

Computational
Neuroscience
Laboratory

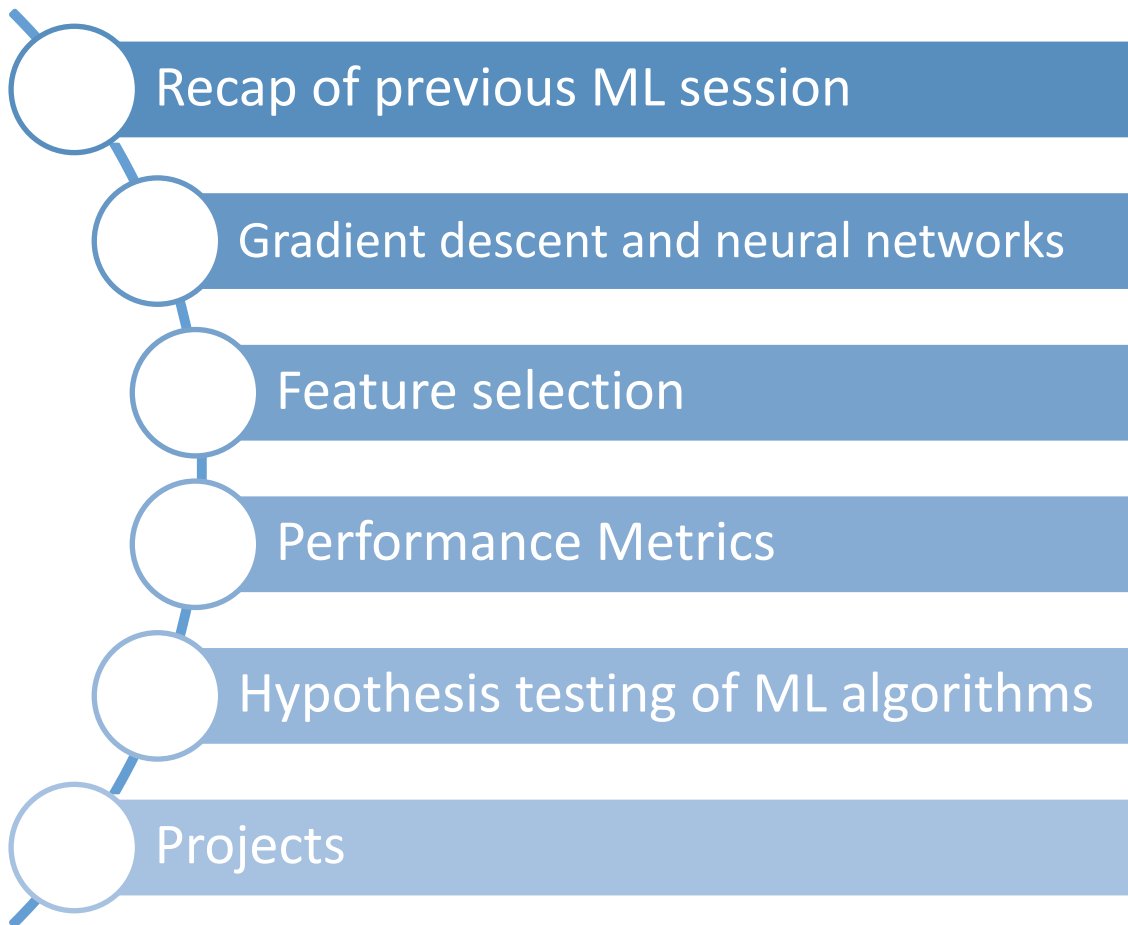
Machine Learning II

Autumn 2023

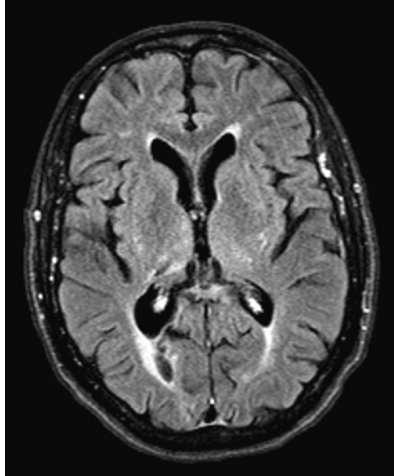
Session 6 – 10/12/2023



Today...



Classification: A core task

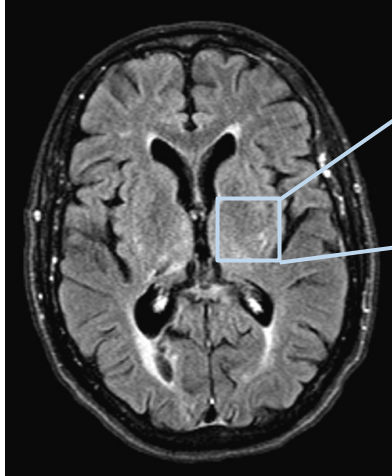


(assume given set of discrete labels)
{AD, MCI, PD, ASD, ...}



Alzheimer's disease

The Problem: Semantic Gap



[105	112	100	111	104	99	106	99	96	103	112	119	104	97	93	87]
[91	98	102	106	104	79	98	103	99	105	123	136	110	105	94	85]
[76	85	90	105	128	105	87	96	95	99	115	112	106	103	99	85]
[99	81	81	93	120	131	127	100	95	98	102	99	96	93	101	94]
[106	91	61	64	69	91	88	65	101	107	109	90	75	84	96	95]
[114	100	85	55	55	69	64	54	64	87	112	129	98	74	84	91]
[133	137	147	103	65	81	88	65	52	54	74	84	102	93	85	82]
[128	137	144	140	109	95	86	70	62	65	63	63	60	73	86	101]
[125	133	140	137	119	121	117	94	65	79	80	65	54	64	72	98]
[127	125	131	147	133	127	126	131	111	96	89	75	61	64	72	84]
[115	114	109	123	150	148	131	118	113	109	100	92	74	65	72	78]
[89	93	90	97	100	147	131	118	113	114	113	109	106	95	77	80]
[63	77	86	81	77	79	102	123	117	115	117	125	125	130	115	87]
[62	65	82	89	78	71	80	101	124	126	119	101	107	114	131	119]
[63	65	75	88	89	71	62	81	120	138	135	105	81	90	110	118]
[87	65	71	87	106	95	69	45	76	130	126	107	92	94	105	112]
[110	97	82	86	117	123	116	66	41	51	95	93	89	95	102	107]
[164	146	112	80	82	120	124	104	76	48	45	66	88	101	102	109]
[157	170	157	120	93	86	114	132	112	97	69	55	70	82	99	94]
[130	120	134	161	139	100	109	118	121	134	114	87	65	53	69	86]
[120	112	96	117	150	144	120	115	104	107	102	93	87	81	72	79]
[123	107	96	86	83	112	153	149	122	109	104	75	80	107	112	99]
[122	121	102	80	82	86	94	117	145	148	153	102	58	78	92	107]
[122	164	148	103	71	56	78	83	93	103	119	139	102	61	69	84]

What the computer sees

An image is just a big grid of numbers between $[0, 255]$:

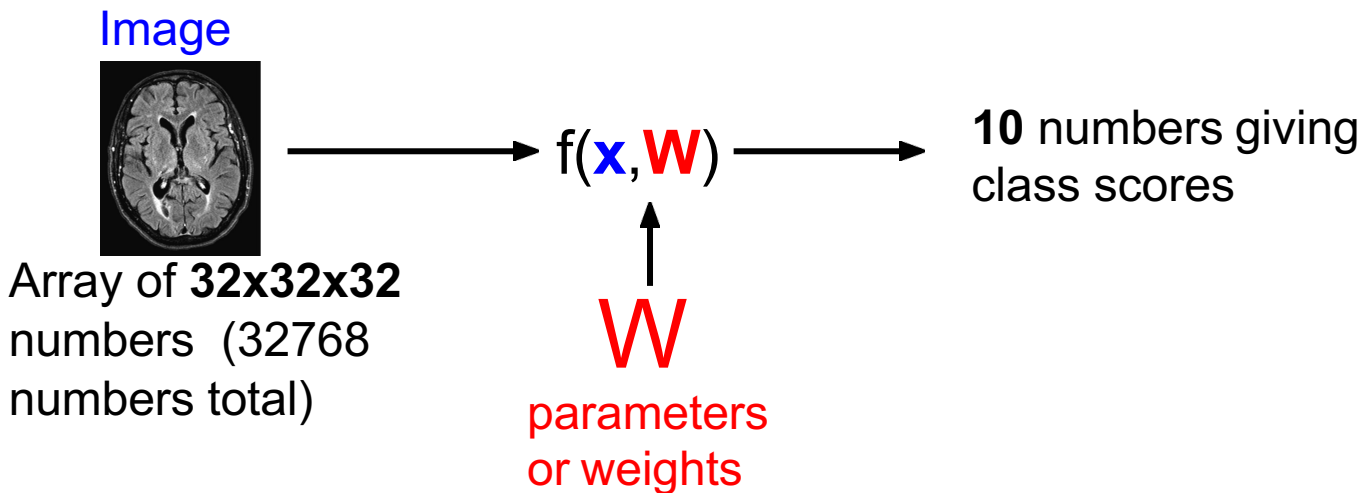
e.g. $800 \times 600 \times 3$
(3 channels RGB)

