# Machine-Learning

An introduction

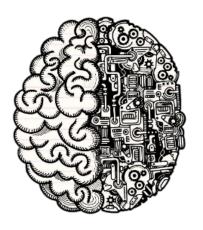
by: Estefany Suárez & Jake Vogel 29/06/2021

# Machine Learning (ML)

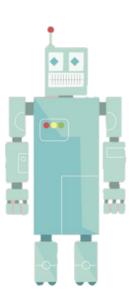
#### Learn from experience



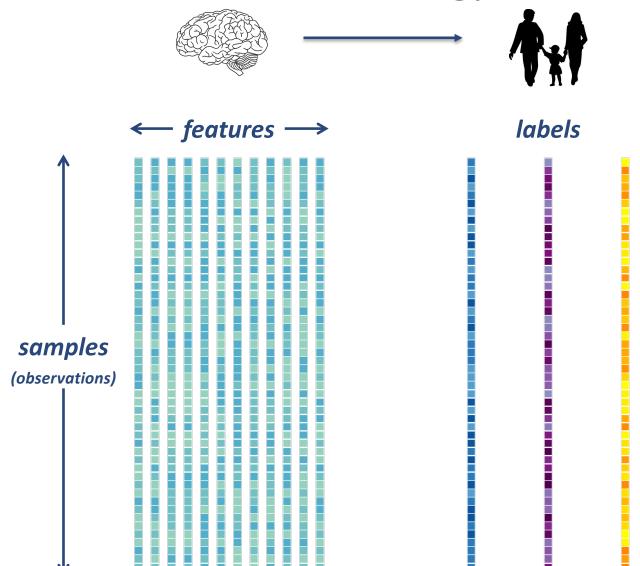




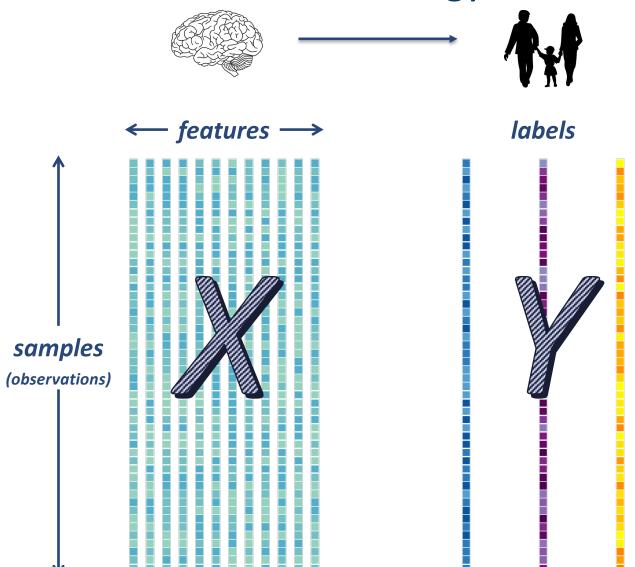
#### Follow instructions



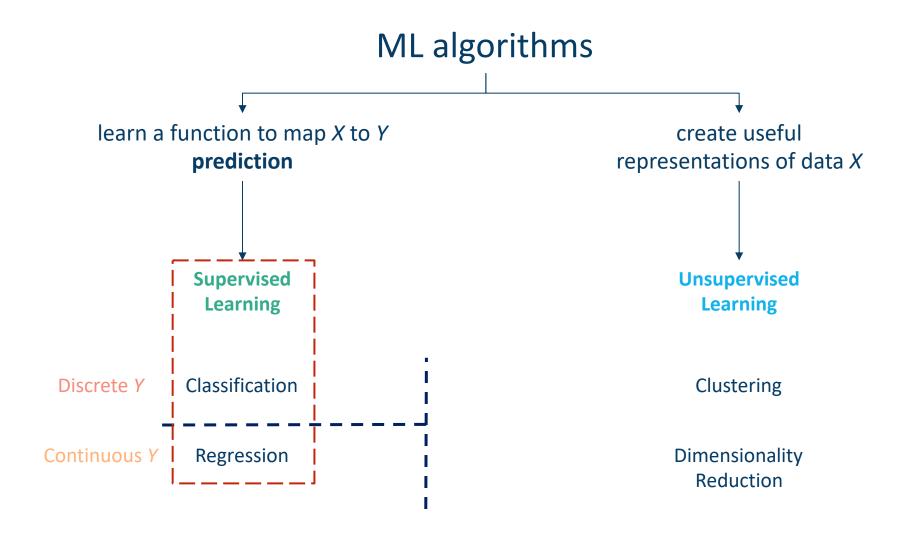
# **Terminology**



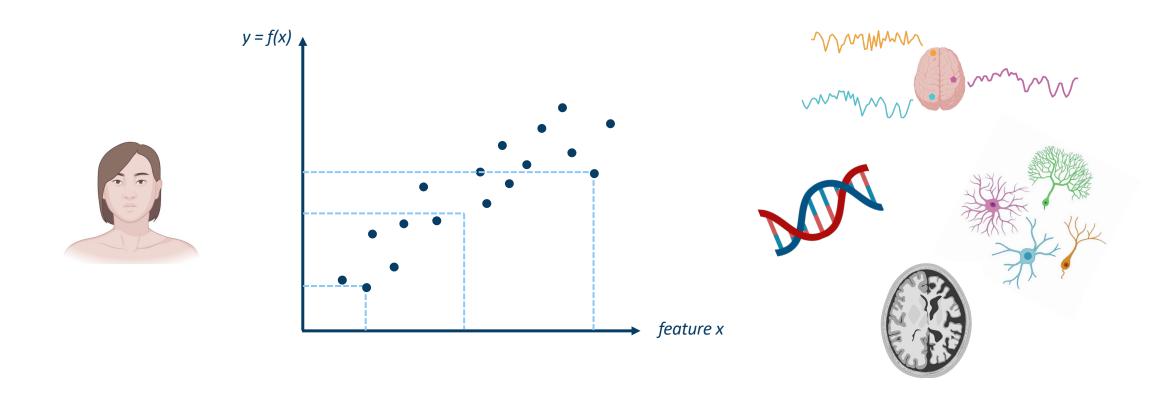
# Terminology



# ML algorithms learn a function to map X to Y create useful prediction representations of data X Unsupervised **Supervised Semi-supervised** Learning Learning Learning x, y $f: x \to y$ y = f(x)

