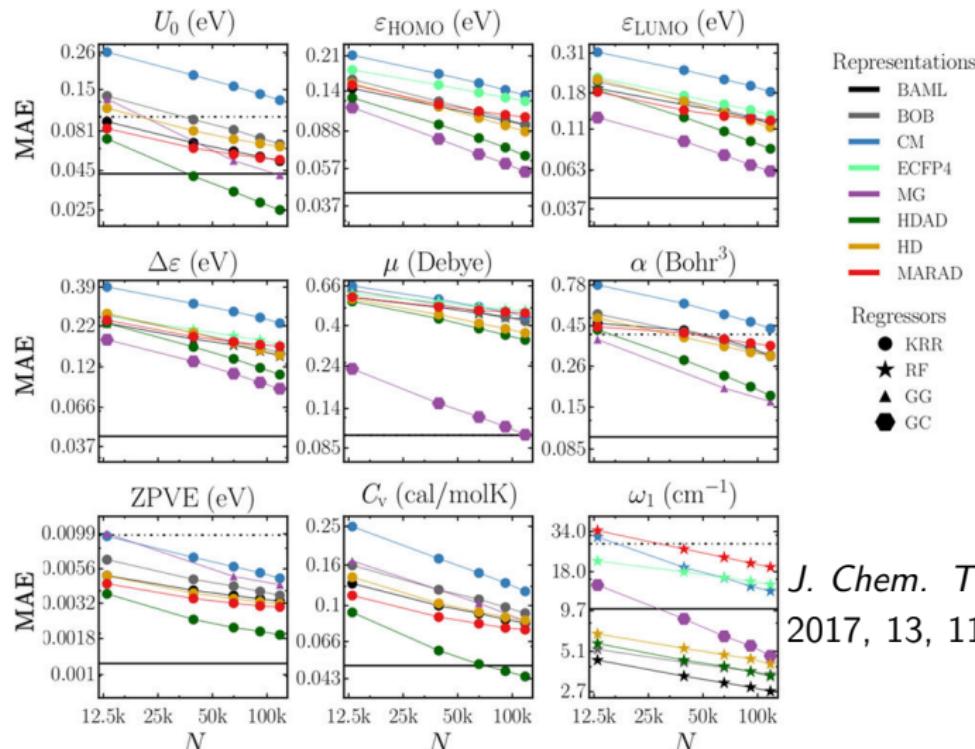
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- 4 check using cross-validation to choose  $\sigma$  and  $\lambda$

# KRR is widely used in chemistry



*J. Chem. Theory Comput.*  
2017, 13, 11, 5255–5264

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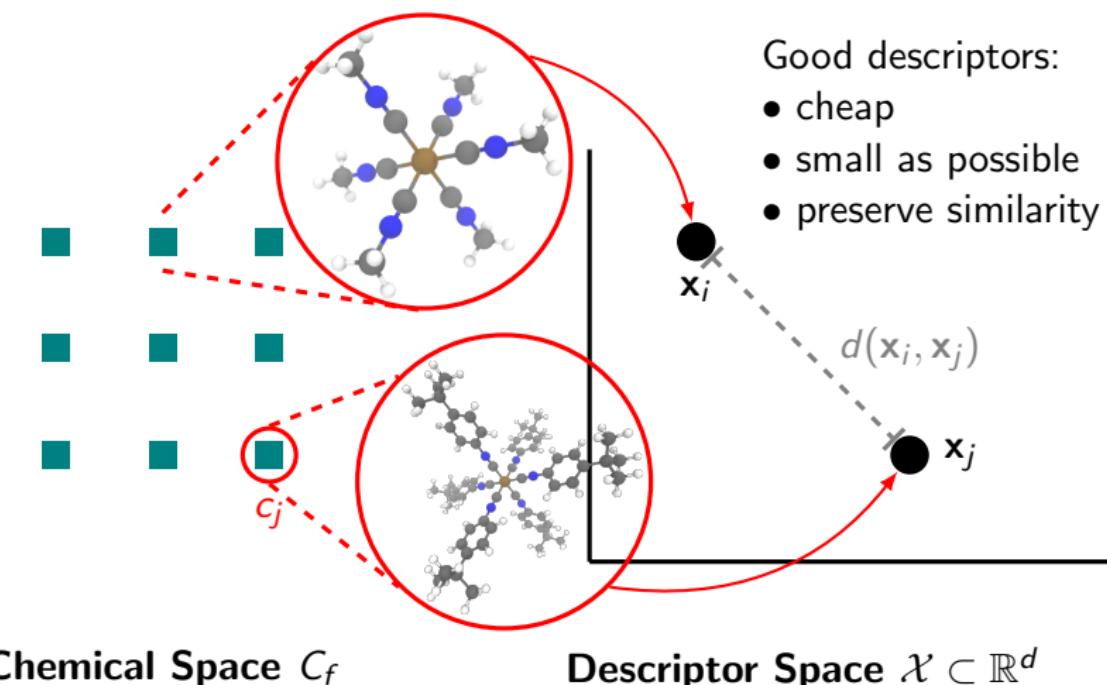
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# KRR example

jupyter notebook: [github.com/jpjanet/ML-chem-workshop/  
blob/master/notebooks/KRR.ipynb](https://github.com/jpjanet/ML-chem-workshop/blob/master/notebooks/KRR.ipynb)

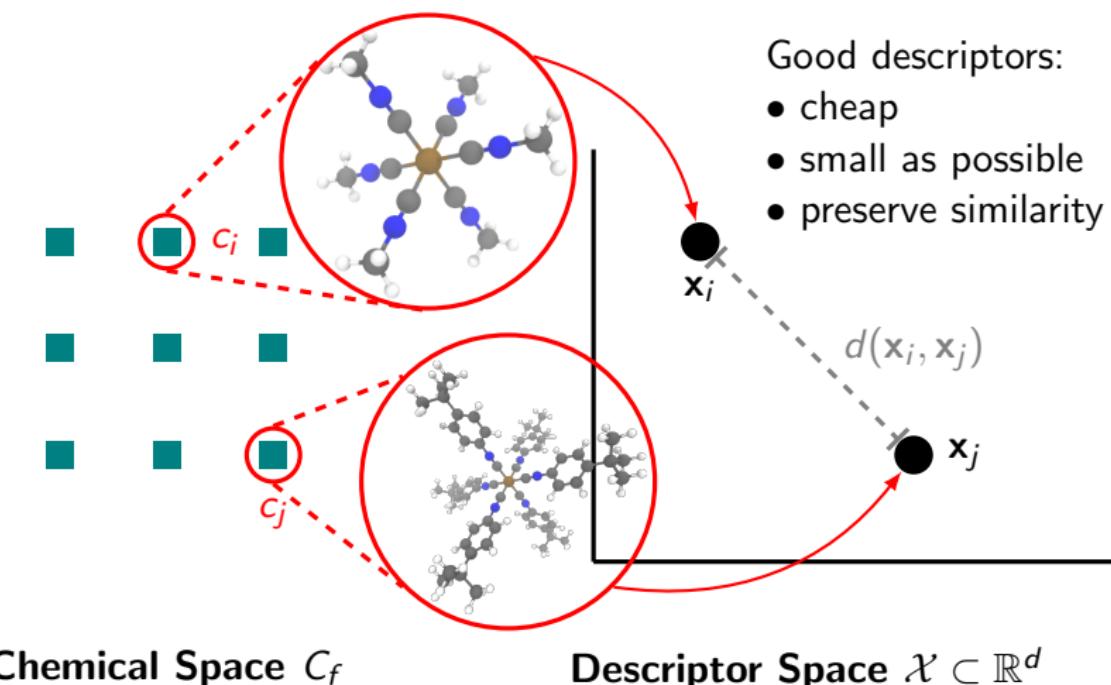
# Purpose



Chemical Space  $C_f$

Descriptor Space  $\mathcal{X} \subset \mathbb{R}^d$

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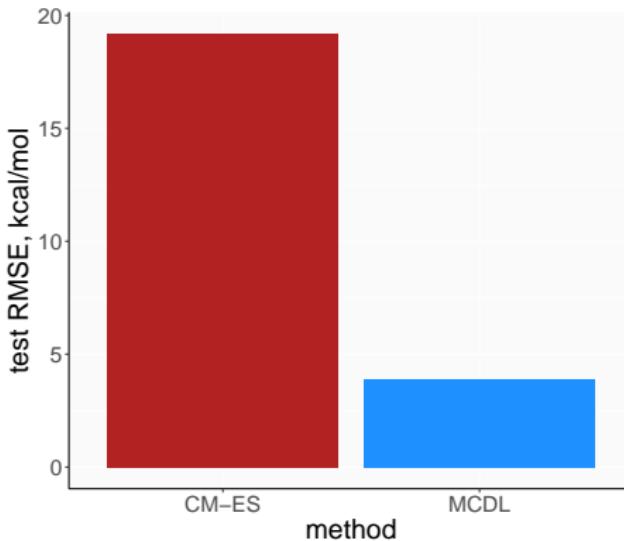


# Why similarity is important

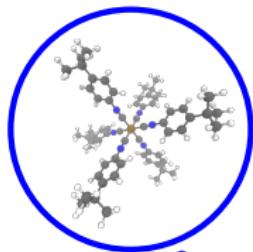
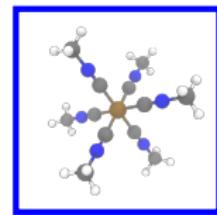
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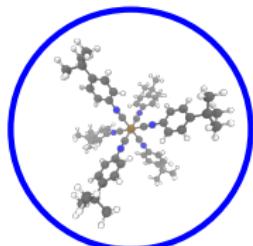
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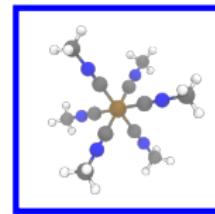
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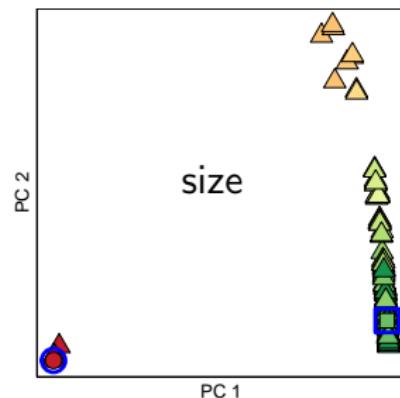
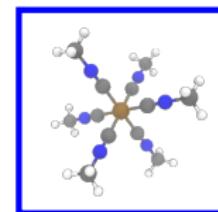
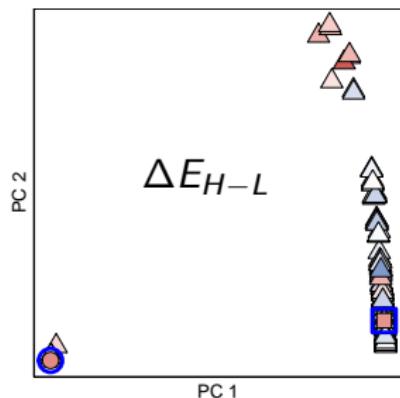
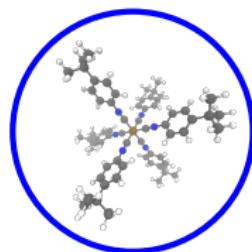
$\Delta E_{\text{H-L}} = 37.7 \text{ kcal/mol}$