Different Challenges

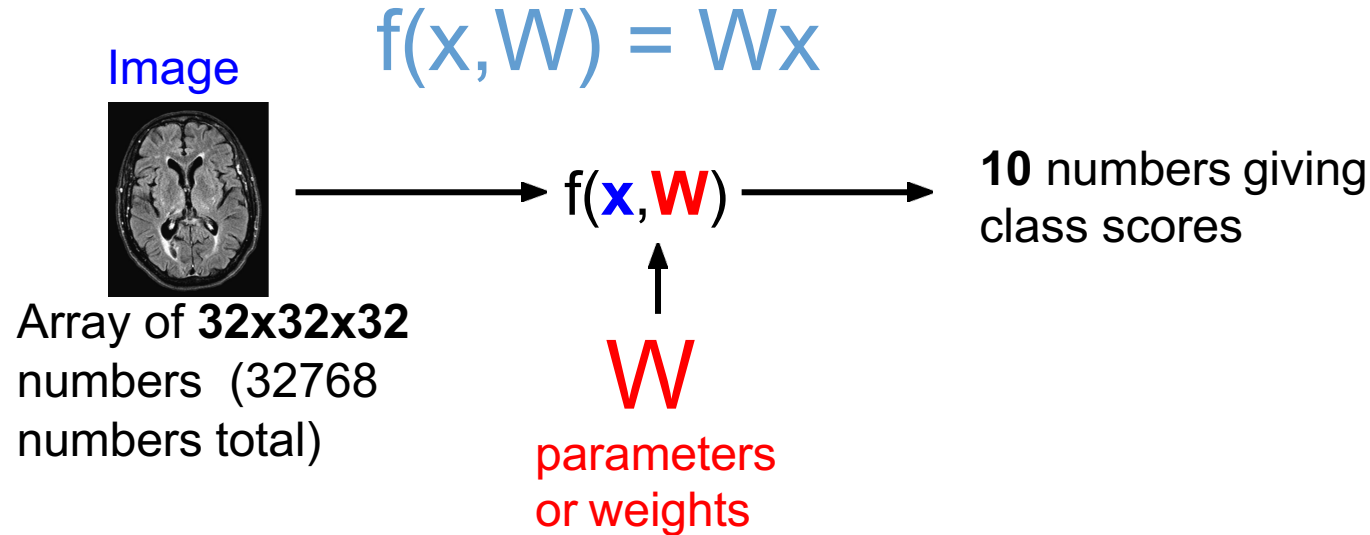
- 3D Volumetric data / 4D Volume+Time / ...
- Scanner and protocol difference
- Intrasubject variabilities of brain structure and function
- Brain disorders/diseases have different pathologies and development patterns
- Data imperfection (motion, blur, missing data, ...)