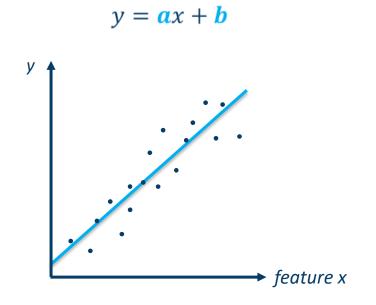


# Regression

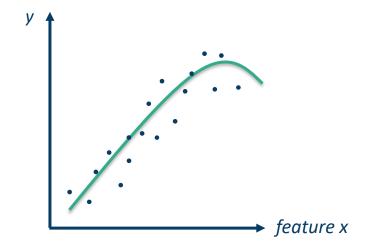


# Regression

$$f: x \to y$$
$$y = f(x)$$



$$y = ax^2 + bx + c$$



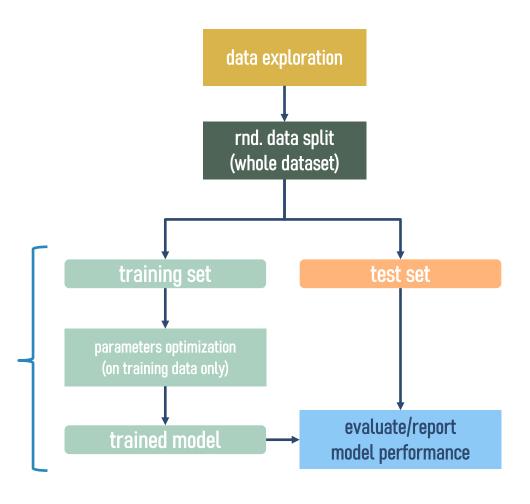
# Typical ML Pipeline

Why cross-validation (CV)?

- Avoid overfitting
- Improve model generalizability

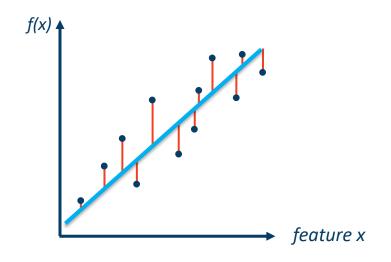
model

fitting



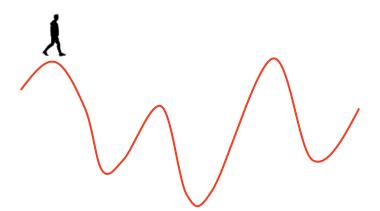
# ML basic ingredients

$$f: x \to y$$
$$y = f(x)$$





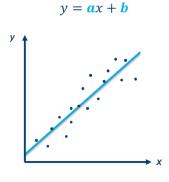
- 2 Loss function  $\longrightarrow MSE = \frac{1}{n} \sum_{i=1}^{n} (y_i \hat{y}_i)^2$
- Optimization algorithm gradient descent



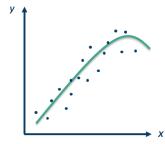
# ML basic ingredients

1 Model (estimator) selection

linear regression



$$y = ax^2 + bx + c$$



polynomial regression

multi-layer perceptron (ANN)



2 Loss (cost) function

mean squared error =  $\frac{1}{n}\sum_{i=1}^{n}(y_i - \hat{y}_i)^2$ 

mean absolute error =  $\frac{1}{n}\sum_{i=1}^{n}|y_i - \hat{y}_i|$ 

 $\max \text{error} = \max(|y_i - \hat{y_i}|)$ 

explained var. =  $1 - \frac{Var(y - \hat{y})}{Var(y)}$ 

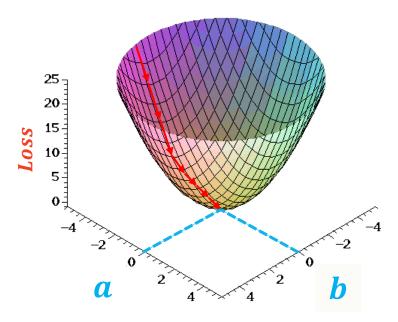
$$R^{2} = 1 - \frac{\sum_{i=1}^{n} (y_{i} - \widehat{y}_{i})^{2}}{\sum_{i=1}^{n} (y_{i} - \overline{y})}$$

# ML basic ingredients

Optimization algorithm for model parameters!

$$y = ax + b$$

$$Loss = f(\mathbf{a}, \mathbf{b})$$



### **Gradient Descent**

Training data

 $y_{\text{train}}$ 

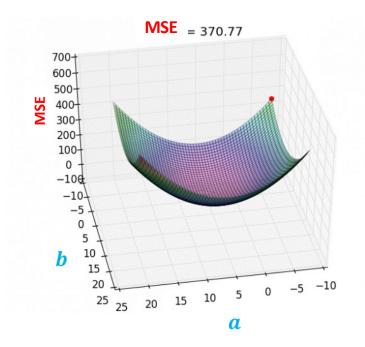
 $\chi_{\rm train}$ 

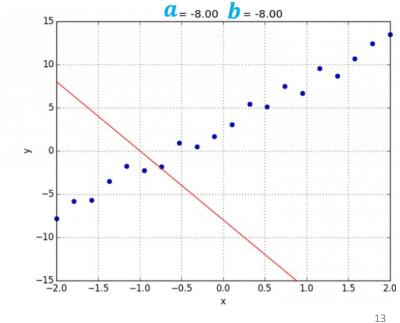
Initialize model parameters (a,b) randomly iterate between:

- 1) Estimate output
- $\widehat{\mathbf{y}}_i = a\mathbf{x}_i + \mathbf{b}$
- 2) Compute *loss*

MSE = 
$$\frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2$$

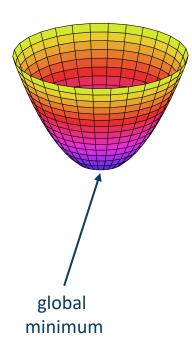
- 3) Compute the gradient
- 4) Update parameters (a,b)



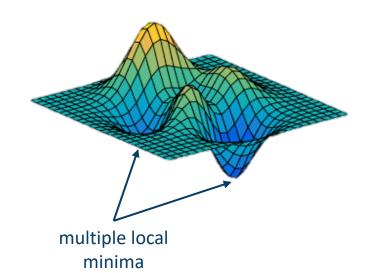


# There is a problem though ...

#### simple loss function



convoluted loss function



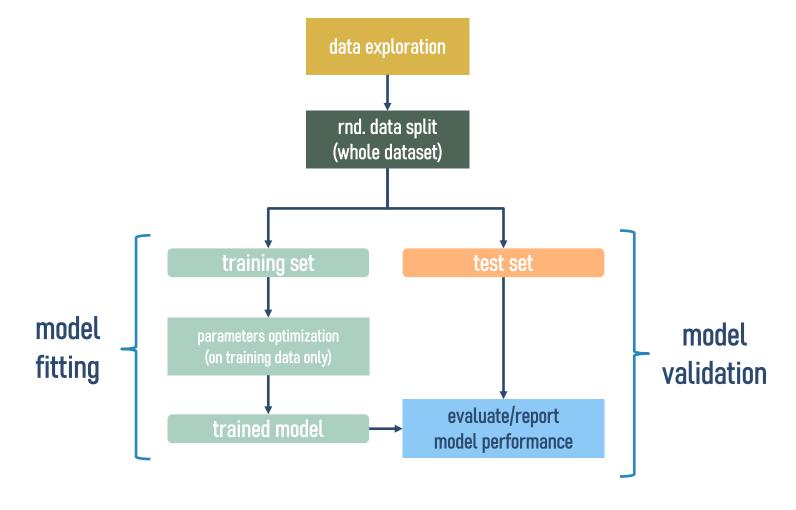
Me WANTS THE DATA



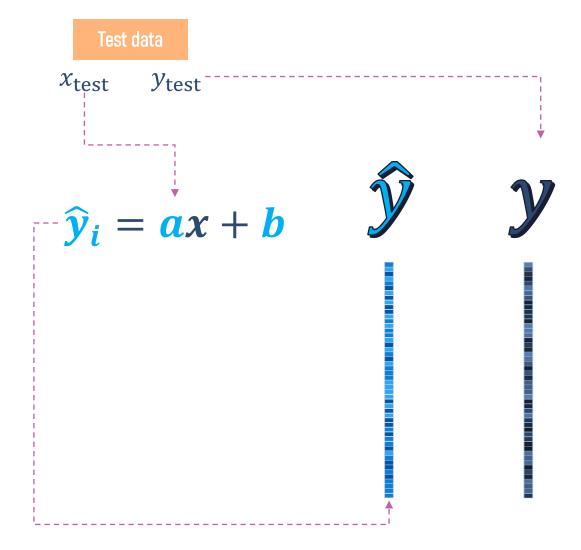




# Typical ML Pipeline



### **Model Validation**



#### score function

mean squared error = 
$$\frac{1}{n}\sum_{i=1}^{n}(y_i - \hat{y}_i)^2$$

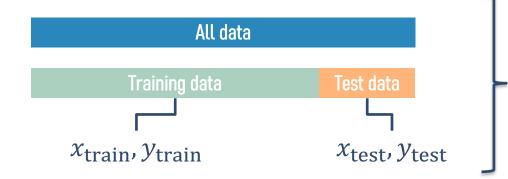
mean absolute error = 
$$\frac{1}{n}\sum_{i=1}^{n}|y_i-\widehat{y}_i|$$

$$\max \operatorname{error} = \max(|y_i - \widehat{y_i}|)$$

explained var. = 
$$1 - \frac{Var(y - \hat{y})}{Var(y)}$$

$$R^{2} = 1 - \frac{\sum_{i=1}^{n} (y_{i} - \widehat{y}_{i})^{2}}{\sum_{i=1}^{n} (y_{i} - \overline{y})}$$

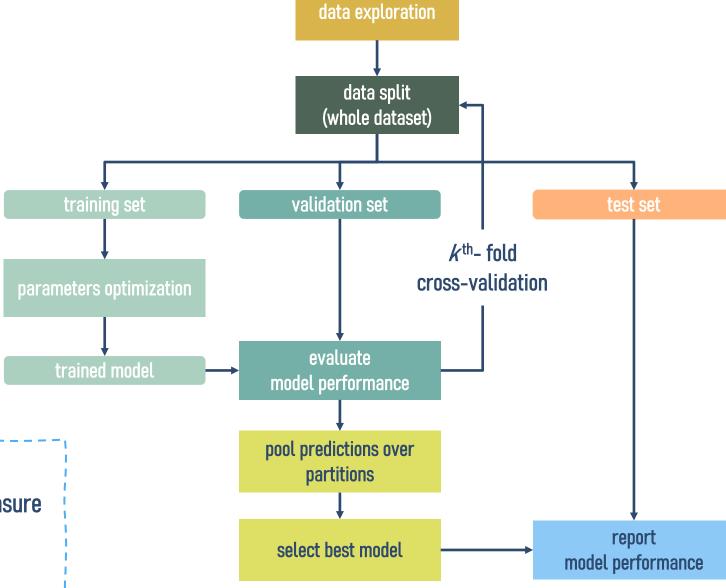
### We have yet another problem ...



BUT ... What if your outlier samples/observations happen to be in the test set?

This would greatly affect the generalizability of your model!!!