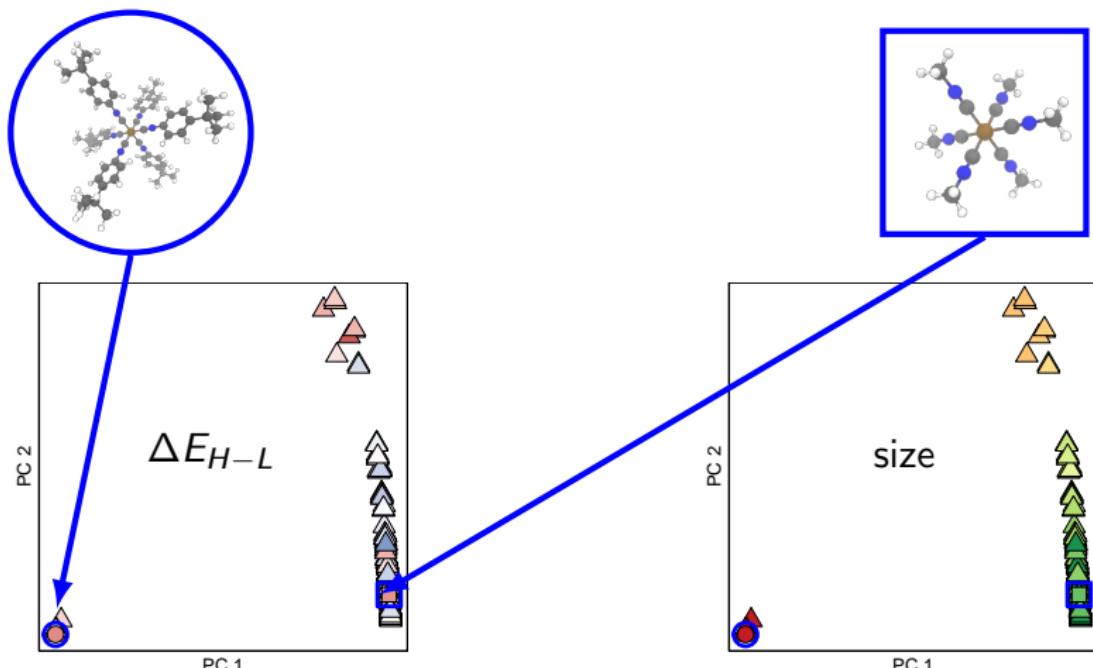
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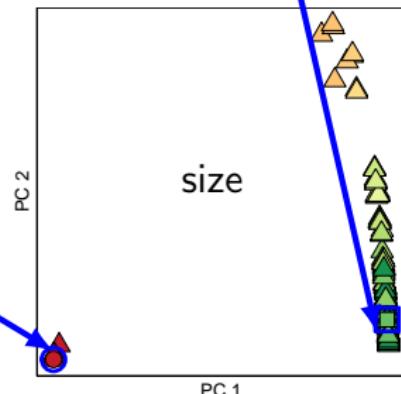
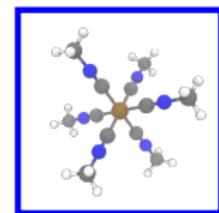
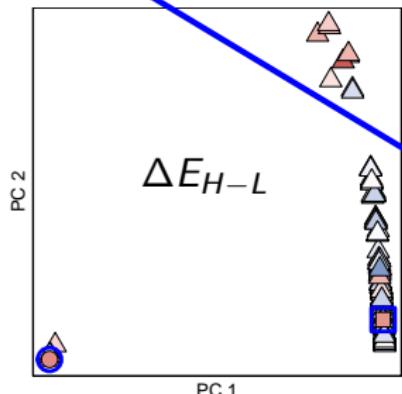
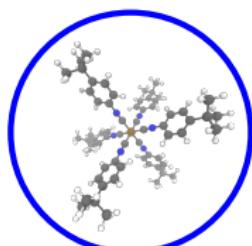
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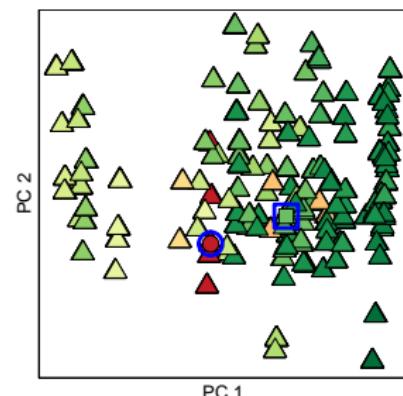
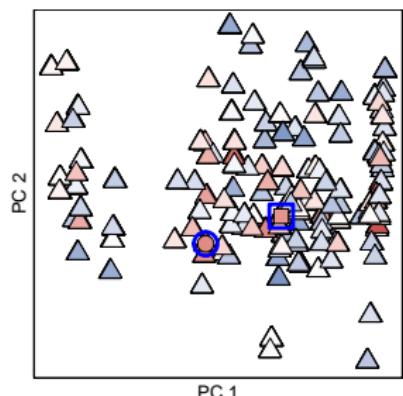
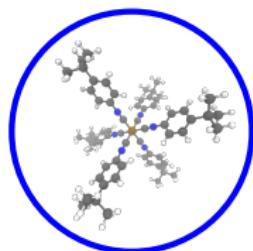
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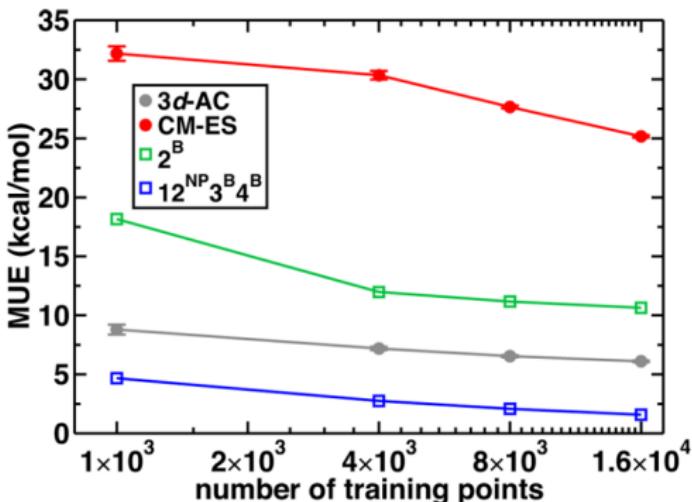


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complexity

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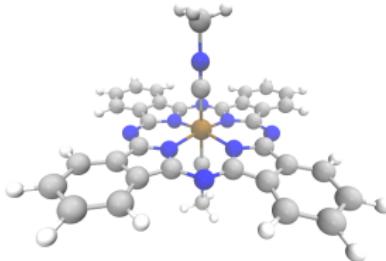


## 3D structure

- fine-grained structural information in 3D
- mimic input to a quantum chemistry code
- expensive to compute, rich information

## Ad-hoc properties

Sometimes, simple lists of atomic properties are sufficient, especially if informed by domain knowledge:

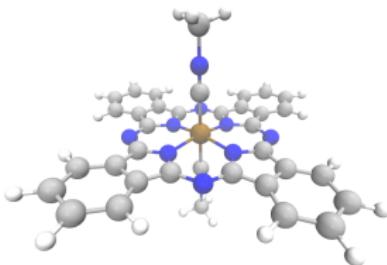


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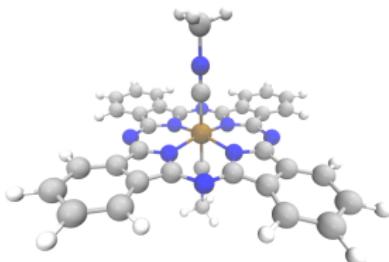
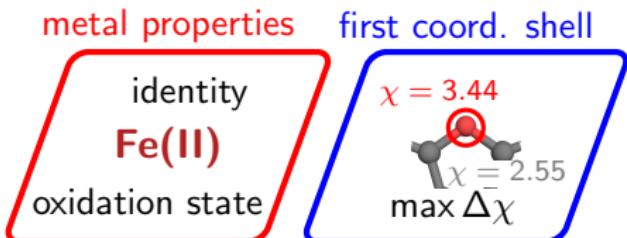
metal properties

- identity
- Fe(II)**
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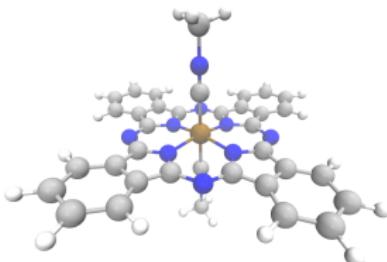
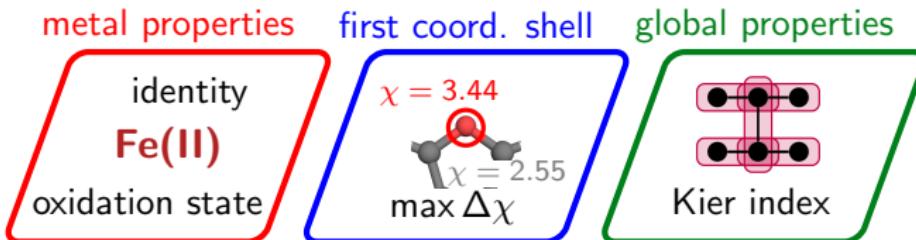
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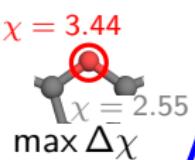
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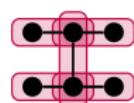
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**Fe(II)**

oxidation state

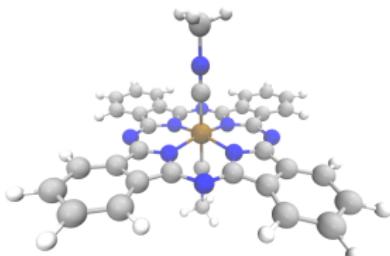
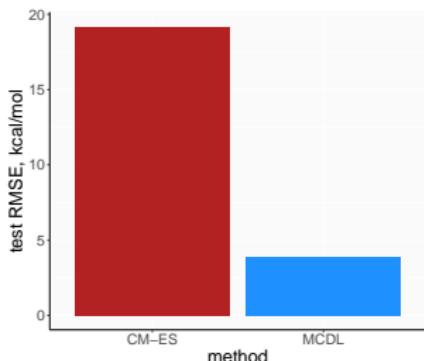
first coord. shell



global properties



Kier index



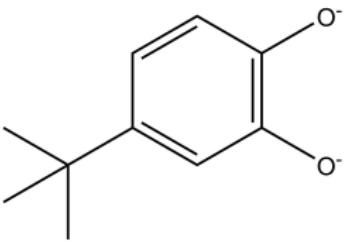
Janet, J.P., and Kulik, H.J. *Chem. Sci.*, 2017, 8, 5137-5152.