Linear Classification

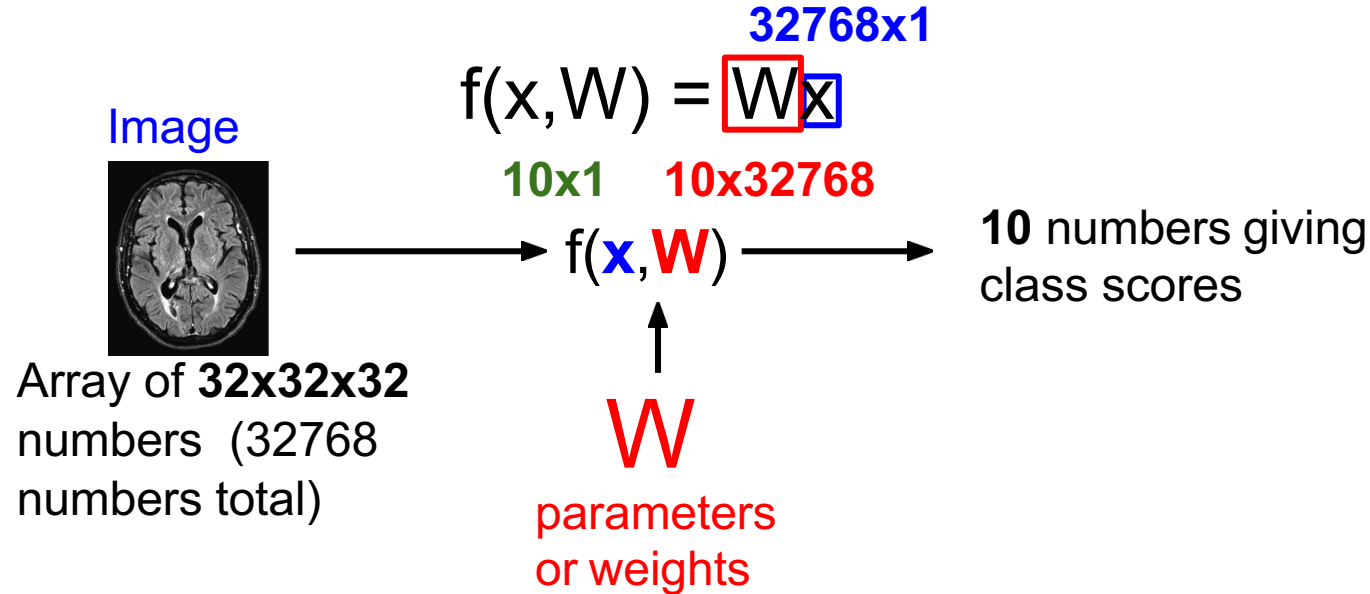
Parametric Approach



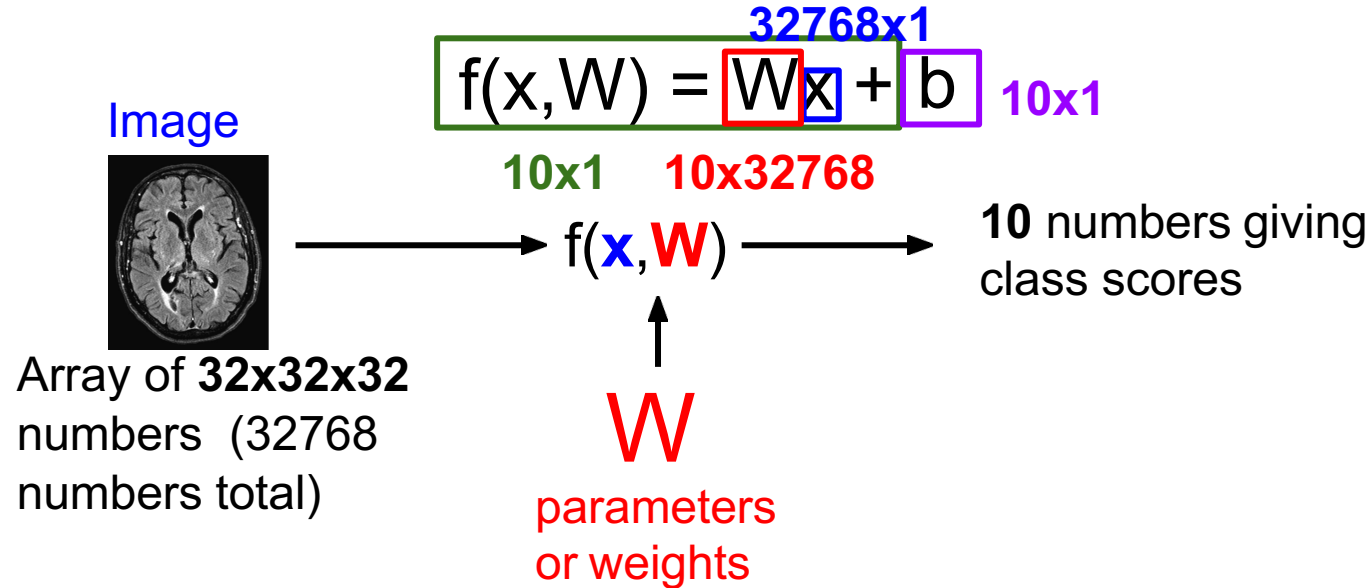
Parametric Approach: Linear Classifier



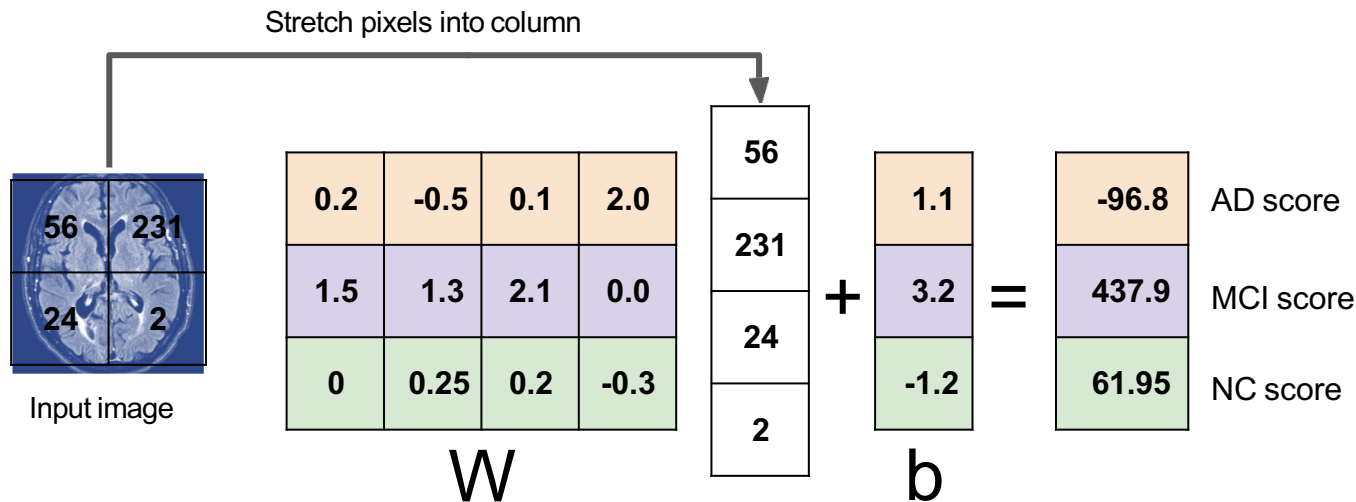
Parametric Approach: Linear Classifier



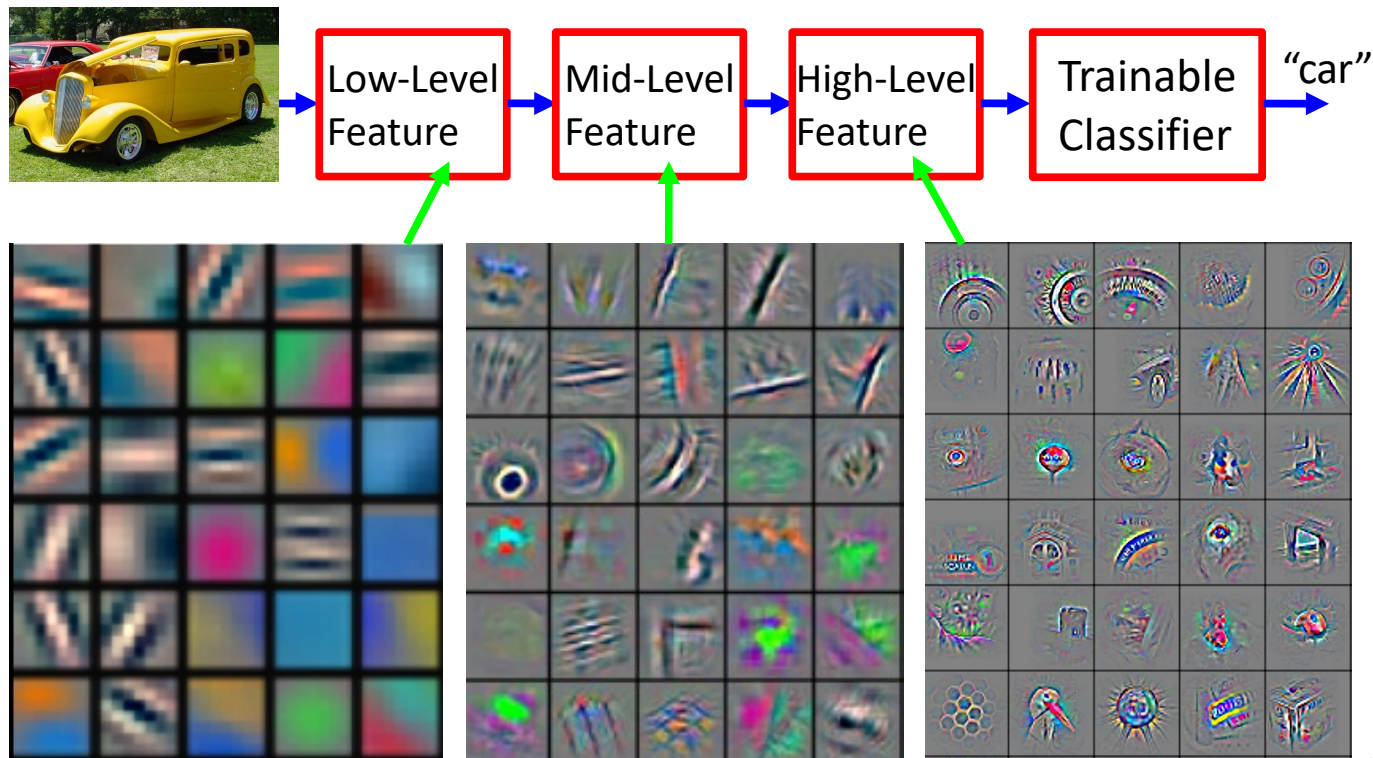
Parametric Approach: Linear Classifier



Example with an image with 4 pixels, and 3 classes (AD/MCI/NC)