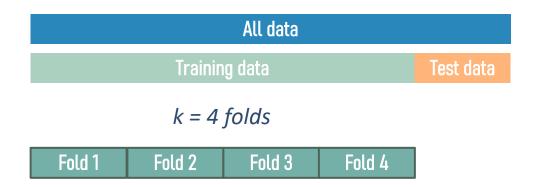
# Typical ML Pipeline

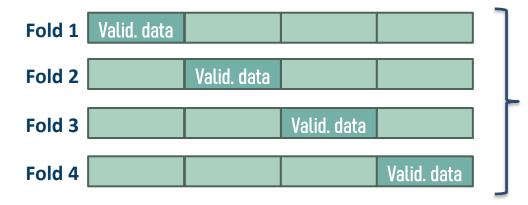


### Why *k<sup>th</sup>*-fold CV?

- Provide a more accurate performance measure
  - [average, standard deviation, range]
- Model selection
- *Hyper*-parameter tuning

# [k-fold] Cross-Validation (CV)





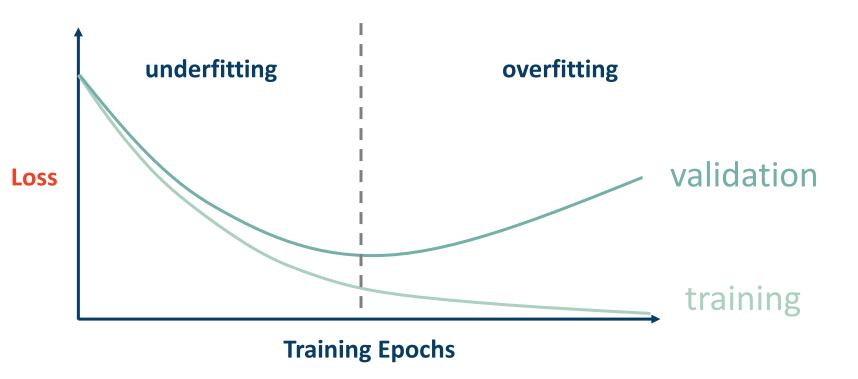
- 1. For each fold:
  - I. Train model parameters with training set
  - II. Evaluate training with <u>validation</u> set
- Based on <u>validation</u> set, select *optimal* model and hyper-parameters
- 3. Only at the very, very **END**, report error on <u>test</u> set