# Fingerprints and the low-information limit

In cheminformatics (esp. drug design literature) fingerprints are binary vectors used to determine molecular similarity. For example, FP2 fingerprint is a 1024 bit fingerprint:

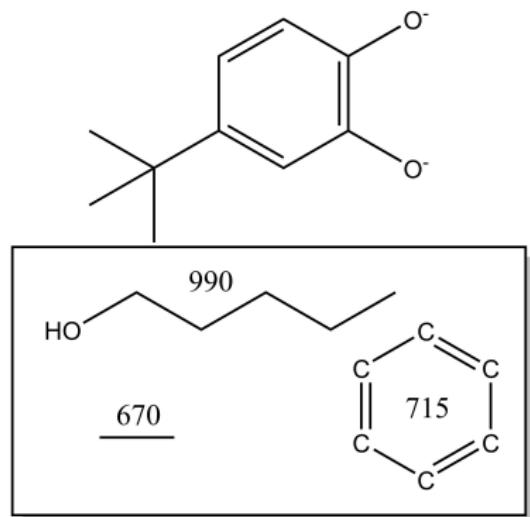
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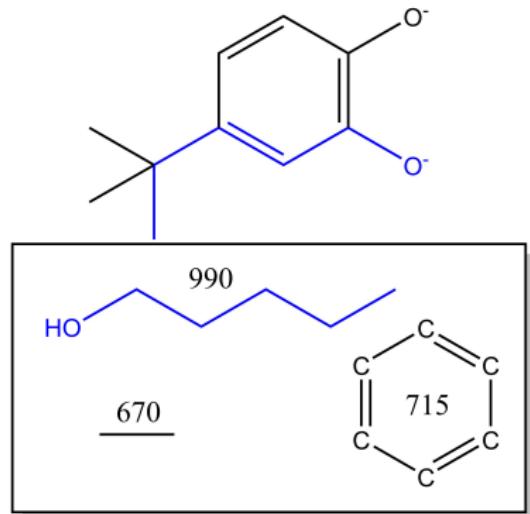
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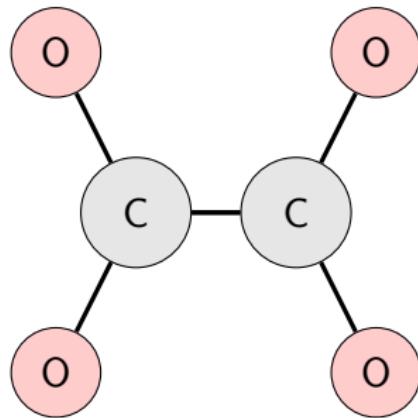
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# Molecular graphs

Based on autocorrelations<sup>1</sup>

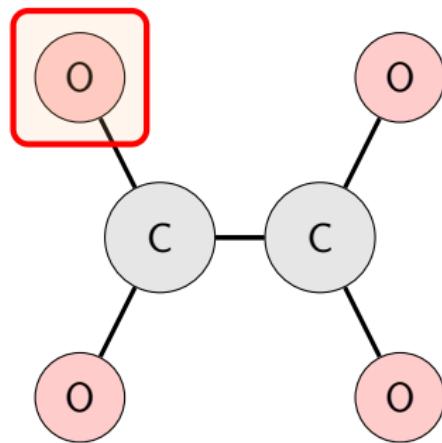


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<sup>1</sup>Broto, P., Moreau, G. and Vandycke, C. *Eur. J. Med. Chem.*, 19(1):71-78, 1984.

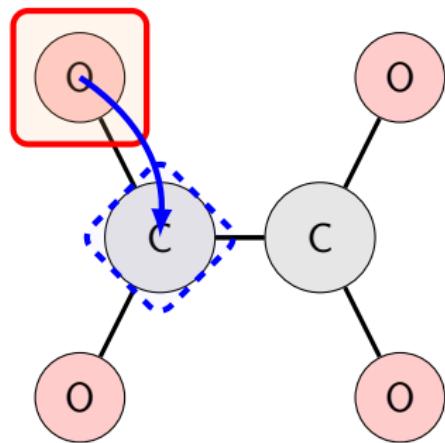
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Based on autocorrelations and modified for TMCs<sup>4</sup>



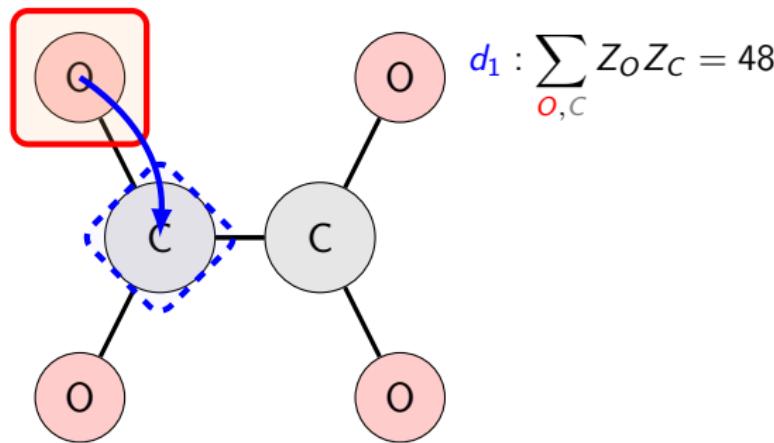
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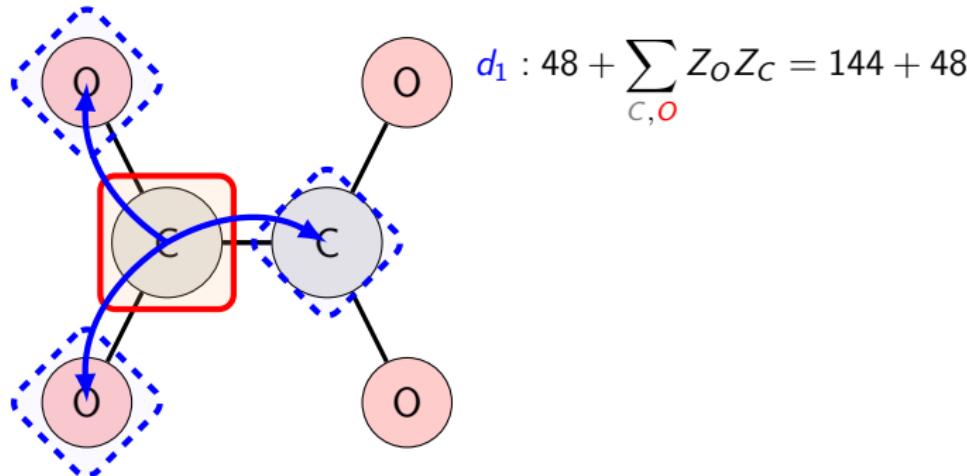
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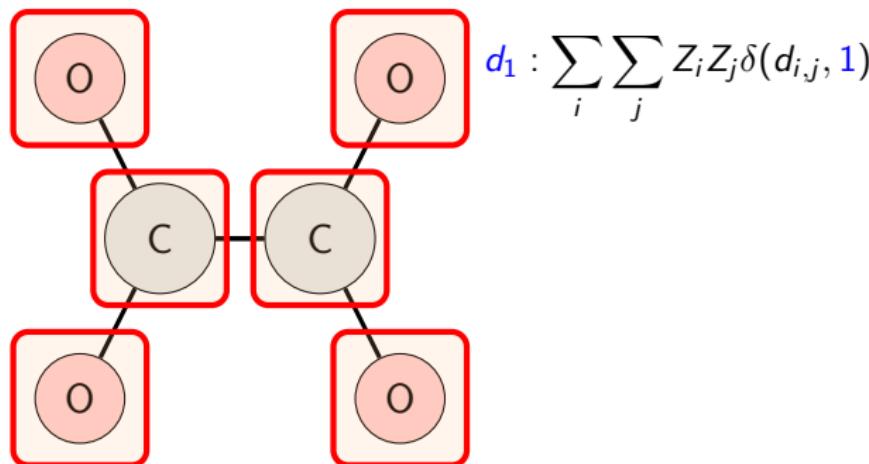
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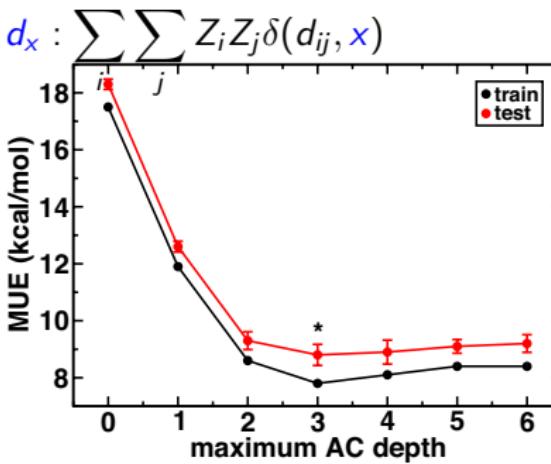
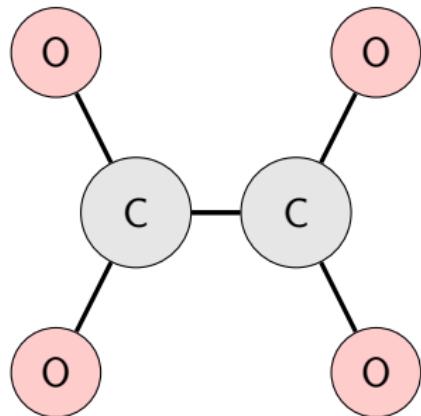
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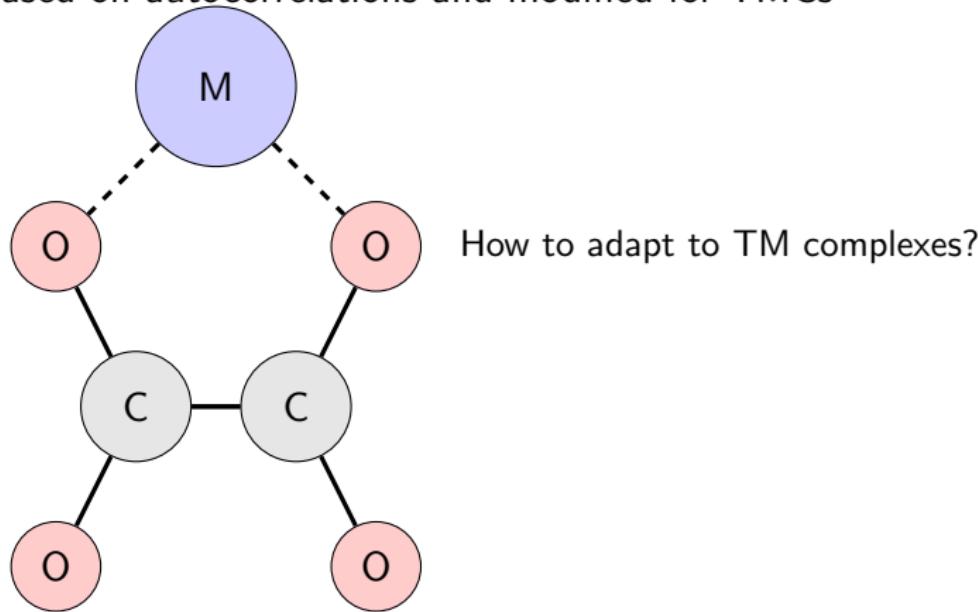
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<sup>4</sup>Janet, J.P., and Kulik, H.J. *J. Phys. Chem. A*, 2017, 121, 46, 8939-8954.