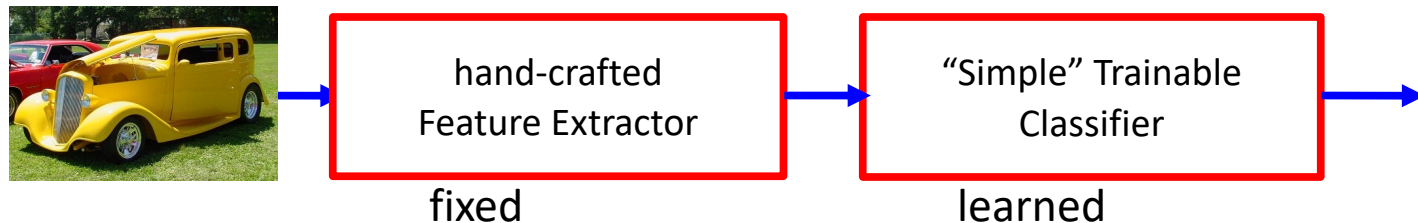


Deep Learning = Hierarchical Compositionality

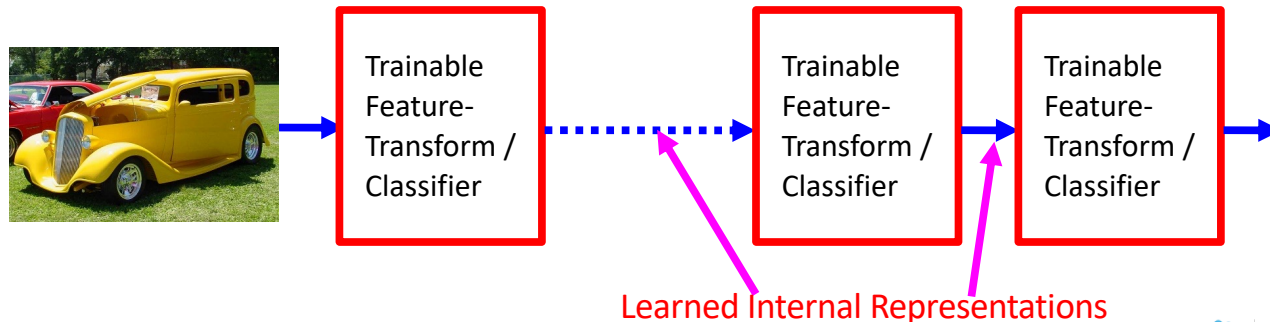


“Shallow” vs. Deep Learning

- “Shallow” models

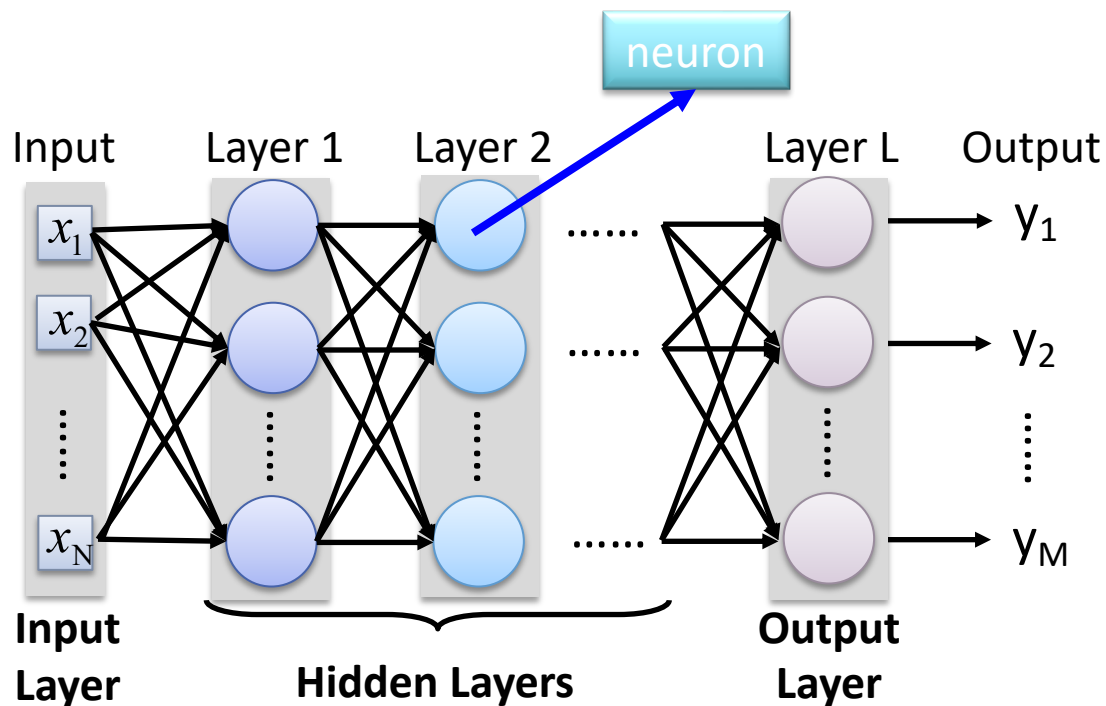


- Deep models



Learned Internal Representations

Neural Network



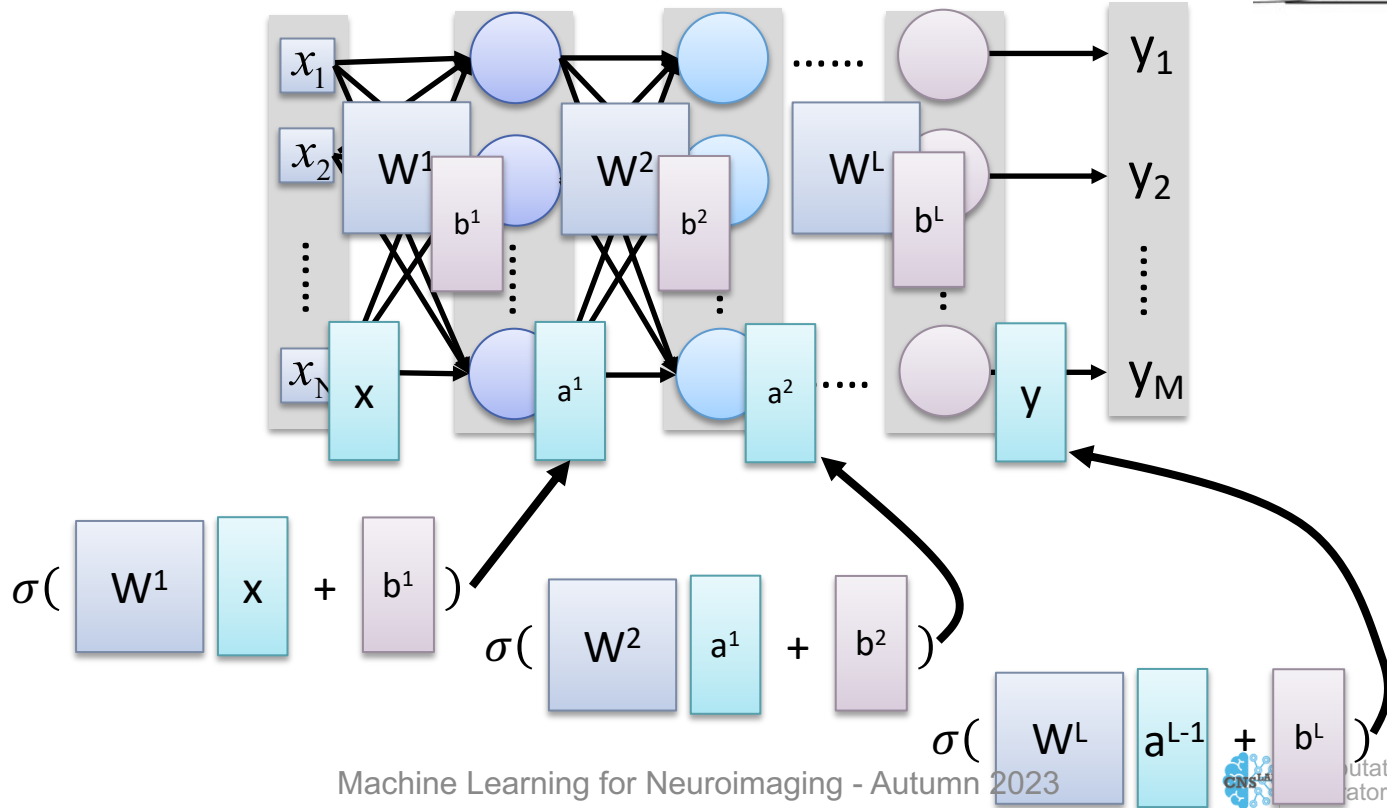
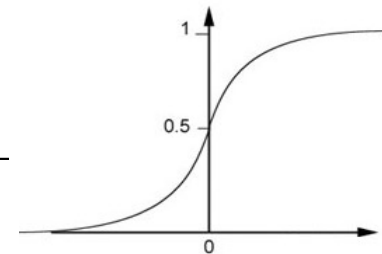
Deep means many hidden layers