# **Overfitting**

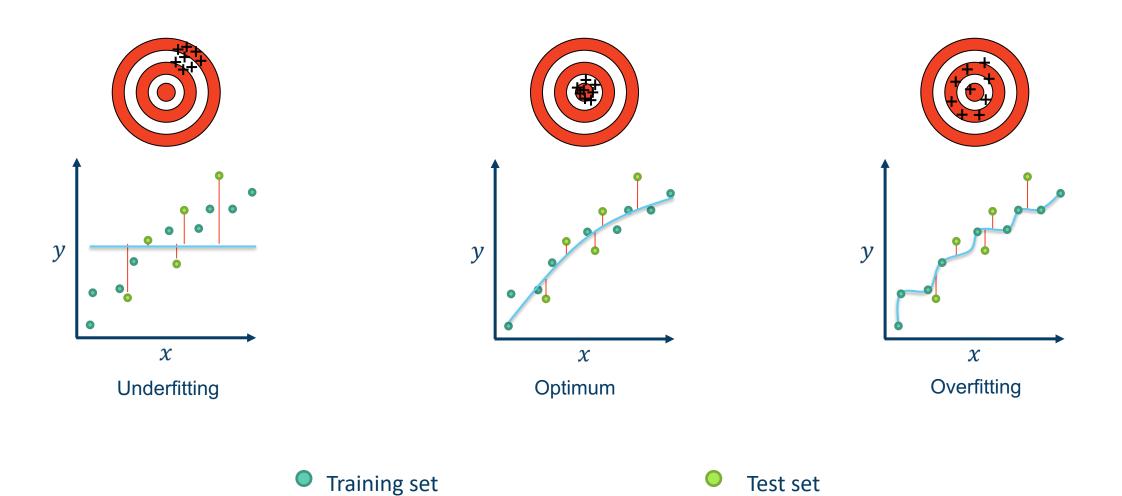
k<sup>th</sup>-fold

#### **How many TE?**



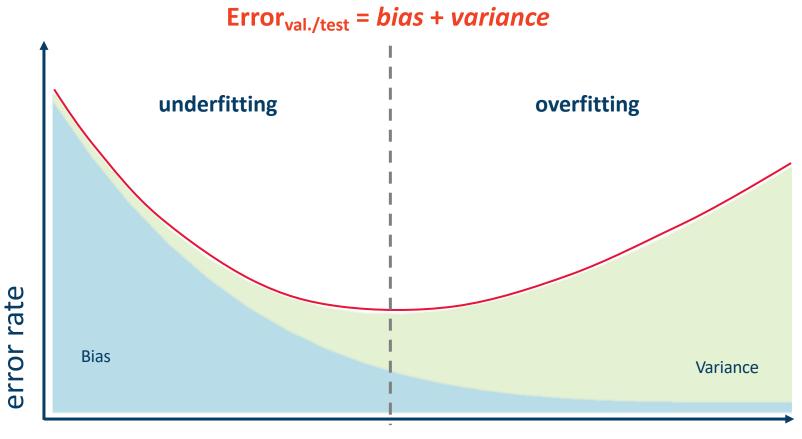


### **Bias-Variance Trade-Off**



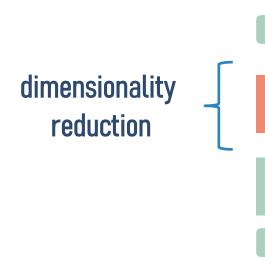
Test set

### Bias-Variance Trade-Off



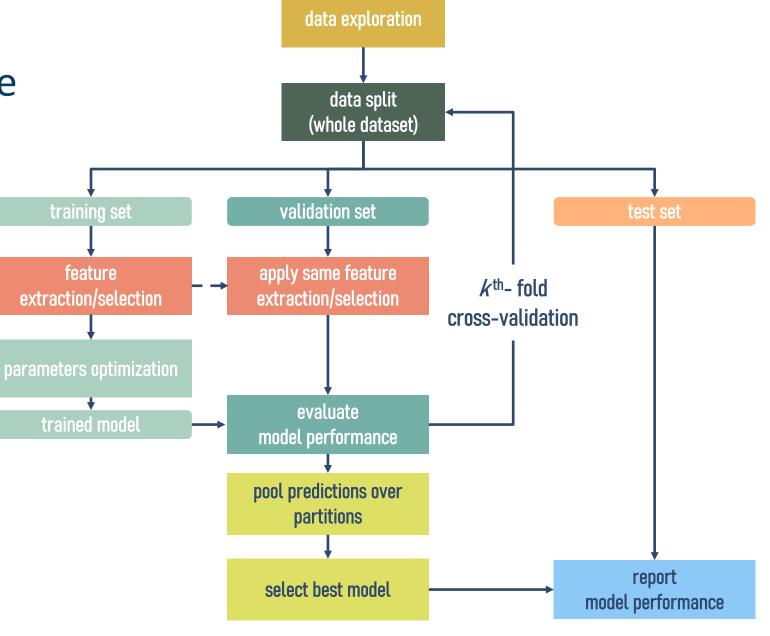
model complexity (~ number of features) learning epochs

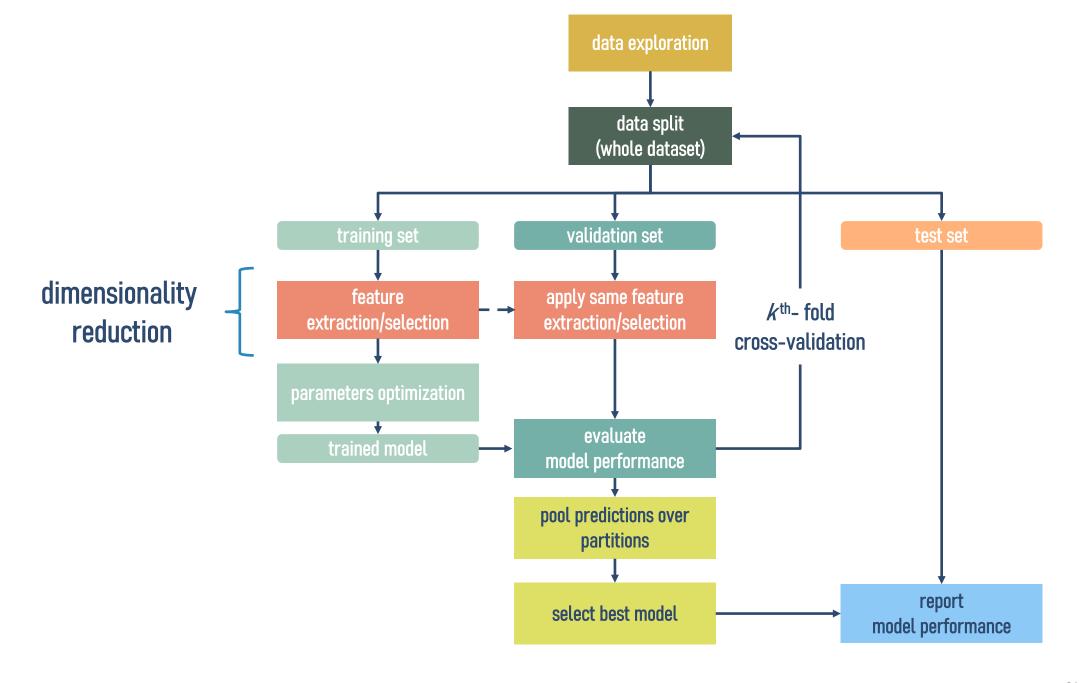
# Typical ML Pipeline



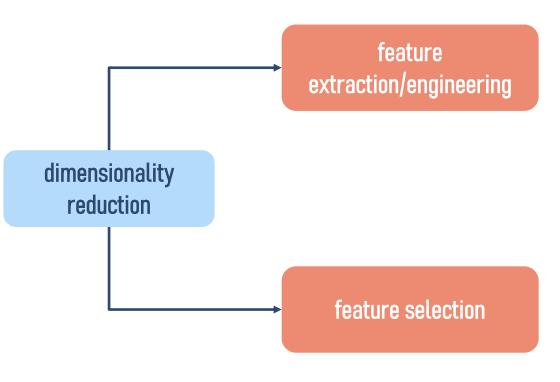
#### WHY DR?

- Curse of dimensionality (more features than samples)
- Intrinsic dimension may actually be small (redundante data)
- Extract "salient" features
- Remove noisy features
- VISUALIZATION!!!!