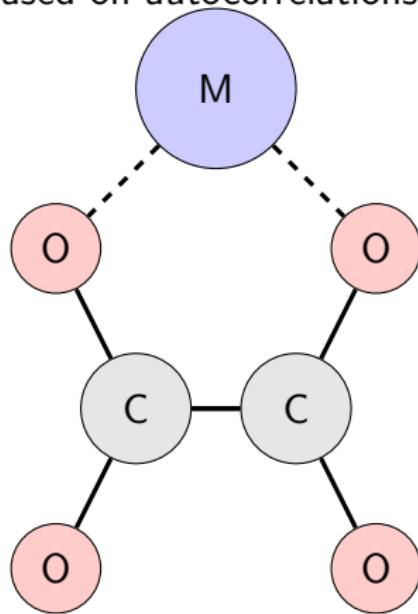
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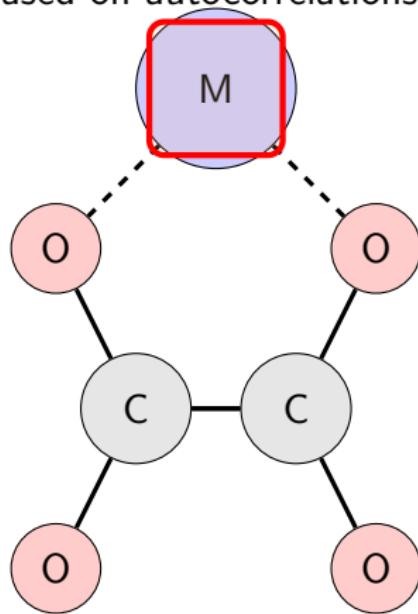
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How to adapt to TM complexes?  
restrict the scope to focus on  
*near-metal atoms*

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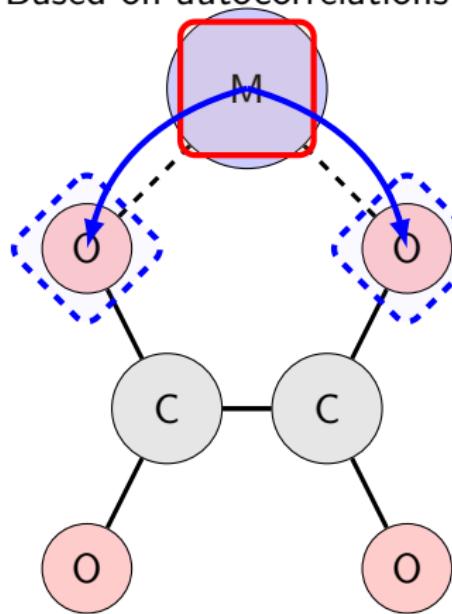
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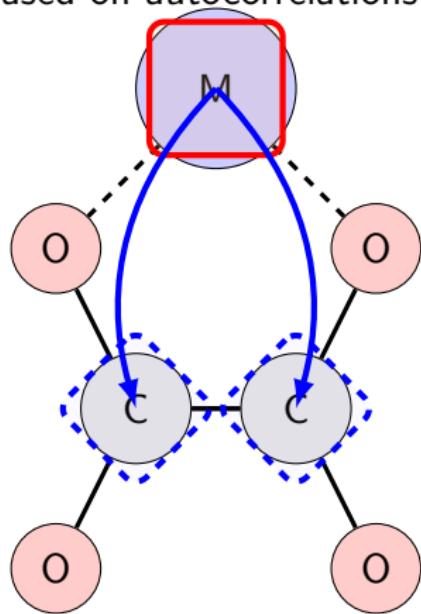


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$$d_1 : \sum_{M,O} Z_M Z_O$$

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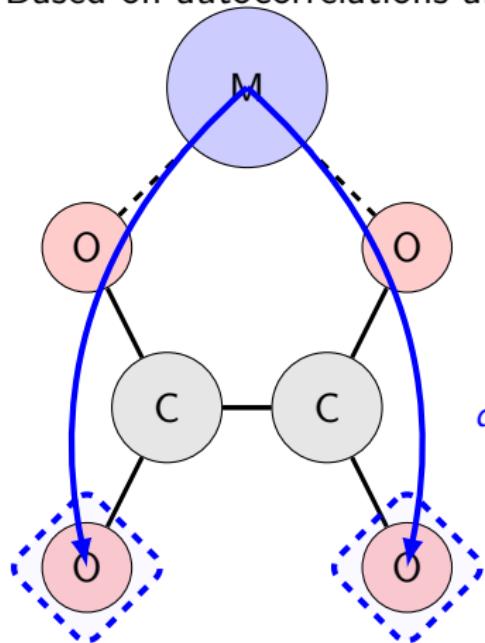


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$$d_2 : \sum_{M,C} Z_M Z_C$$

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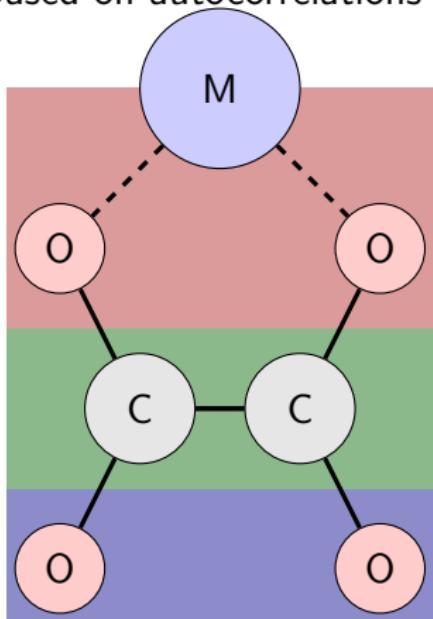


How to adapt to TM complexes?  
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$$d_3 : \sum_{M,O} Z_M Z_O$$

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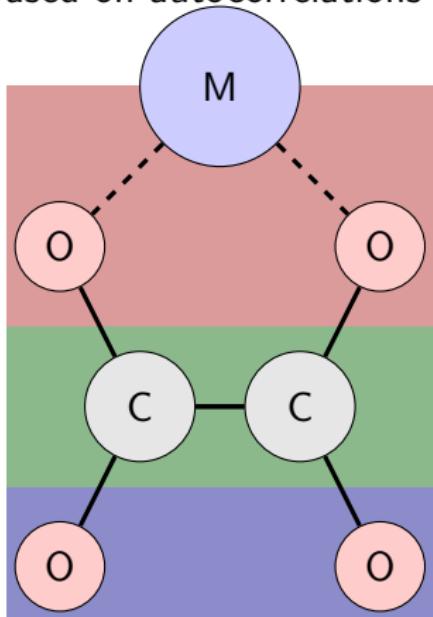


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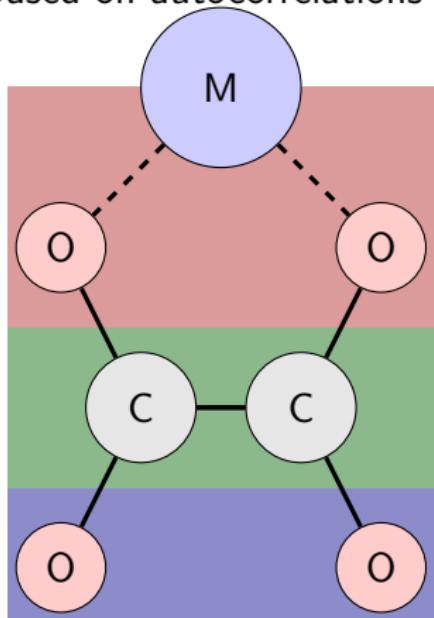
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$$(Z_i - Z_j)$$

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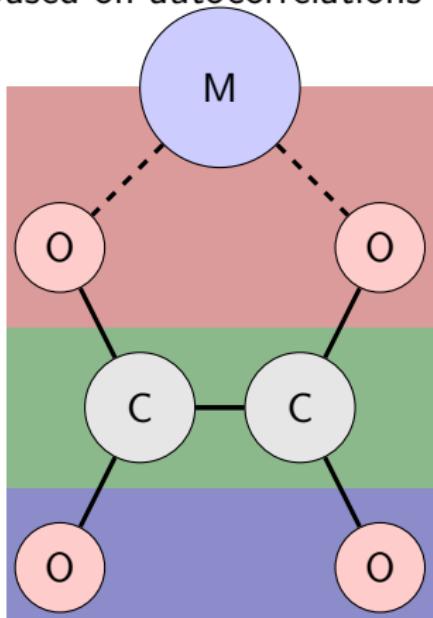
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$$d_3 : \sum_{M,O} Z_M Z_O (Z_i - Z_j)$$

properties:  $T, \chi, Z, I, S$

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$$d_3 : \sum_{M,O} Z_M Z_O (Z_i - Z_j)$$

~ 160 features in total

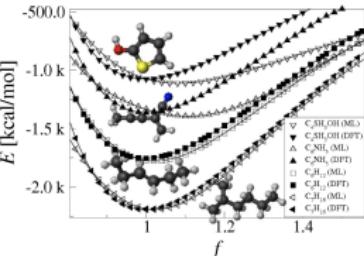
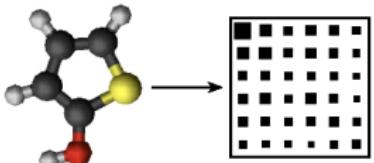
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# Coulomb matrices

One family of 3D descriptors attempt to copy information used in quantum chemistry codes, e.g. Coulomb Matrices:

Montavon, G. et al.. Learning Invariant Representations of Molecules for Atomization Energy Prediction, NIPS 25, 2012

$$M_{I,J} = \begin{cases} 0.5Z_I^{2.4} & \text{for } I = J \\ \frac{Z_I Z_J}{|R_I - R_J|} & \text{for } I \neq J \end{cases}$$

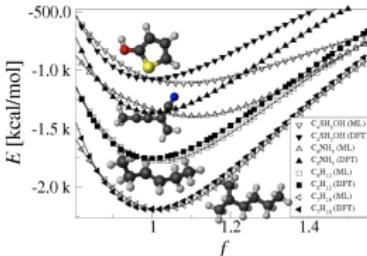
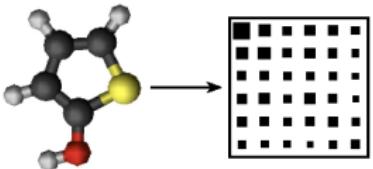