Machine Learning for Neuroimaging - Autumn 2023

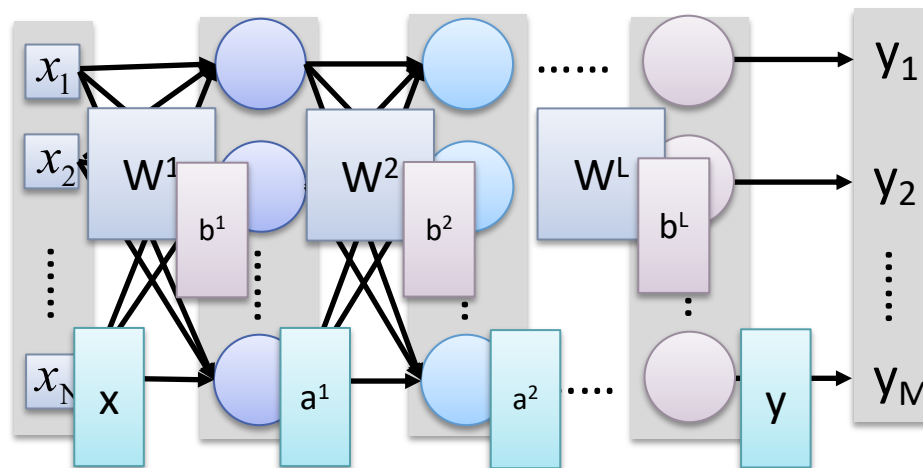
Neural Network

Sigmoid
Function

$$\sigma(z) = \frac{1}{1 + e^{-z}}$$



Neural Network



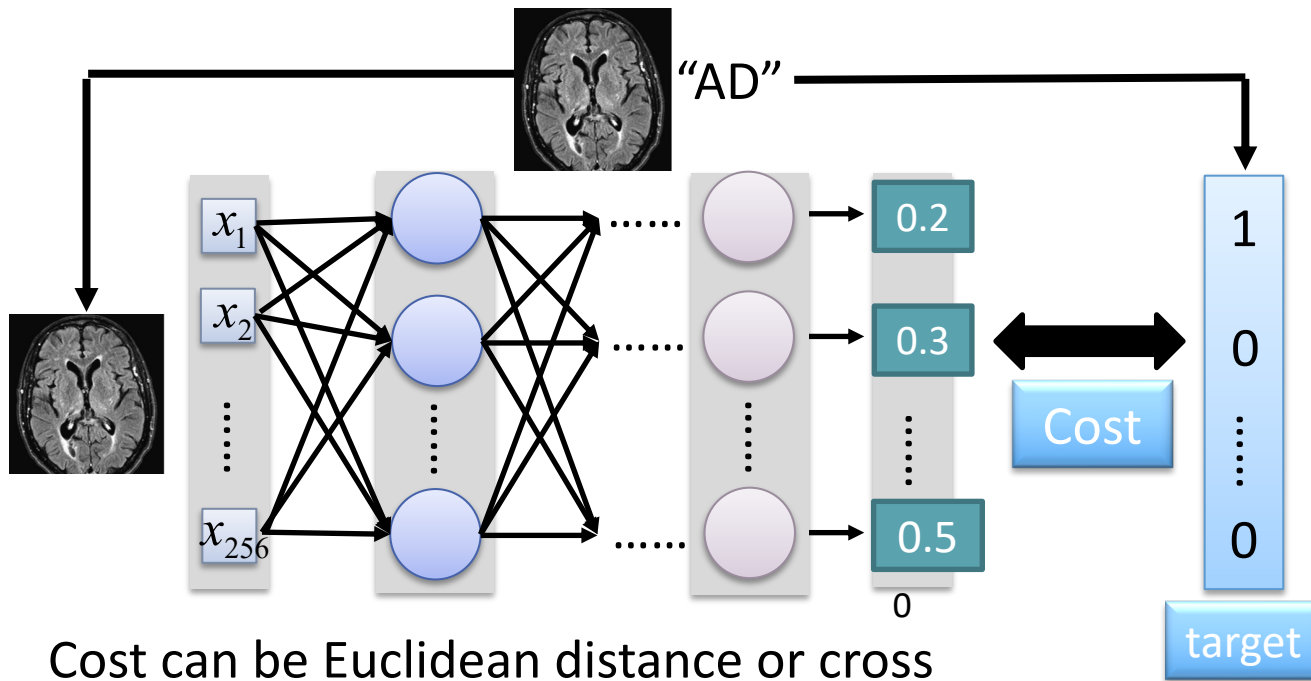
$$y = f(x)$$

Using parallel computing techniques to speed up matrix operation

$$= \sigma(W^L \dots \sigma(W^2 \sigma(W^1 x + b^1) + b^2) \dots + b^L)$$

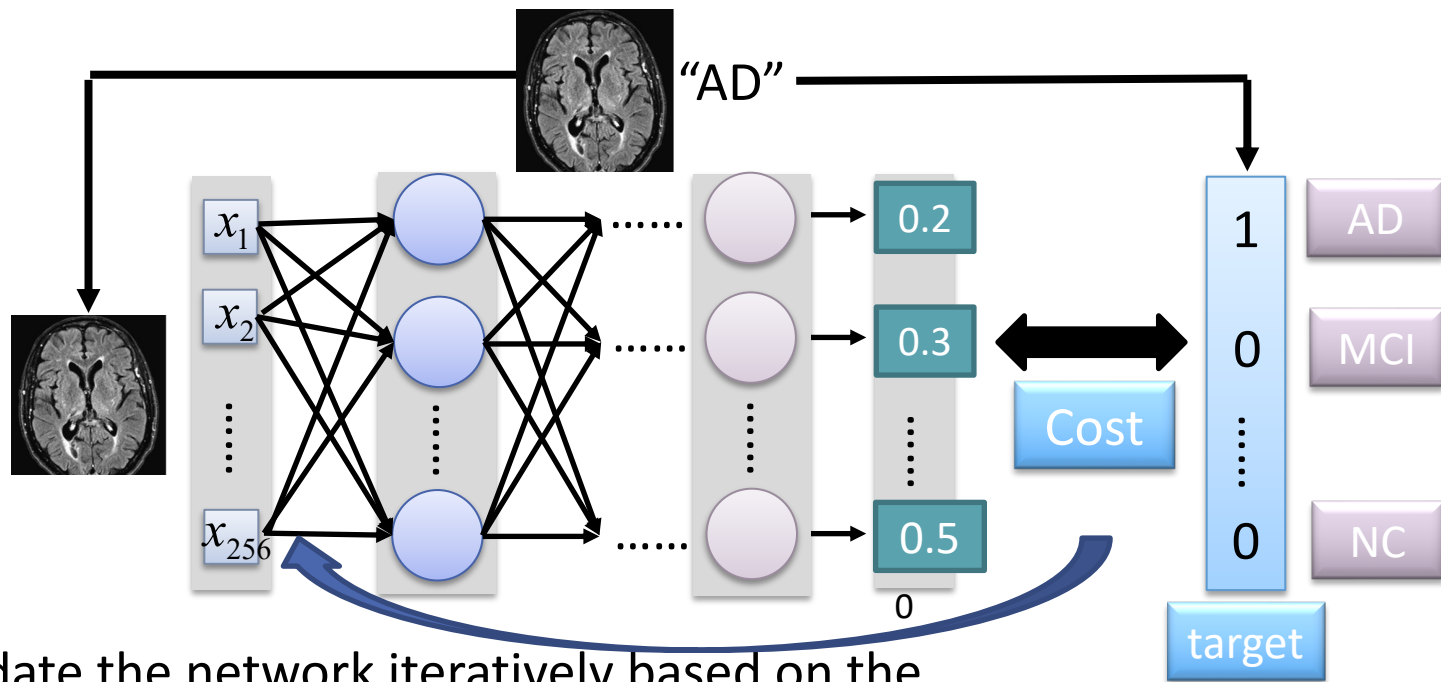
Cost

Given a set of network parameters, each example has a cost value.



Cost can be Euclidean distance or cross entropy of the network output and target

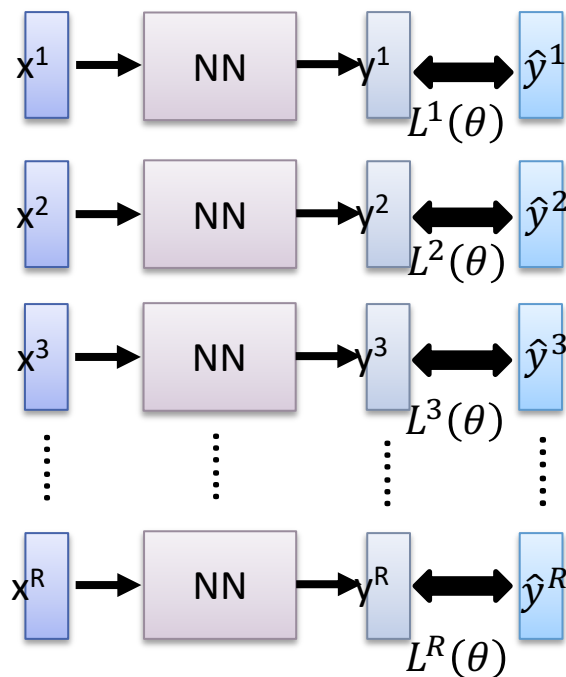
Training



Update the network iteratively based on the gradient of the cost until the network converges

Total Cost

For all training data ...