### Tip 1: Dimensionality Reduction



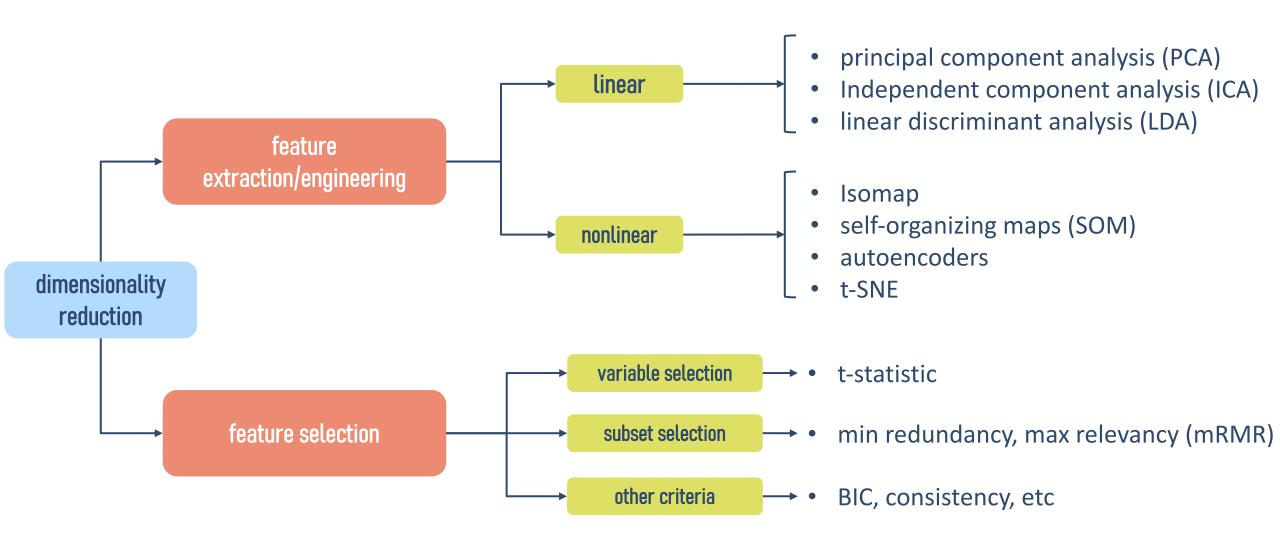
- Compact representation of the data
- Maps input features into a lower dimensional space

e.g. If features are 
$$X = [x_1, x_2, x_3, x_4]$$
  
then  $Z = T(X) = [c_1x_1 + c_2x_2, x_3 * x_4]$ 

- Selection of a subset of input features
- Features are still in original space

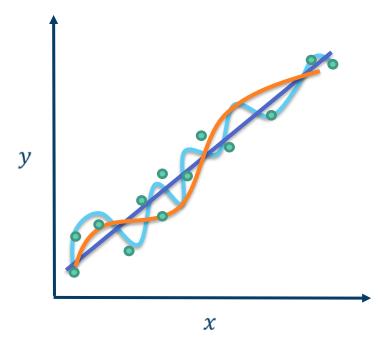
e.g. 
$$Z = S(X) = [x_2, x_3]$$

### Tip 1: Dimensionality Reduction



### Tip 2: Regularization

$$y = \beta_0 + \beta_1 x + \beta_2 x^2 + \dots + \beta_p x^p$$



Penalties on the *LOSS* function to prevent overfitting!

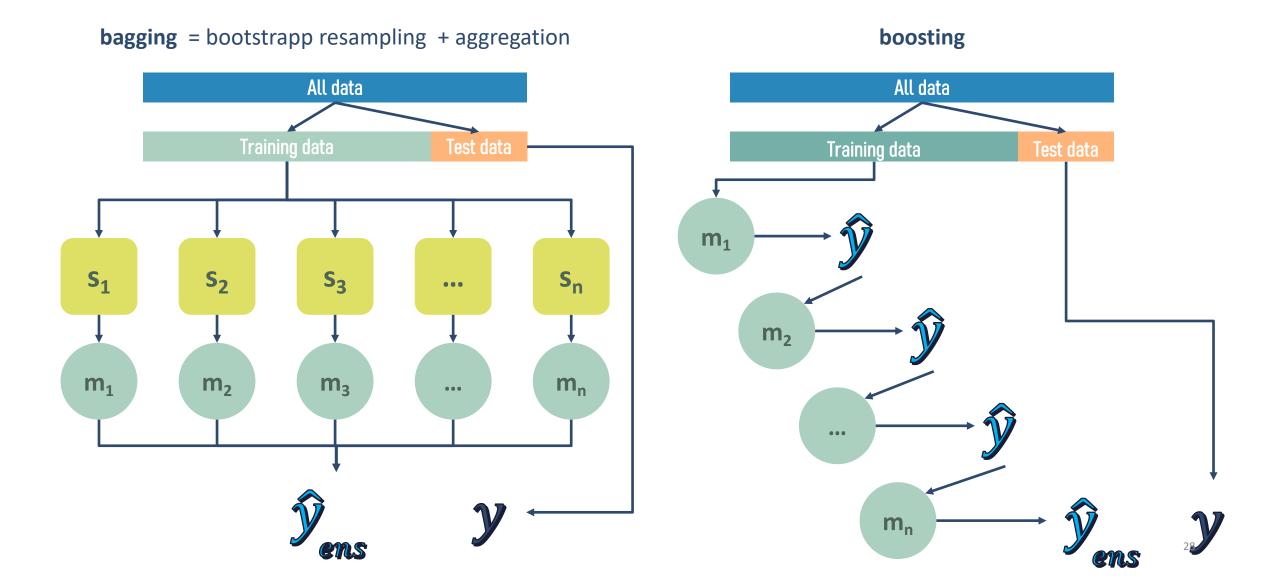
1) L1/Lasso: constrains parameters to be sparse

$$MSE = \sum_{i=1}^{n} \left( y_i - \sum_{j=1}^{p} x_{ij} \beta_j \right)^2 + \lambda \sum_{j=1}^{p} |\beta_j|$$

2) L2/Ridge: constrains parameters to be small

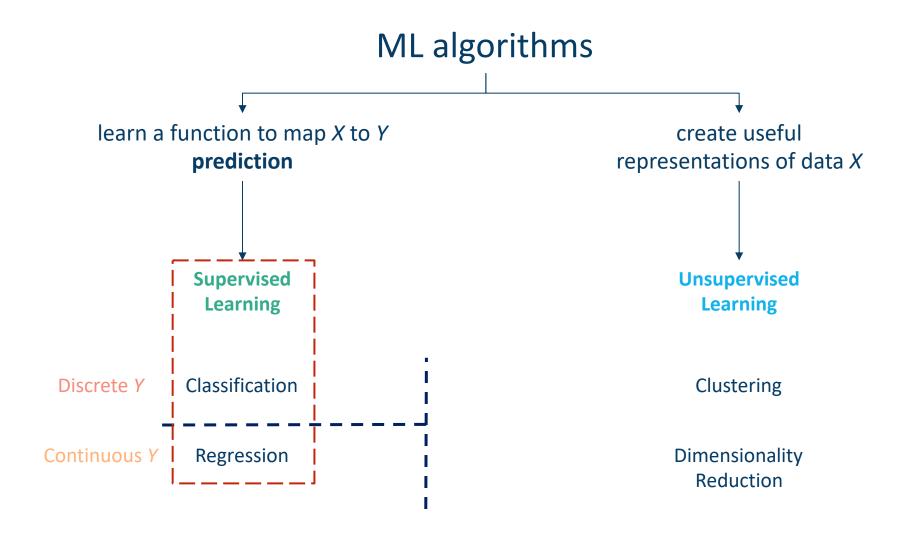
$$MSE = \sum_{i=1}^{n} \left( y_i - \sum_{j=1}^{p} x_{ij} \beta_j \right)^2 + \lambda \sum_{j=1}^{p} \beta_j^2$$