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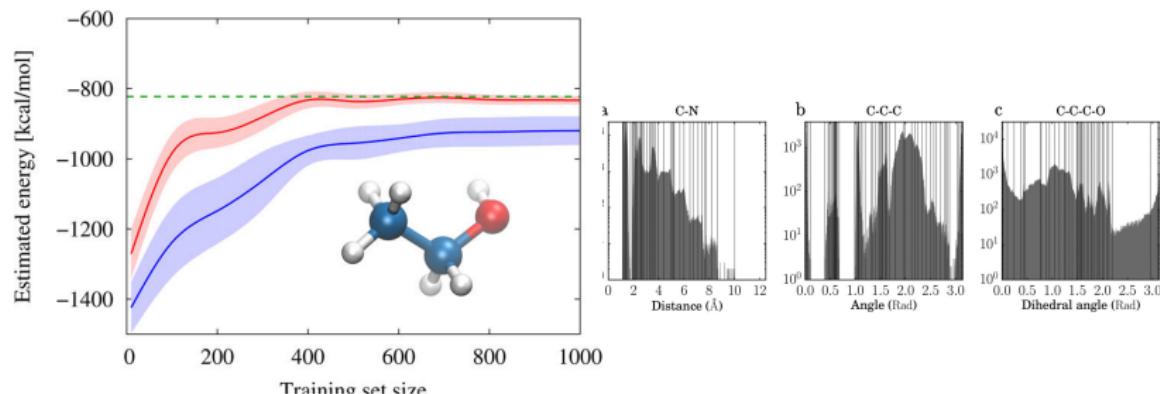


rotational and translational invariance

# HDAD and beyond-CM

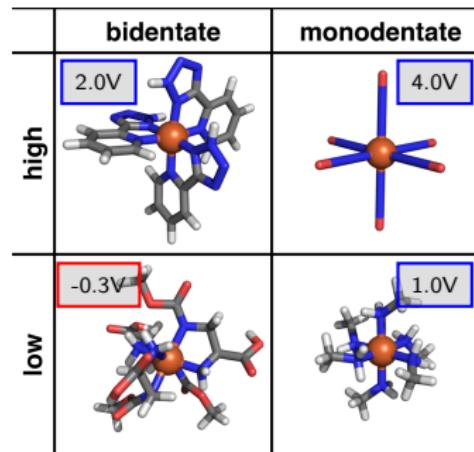
Subsequent work adds descriptors derived from geometric parameters, i.e. bonds, angles, and dihedral angles:

Faber, F. et al.. Prediction Errors of Molecular Machine Learning Models Lower than Hybrid DFT Error, *J. Chem. Theory Comput.* 2017, 13, 11, 5255-5264



# Why do feature selection?

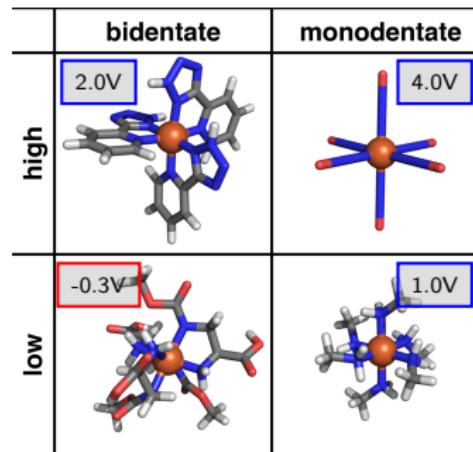
This can help explain what factors are important<sup>5</sup>:



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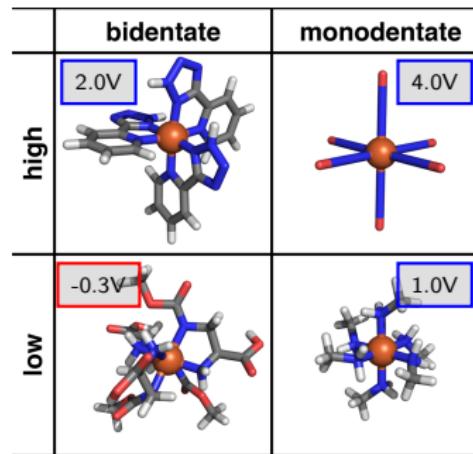


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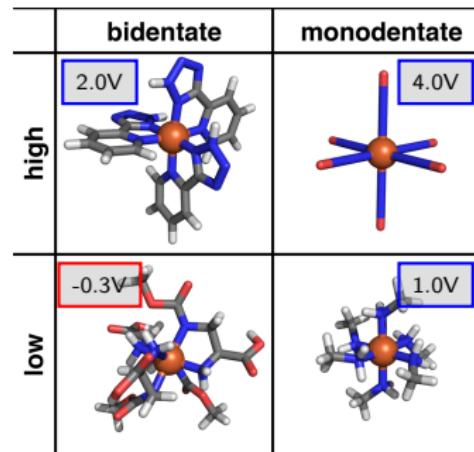


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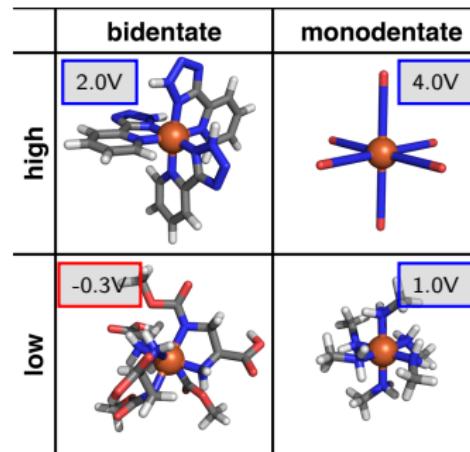
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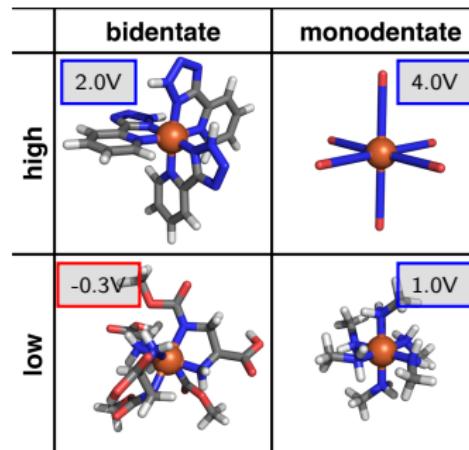
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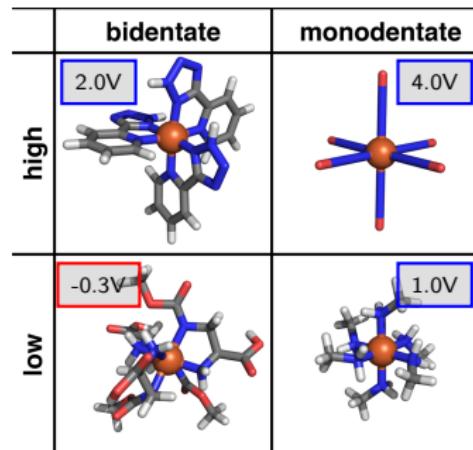
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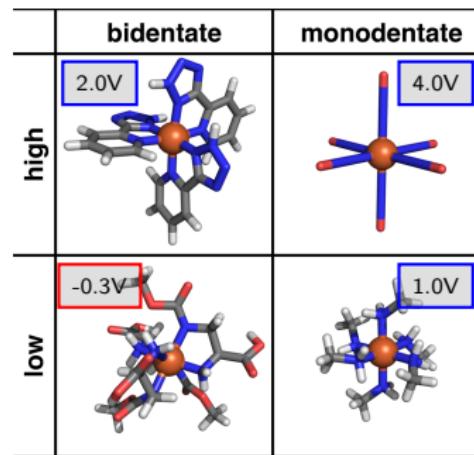
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Can use *autocorrelation* functions to describe how ligands and atoms are connected

$$\eta_{i,0} = p_i p_i$$

$$\eta_{i,k} = \sum_{i \neq j} p_i p_j \delta(d_{i,j} - d_k), \quad k \neq 0$$

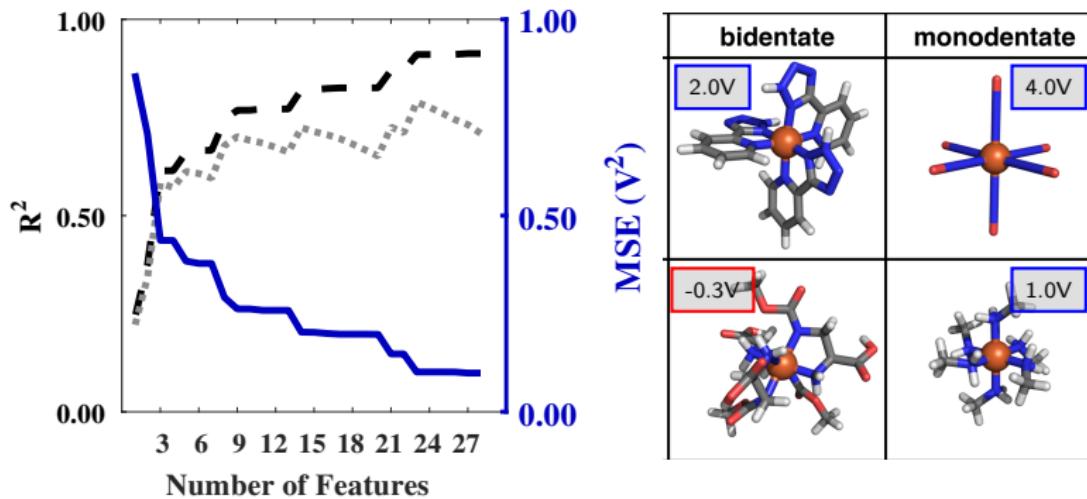
$$\text{AC}_k = \sum_i \eta_{i,k}$$

4 properties,  $k \in [0, 5]$   
 $\implies$  28 variables.

How to choose?

# Why do feature selection?

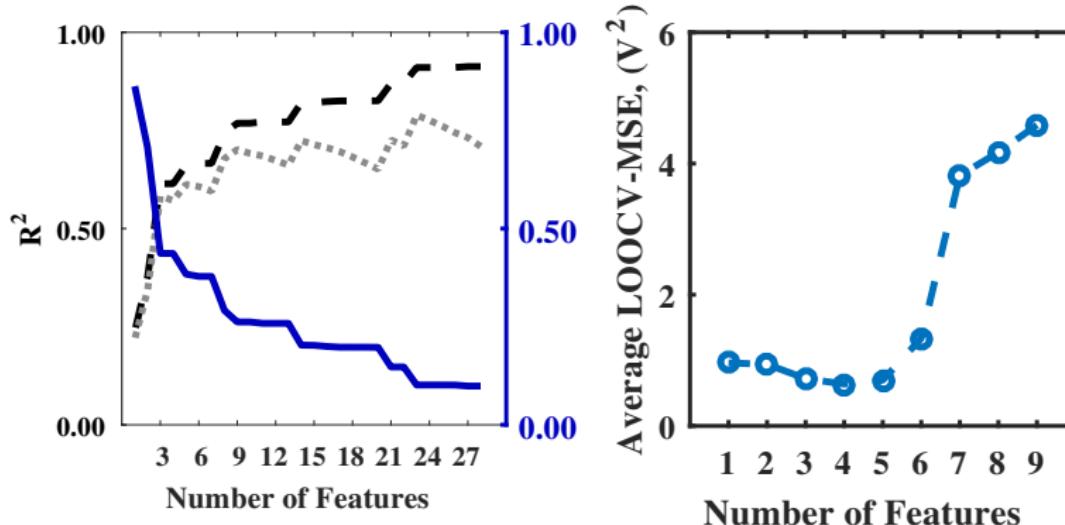
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## How to pick important features?