Total Cost:

$$C(\theta) = \sum_{r=1}^R L^r(\theta)$$

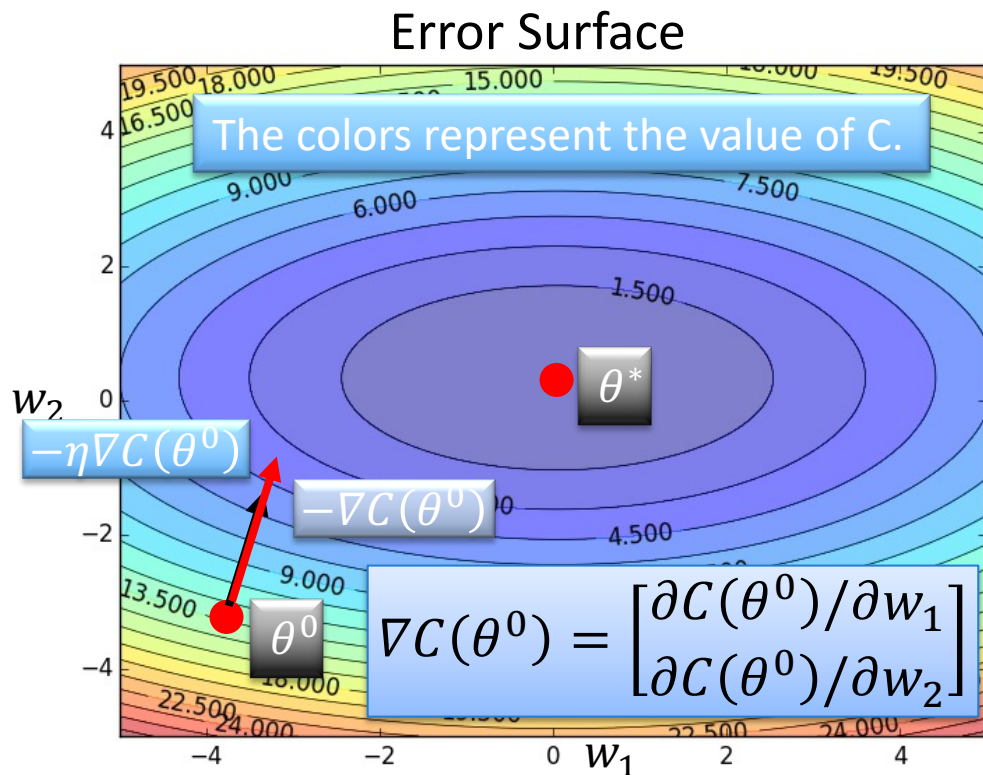
How bad the network parameters θ is on this task

Find the network parameters θ^* that minimize this value

Gradient Descent

Assume there are only two parameters w_1 and w_2 in a network.

$$\theta = \{w_1, w_2\}$$



Randomly pick a starting point θ^0

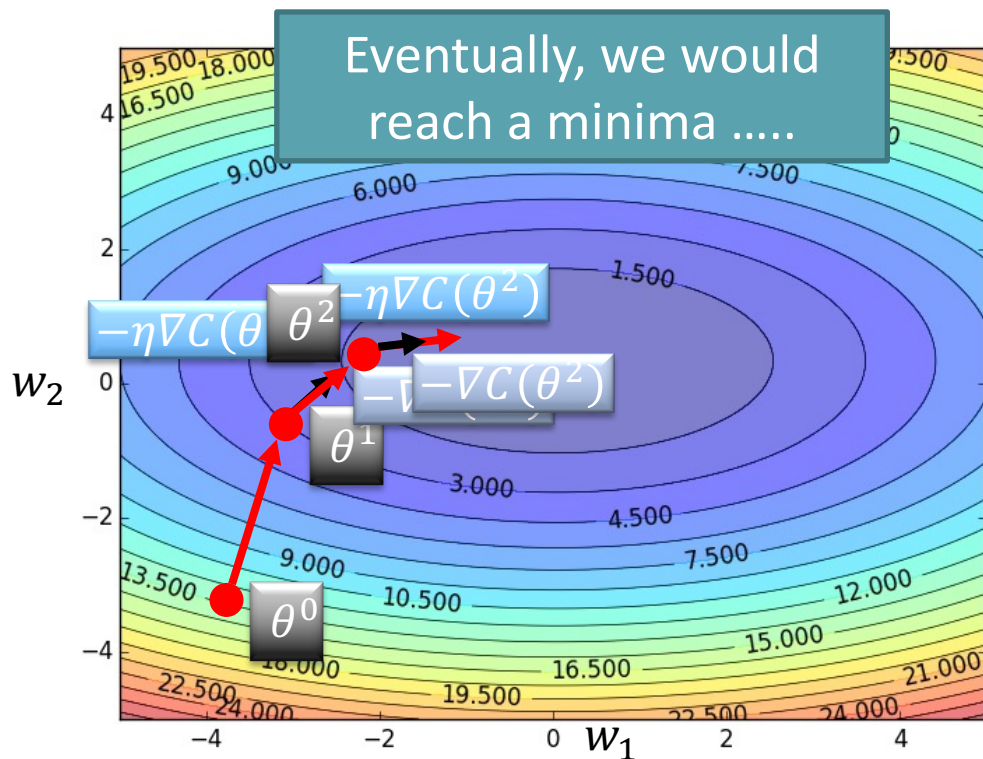
Compute the negative gradient at θ^0

➡ $-\nabla C(\theta^0)$

Times the learning rate η

➡ $-\eta \nabla C(\theta^0)$

Gradient Descent



Randomly pick a starting point θ^0

Compute the negative gradient at θ^0

$\rightarrow -\nabla C(\theta^0)$

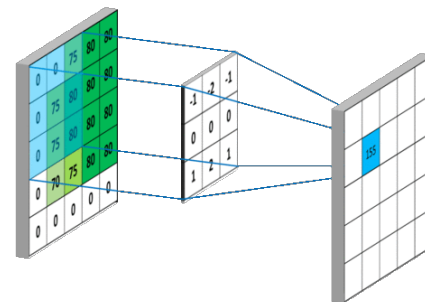
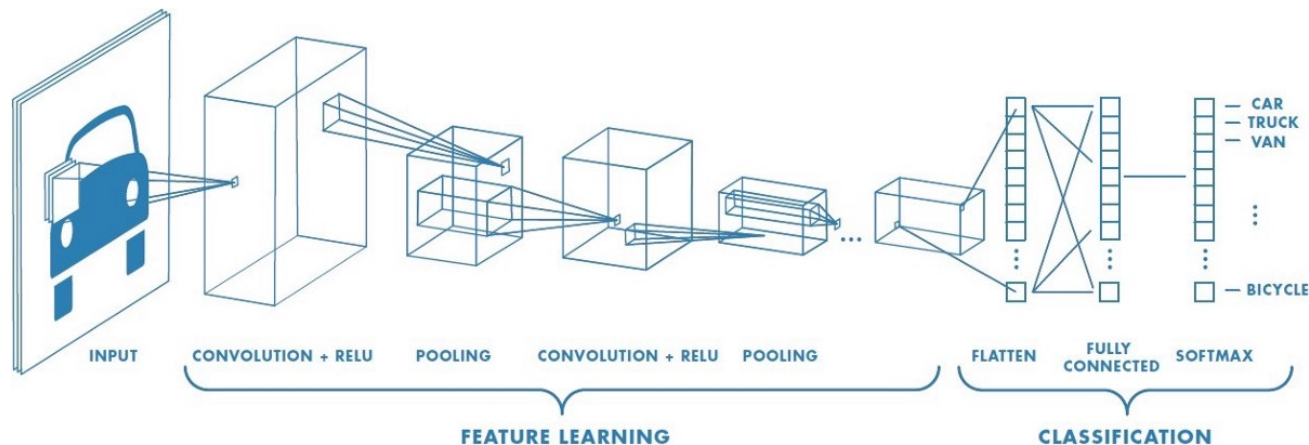
Times the learning rate η