# Tip 3: Ensemble Methods

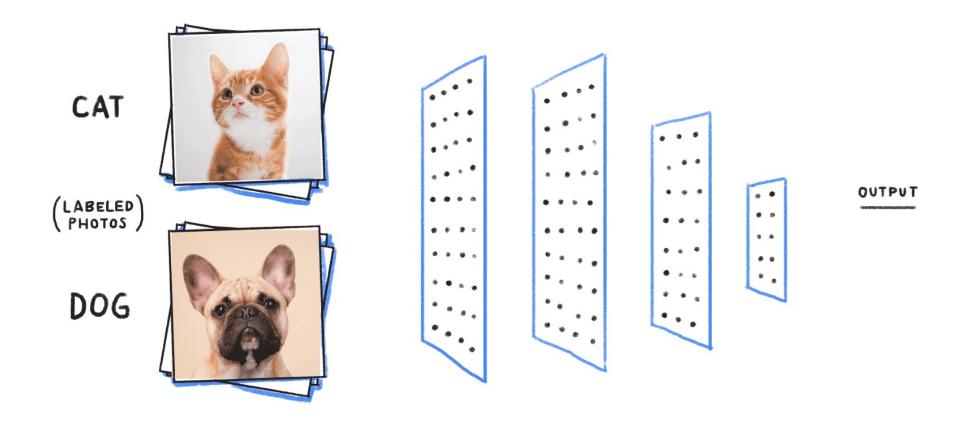


# Sources of Bias

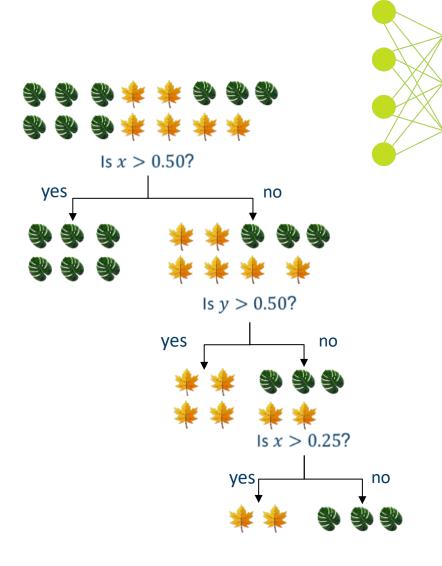
Type	DO NOT	8	Sexy name	DO	
<i>k</i> -hacking	Try many <b>k's in k-fold CV</b> (or different training %) and report only the best		<i>k</i> -hacking	Pick k=10, repeat It many times (>200 or as many as possible!), and report the full distribution (NOT boxplots!)	
<i>metric</i> -hacking	Try different performetrics (e.g., accurate, e.g., arcurate, e.g., accurate, e	uracy, F1, etc.) and	<i>m</i> -hacking	appropriate a metric for the AUC for	the most nd recognized problem (e.g., r binary cation)
feature/dataset-hacking	Try subsets of feat subsamples of da but report only t	itaset(s),	<i>d</i> -hacking	everything: a	report on Il analyses on tasets



### Classification



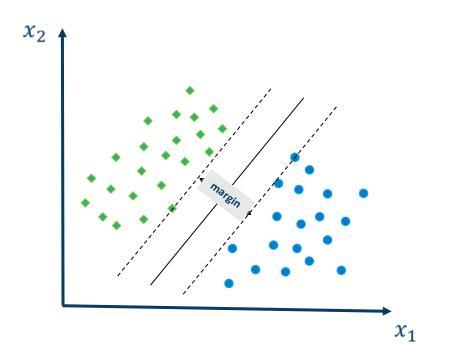
### Classification



Support Vector Machine (SVM)

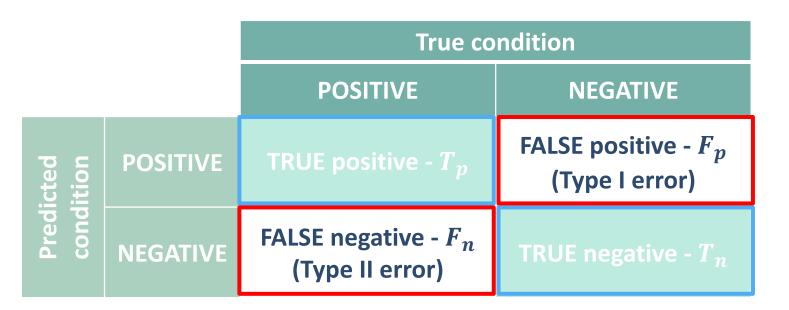
- Artificial Neural Networks
- Logistic regression
- Decision Trees
- Random Forests

probabilistic classifiers



### Performance Metrics – Binary Classification

#### confusion matrix



#### score function

$$accuracy = \frac{T_p + T_n}{T_p + T_n + F_p + F_n}$$

$$precision = \frac{T_p}{T_p + F_p}$$

$$recall = \frac{T_p}{T_p + F_n}$$

$$accuracy = \frac{T_p}{T_p + F_n}$$

$$Appropriate when classes are imbalanced!$$

$$F1 \ score = \frac{2T_p}{2T_p + F_p + F_n}$$

# Multiclass Prediction (≠ Multilabel Prediction)

		True class		
		CAT	DOG	BIRD
Predicted class	CAT	13	0	0
	DOG	0	10	6
	BIRD	0	0	9

- To extend a binary metric to multiclass problems, the data is treated as a collection of binary problems, one for each class.
- The binary metric is then averaged across the set of classes, each of which may be useful in some scenario.