Using unnecessary features can degrade model performance, so we want to able to pick the subset of variables that is best correlated with our objective, formally:

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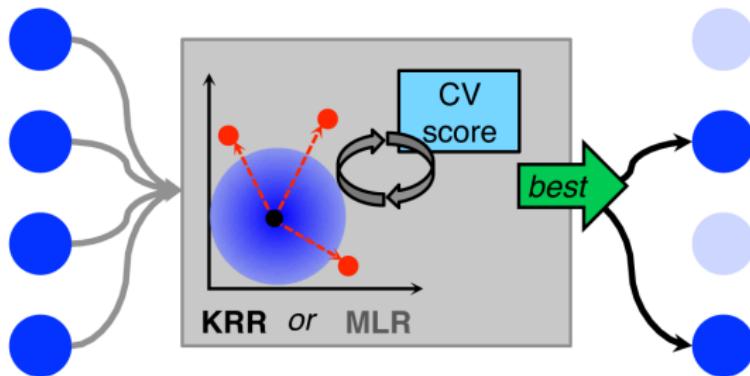
We don't know the optimal number upfront, and this is a combinatorial problem – possible for  $\leq 30$  dimensions or so, but rapidly becomes unfeasible.

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- $\lambda$  comes from cross-validation

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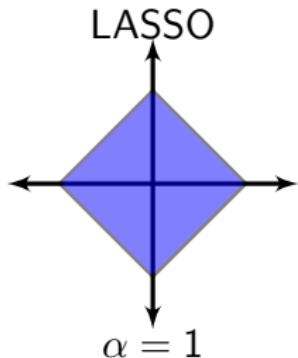
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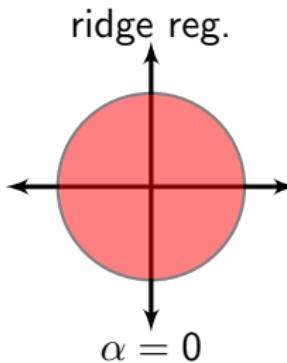
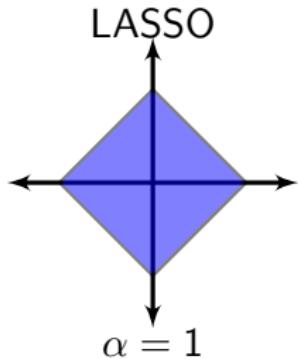
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$$w = \arg \min_{w \in \mathbb{R}^p} \|Xw - y_{data}\|_2^2 + \lambda (\alpha \|w\|_1 + (1 - \alpha) \|w\|_2^2)$$



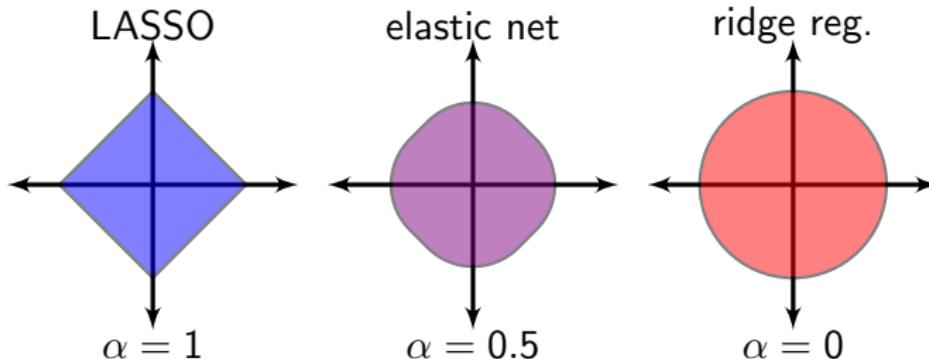
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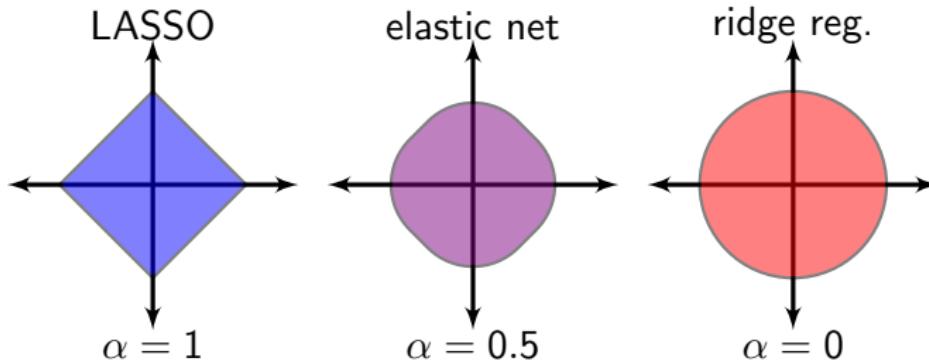
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Using even a small  $\alpha > 0$  ensures the minimization is stable.

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compare to

$$\|W\|_1^1 = \begin{vmatrix} 1 \\ 0 \end{vmatrix}_1 = |1| + |1| = 1$$

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- 3 Feature selection techniques can help identify important features, for modeling and for interpretation

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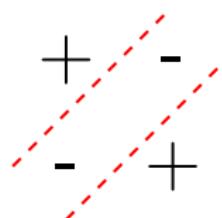
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Decision boundary

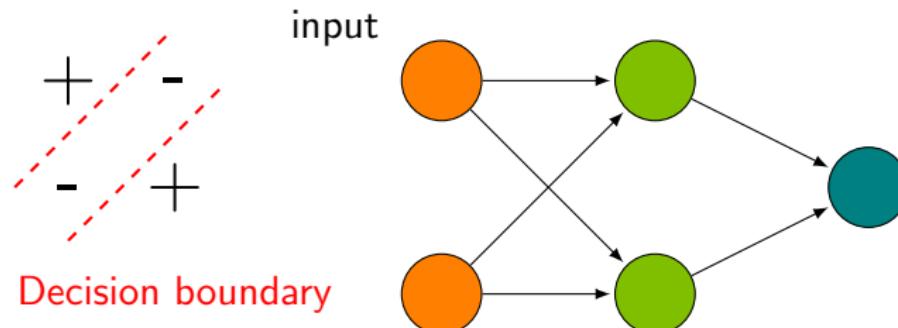
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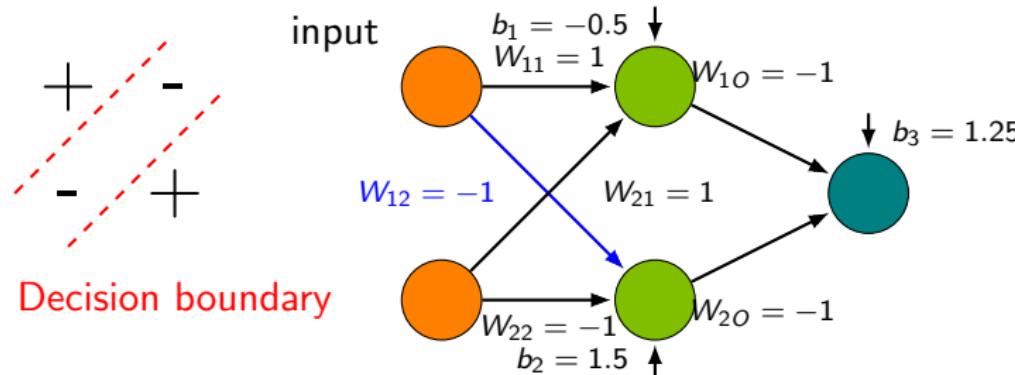
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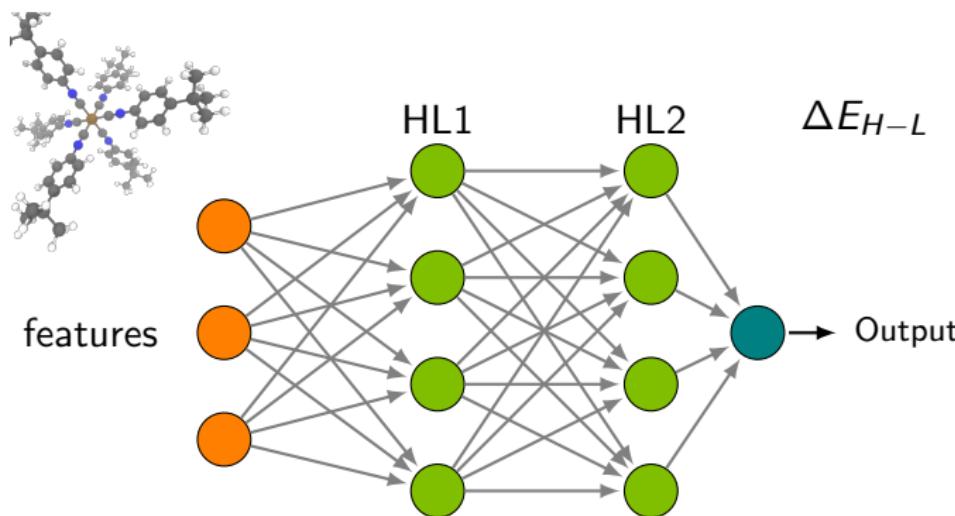
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# What do they look like?