$\rightarrow -\eta \nabla C(\theta^0)$

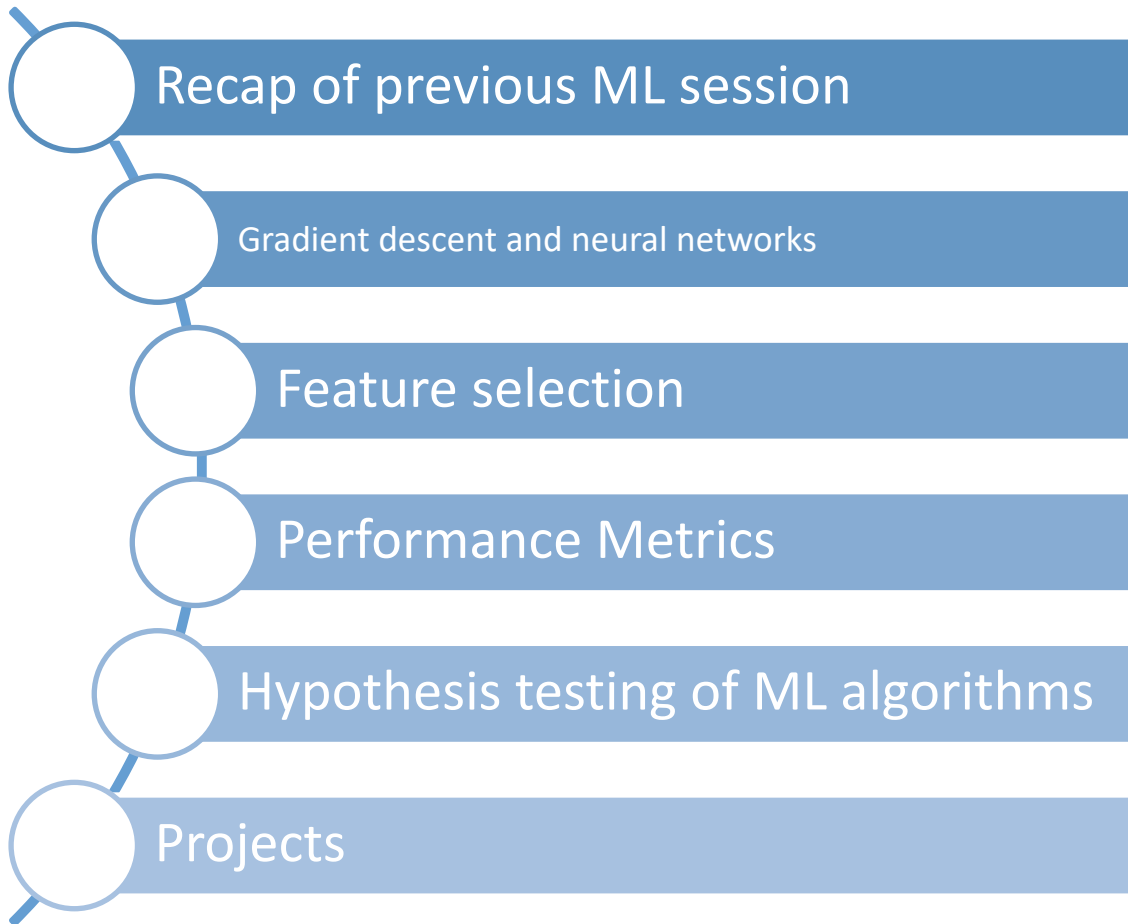
Reading assignment

■ Convolutional Neural Networks (CNN)

- <https://towardsdatascience.com/a-comprehensive-guide-to-convolutional-neural-networks-the-eli5-way-3bd2b1164a53>
- <https://www.kaggle.com/code/shivamb/3d-convolutions-understanding-use-case>

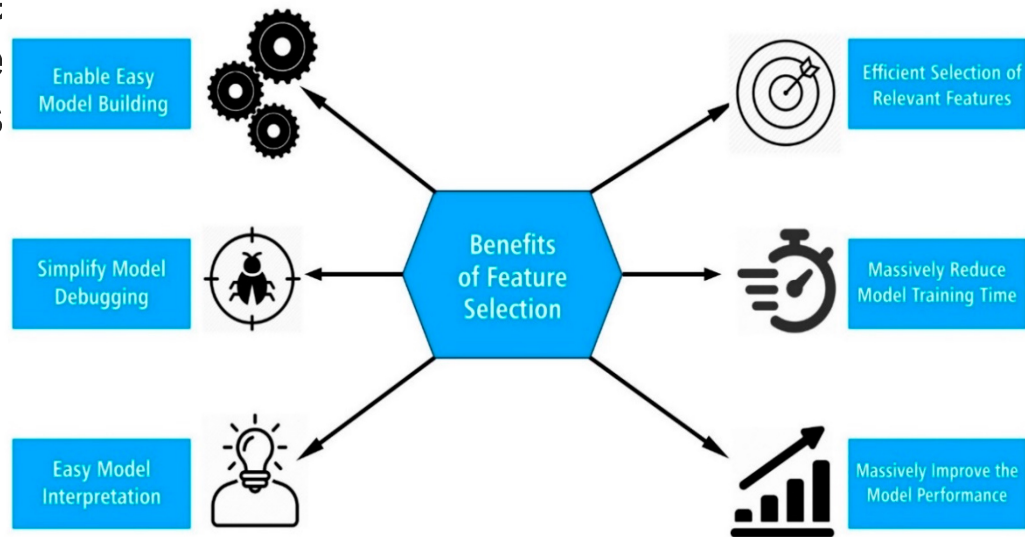


Today...



Why Feature Selection

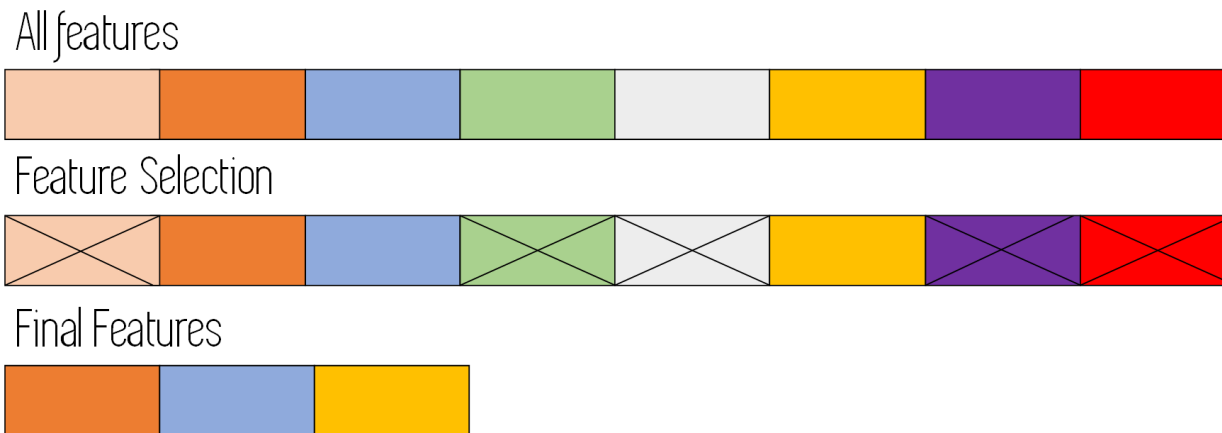
- In machine learning, one of the key challenges is to **select the right set of features** as inputs to a model.
- The features that are used to train a model will have a huge influence on the achieved performance.
- Irrelevant or partially relevant features can negatively impact the performance of a model.



heavy.ai

What is Feature Selection?

- Feature selection is the process of selecting a subset of relevant features used to train a machine learning model.



Why is feature selection important?

- Enhanced generalization by reducing overfitting
- Reduces training times
- Increase model interpretability
- Variable redundancy
- Reduces prediction time

Methods

- **Filter Methods** (fast, no/min feature interaction)
 - In this method, the selection of features is done independently of a machine learning algorithm. This method relies on the characteristics of the data to filter features based on a given metric.
 - Examples:
 - Chi-square test
 - Pearson Correlation
 - Mutual Information
 - **Minimum Redundancy-Maximum Relevance (mRMR)**



Minimum Redundancy-Maximum Relevance

Objective Function:

$$Rel = \sum_{x_i \in X} I(x_i; C)$$

$$Red = \sum_{\substack{x_i, x_j \in X, \\ \text{and } i \neq j}} I(x_i; x_j)$$

- X is the selected feature subset
- x_i, x_j : feature in X
- C is the class labels
- Rel : relevance between X and c
- Red : redundancy within X

$$\begin{aligned} I(X; Y) &= H(X) - H(X|Y) = H(Y) - H(Y|X) \\ &= \sum_{x \in X, y \in Y} p(x, y) \log_2 \frac{p(x, y)}{p(x)p(y)} \end{aligned}$$

Minimum Redundancy-Maximum Relevance

- S is the feature subset, Ω is the pool of all candidate features, the **minimum redundancy condition** is:

$$\min_{S \subset \Omega} \frac{1}{|S|^2} \sum_{i,j \in S} I(x_i, x_j)$$

where $|S|$ is the number of features in S .