### Performance Metrics – Binary Classification

		True condition		
		POSITIVE	NEGATIVE	
icted	POSITIVE TRUE positive - T		FALSE positive - $F_p$ (Type I error)	
Predicted condition	NEGATIVE	FALSE negative - $F_n$ (Type II error)	TRUE negative - $T_n$	

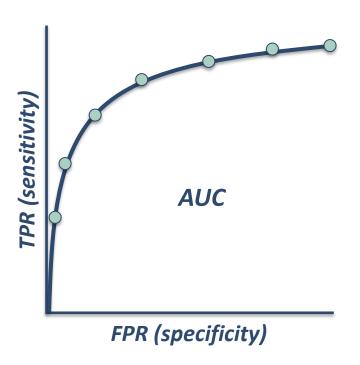
Sensitivity (or recall) 
$$TPR = \frac{T_p}{P} = \frac{T_p}{T_p + F_n}$$
  $FPR = \frac{F_p}{N} = \frac{F_p}{T_n + F_p}$  Specificity (or selectivity)

$$FNR = \frac{F_n}{P} = \frac{F_n}{T_p + F_n}$$
  $TNR = \frac{T_n}{N} = \frac{T_n}{T_n + F_p}$ 

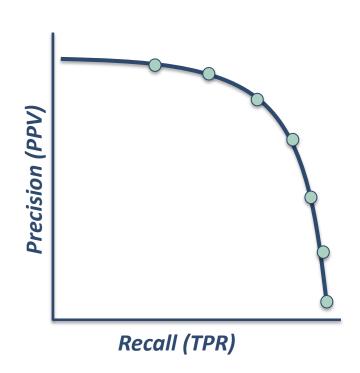
### Performance Metrics – Binary Classification

(probabilistic classifiers)

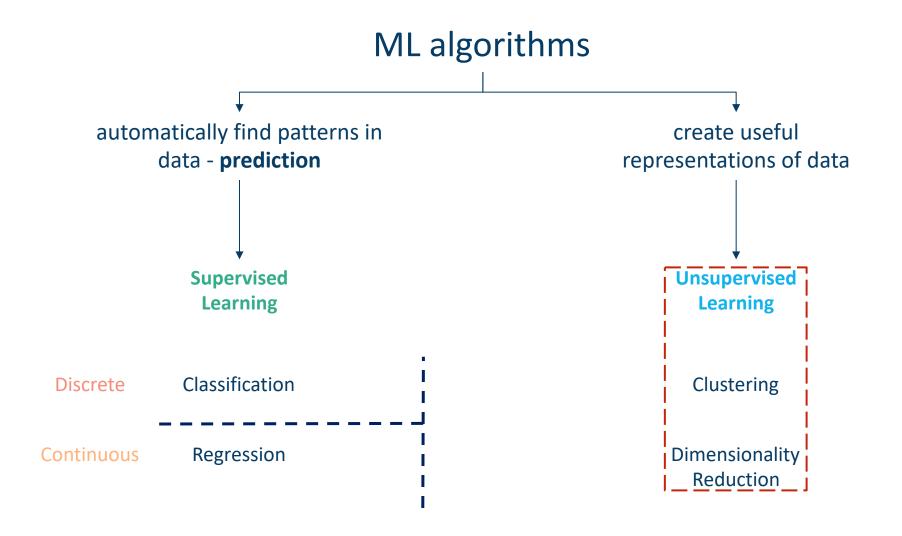
#### **ROC** curve



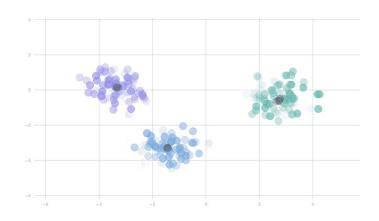
#### precision-recall curve



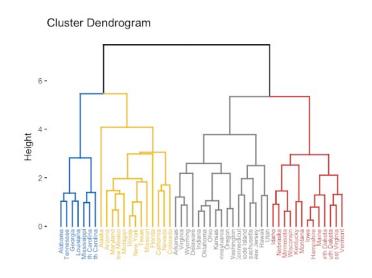
Appropriate when classes are imbalanced!



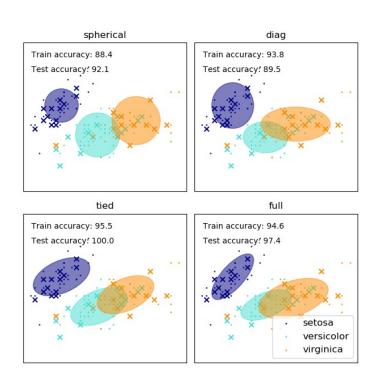
### Clustering



*K*-means



Hierarchical clustering



Gaussian mixture models

# Inference vs Prediction – Linear Model $\mathbf{Y} = \boldsymbol{\beta} \mathbf{X} + \boldsymbol{\epsilon}$

#### statistical inference

#### pattern recognition

#### Goal:

• Identify significant contributing variables (statistical null-hypothesis testing, *p*-values)

#### **Uses:**

- Scientific discovery. Ideal to uncover characteristics or true properties of the biological processes of the studied phenomenon.
- Useful to judge the individual relevance of each quantitative measure in impacting the response of interest.

#### Goal:

 Identify most predictive variable sets (out-of-sample prediction performance)

#### **Uses:**

- Pragmatic forecasting of biological processes.
- Tends to concern less regarding the data-generating process.

### Diagnosing Features (≈ Interpretability)

#### Local Interpretable Model-Agnostic Explanations [LIME]