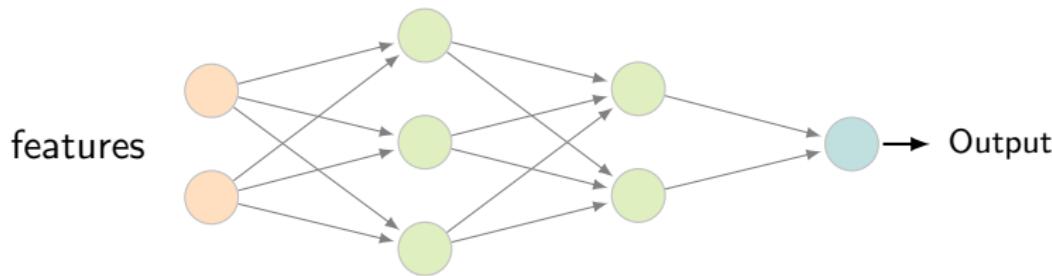


# Backpropagation (chain rule!)

Back-propagation updates neural network weights → example

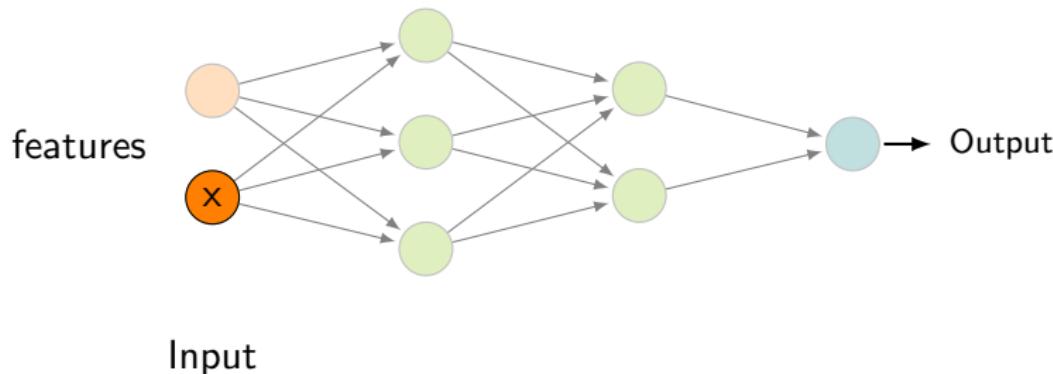
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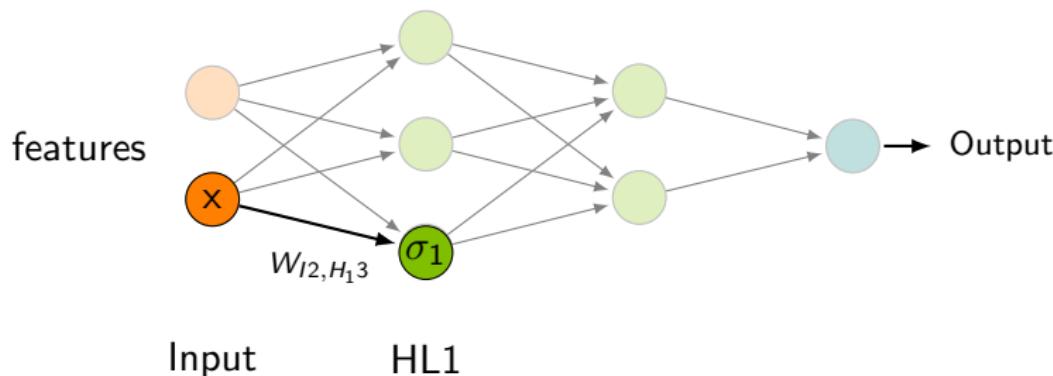
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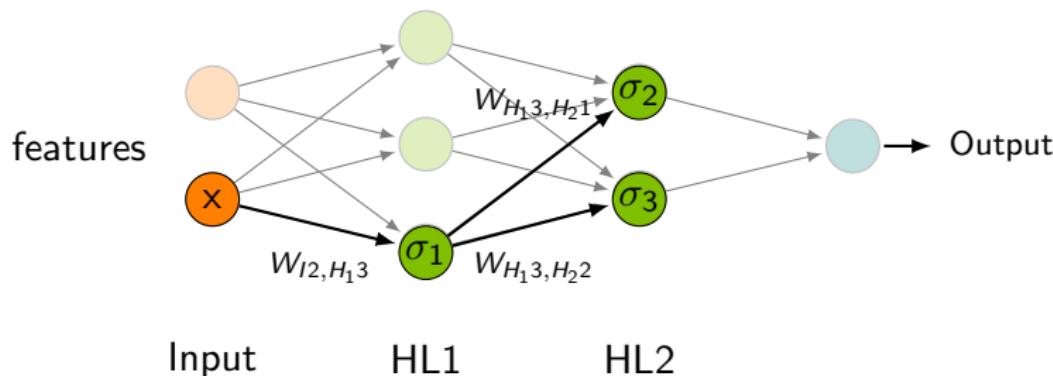
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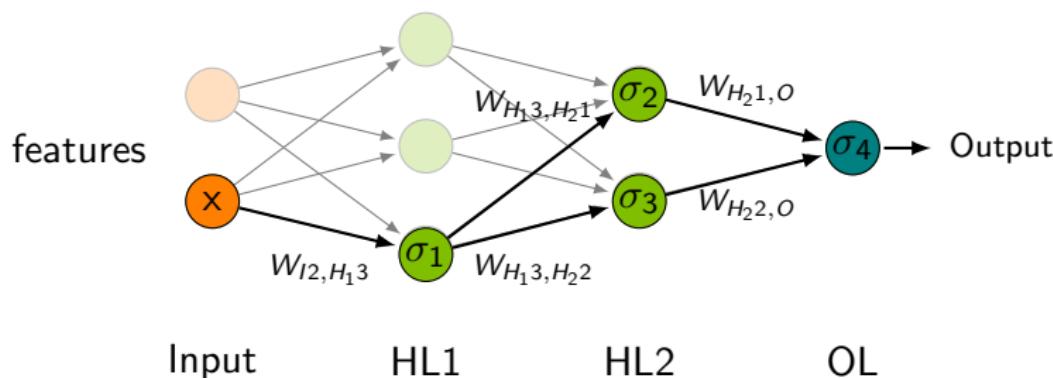
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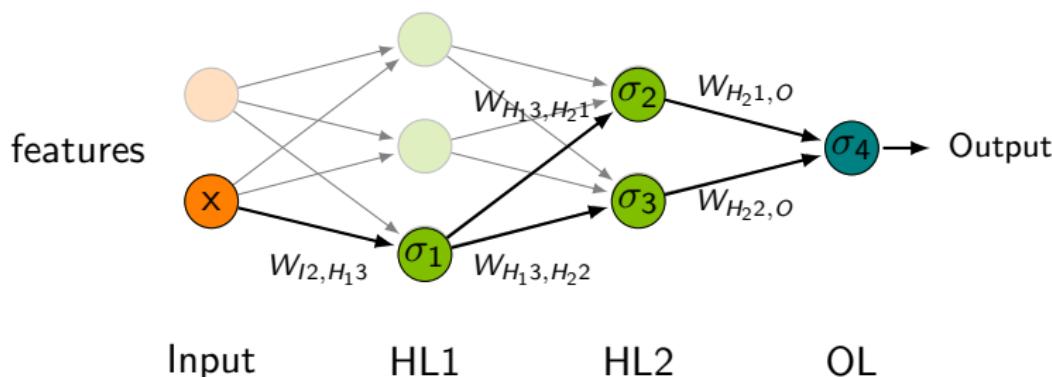
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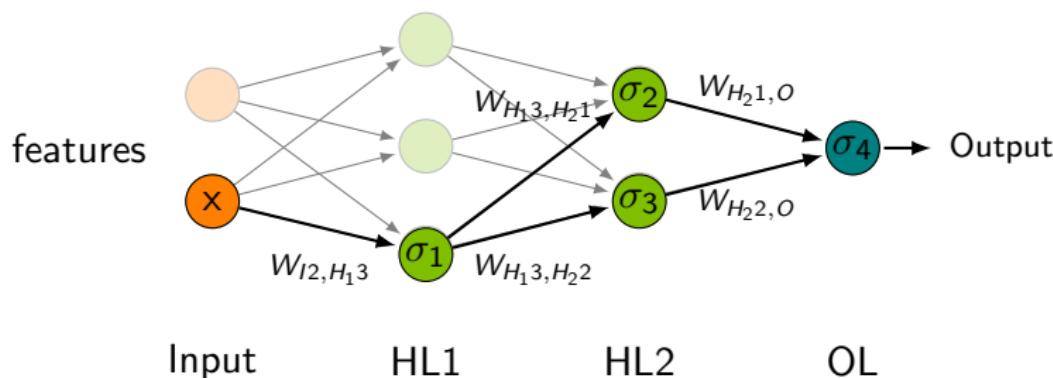
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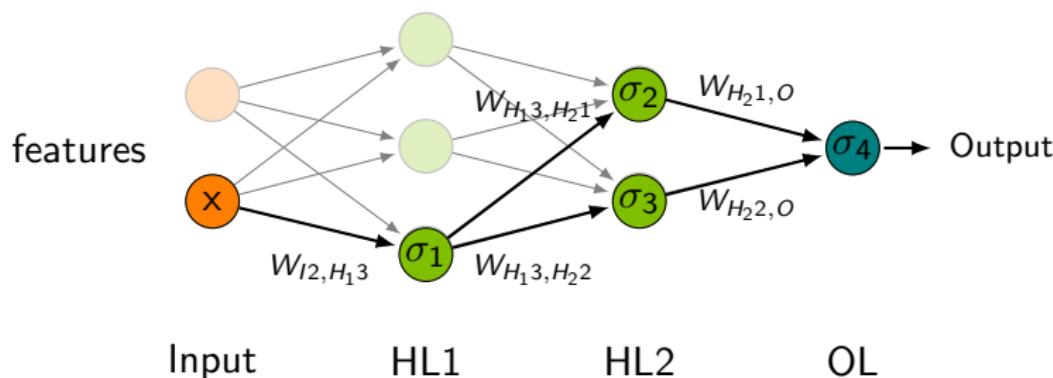
$$\text{Loss} \rightarrow \mathcal{L}(W) = \sum_{i=1}^N ((y_i - y_{pred})^2) + \lambda (\sum_{l=1}^L \|W_l\|^2)$$



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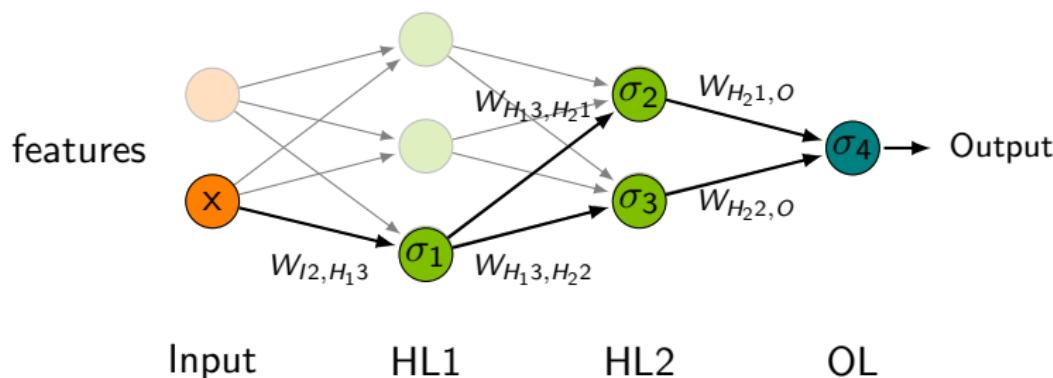
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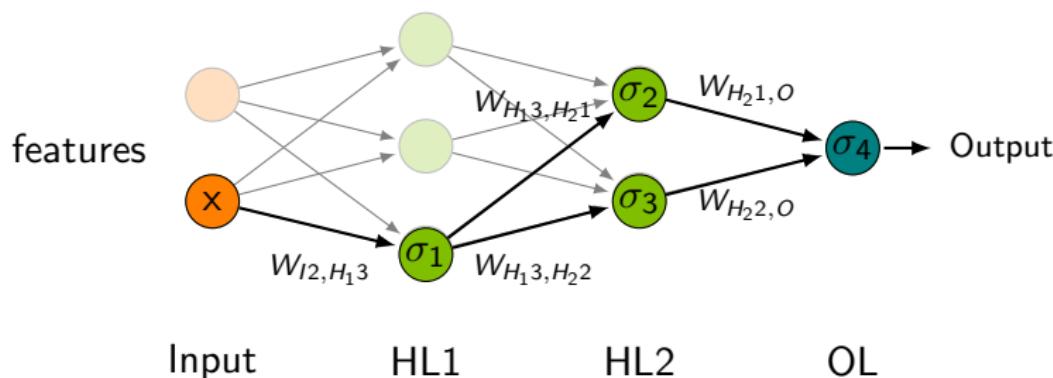
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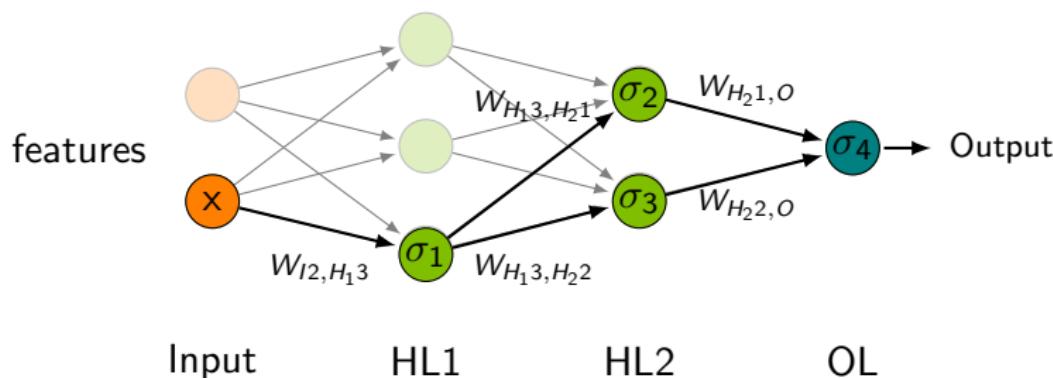
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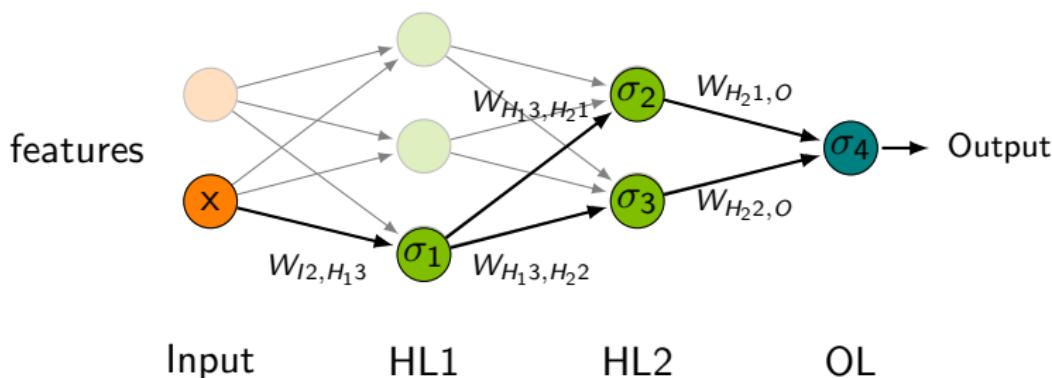


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$$\left( \frac{\partial \sigma_1}{\partial z_1} \right) \left( \frac{\partial z_1}{\partial W_{I2, H_13}} \right) + \left( \frac{\partial z_4}{\partial W_{H_22, O}} \right) \left( \frac{\partial W_{H_22, O}}{\partial \sigma_3} \right) \left( \frac{\partial \sigma_3}{\partial z_3} \right) \left( \frac{\partial z_3}{\partial W_{H_13, H_22}} \right) \left( \frac{\partial W_{H_13, H_22}}{\partial \sigma_1} \right) \left( \frac{\partial \sigma_1}{\partial z_1} \right) \left( \frac{\partial z_1}{\partial W_{I2, H_13}} \right)$$



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$$SGD \rightarrow \left( \frac{\partial Loss}{\partial W} \right) \rightarrow \text{one/few examples} \rightarrow \text{MUCH noisier!} \rightarrow \text{less stuck}$$

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**Recurrent** → Layers that “store” information about previous times, thus commonly used in speech or handwriting recognition

# Interpretation as representation learning

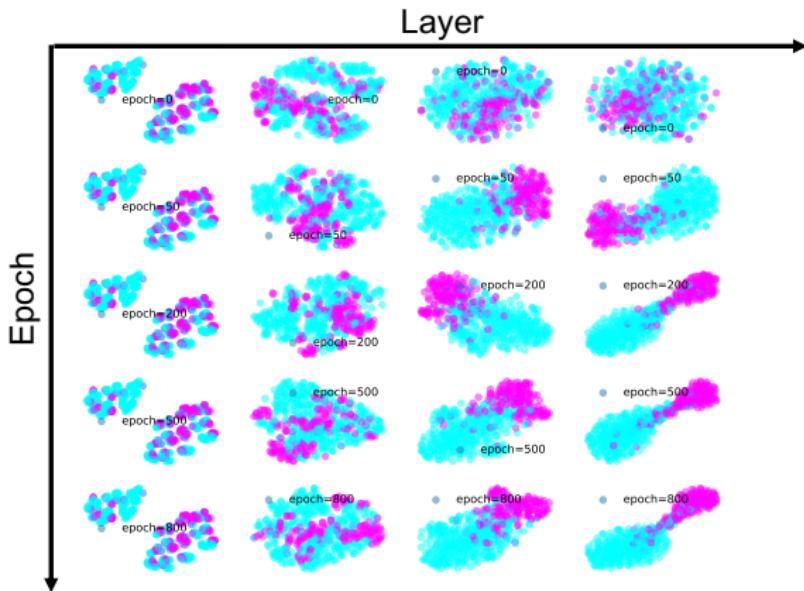
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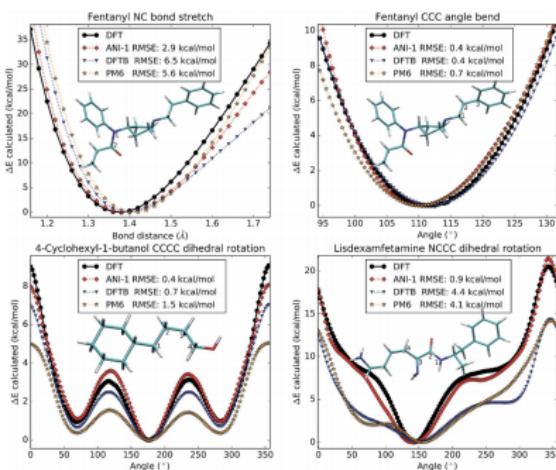
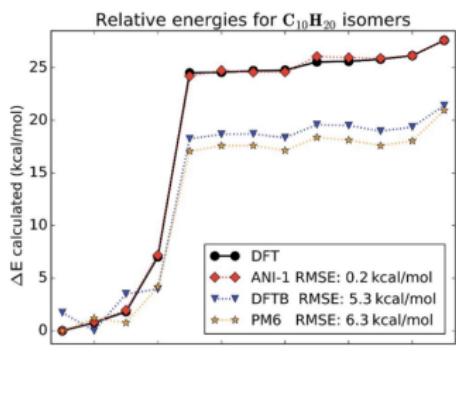
- 1 Neural network models provide model complexity ‘on tap’
- 2 Backpropagation allows easy access to derivatives
- 3 We can understand neural networks as automatic feature selection/transformation, followed by linear regression

# ANN example

jupyter notebook: [github.com/jpjanet/ML-chem-workshop/  
blob/master/notebooks/ANN.ipynb](https://github.com/jpjanet/ML-chem-workshop/blob/master/notebooks/ANN.ipynb)

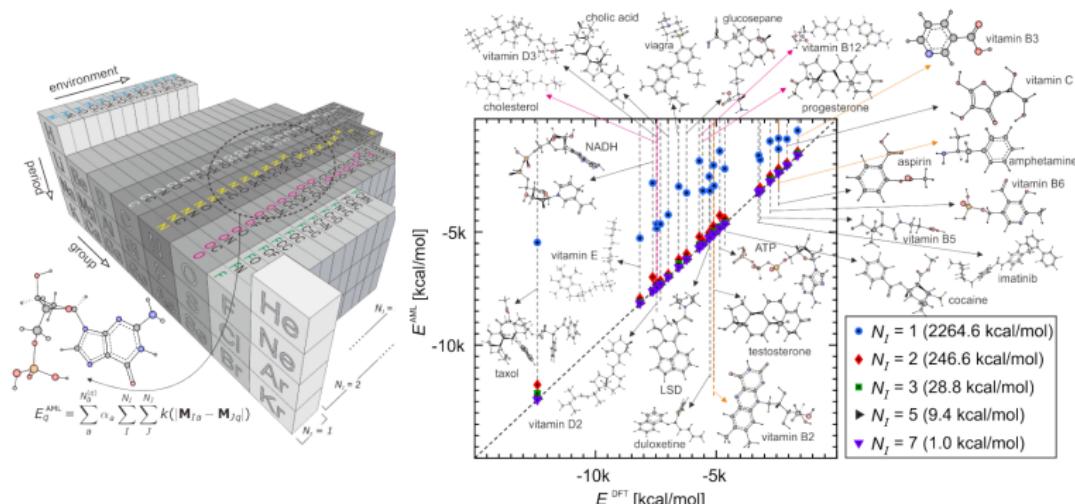
# Neural network potentials

Smith, J. S., Isayev, O., and Roitberg, A. E. ANI-1: an extensible neural network potential with DFT accuracy at force field computational cost. *Chem. Sci.*, 2017, 8, 3192-3203.



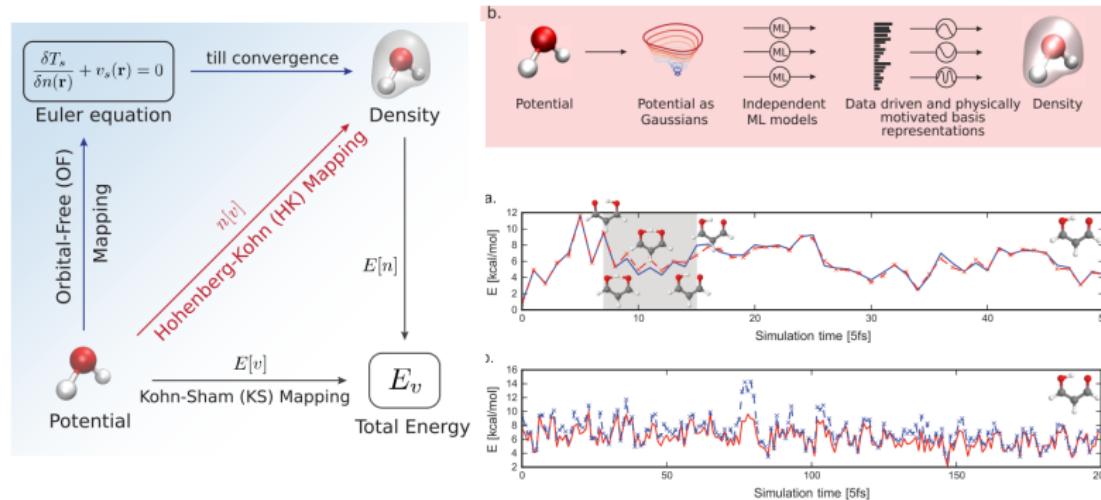
# Property predictions

Huang, B. & von Lilienfeld, O.A. *arXiv* 1707.04146, "The 'DNA' of chemistry: Scalable quantum machine learning with amons", 2017.



# Accelerating quantum chemistry

Bogojeski, M. *et al.*, Burke, K. and Müller, K.R. *arXiv* 1811.06255, Efficient prediction of 3D electron densities using machine learning, 2018.



# Controlling calculations on the fly

Chenru Duan et. al. *ChemRxiv* .7616009, "Learning from Failure: Predicting Electronic Structure Calculation Outcomes with Machine Learning Models", 2019.