- For classes $c=(c_i, \dots, c_k)$ the maximum relevance condition maximizes the total relevance of all features in S :

$$\max_{S \subset \Omega} \frac{1}{|S|} \sum_{i \in S} I(x_i, c)$$

H.C. Peng, F.H. Long, and C. Ding, Feature Selection Based on Mutual Information: Criteria of Max-Dependency, Max-Relevance, and Min-Redundancy, *IEEE Trans. Pattern Analysis and Machine Intelligence*, vol. 27, no. 8, 2005, pp. 1226–1238.

Minimum Redundancy-Maximum Relevance

- The mRMR feature set optimizes these two conditions simultaneously, either in quotient form:

$$\max_{s \subset \Omega} \left\{ \frac{\sum_i I(x_i, c)}{\frac{1}{|S|} \sum_{i,j \in S} I(x_i, x_j)} \right\}$$

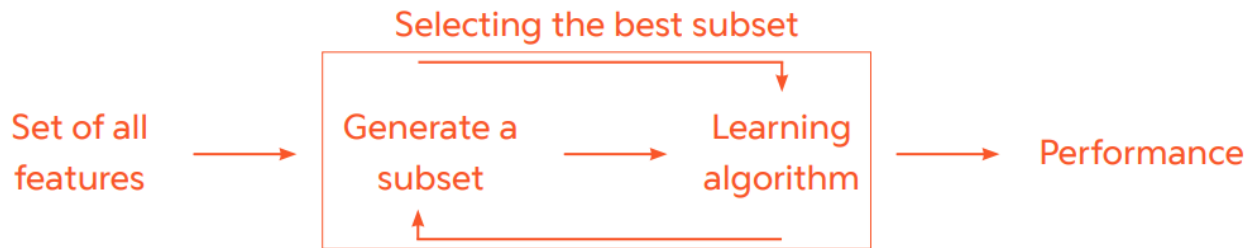
or in difference form:

$$\max_{s \subset \Omega} \left\{ \sum_i I(x_i, c) - \frac{1}{|S|} \sum_{i,j \in S} I(x_i, x_j) \right\}$$

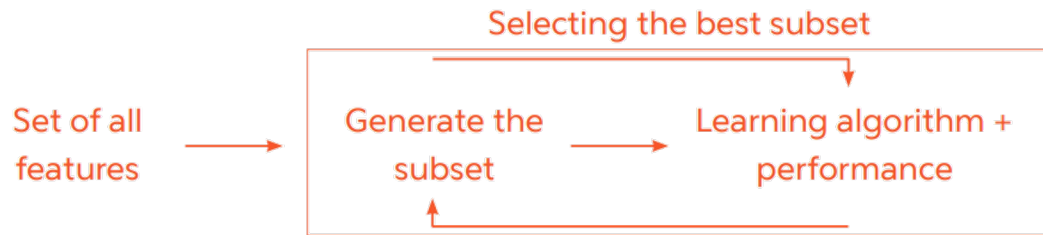
H.C. Peng, F.H. Long, and C. Ding, Feature Selection Based on Mutual Information: Criteria of Max-Dependency, Max-Relevance, and Min-Redundancy, *IEEE Trans. Pattern Analysis and Machine Intelligence*, vol. 27, no. 8, 2005, pp. 1226–1238.

Methods

- Filter Methods (fast, no/min feature interaction)
- **Wrapper Methods** (slow, more accurate, considers feature interaction)
 - In this method, feature selection is based on a search criteria where a model is initially trained on a subset of features. Based on the inferences drawn from the previous model, we decide to either add or remove features.
 - Examples:
 - Recursive Feature Elimination (greedy)



Methods



- Filter Methods (fast, no/min feature interaction)
- Wrapper Methods (slow, more accurate, considers feature interaction)
- **Embedded Methods** (more reliable feature estimates, reduce overfitting, robust to outliers, higher computational costs)
 - Embedded methods use the qualities of both the filter and wrapper methods. With this method, feature selection is embedded within the ML algorithm.
 - **LASSO regularization:** Lasso uses **L1 regularization/penalty**. It shrinks some parameters or feature coefficients to zero. It uses logistic regression to train a model with **L1 penalty** term to evaluate the coefficients of different variables and remove those variables with zero coefficients.

$$\sum_{i=1}^n (Y_i - \sum_{j=1}^p X_{ij} \beta_j)^2 + \lambda \sum_{j=1}^p |\beta_j|$$

Methods

Set of all
features



Generate the
subset



Learning algorithm +
performance

Selecting the best subset

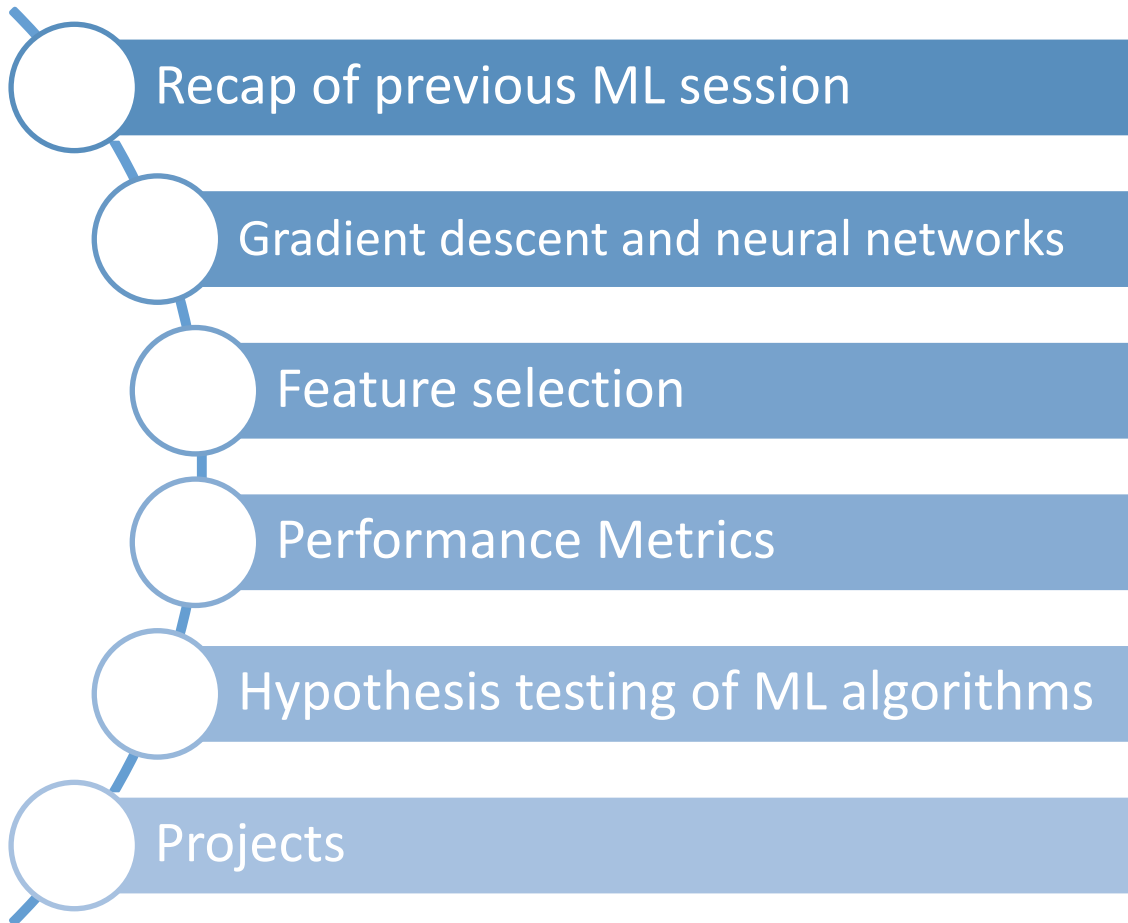


- Filter Methods (fast, no/min feature interaction)
- Wrapper Methods (slow, more accurate, considers feature interaction)
- **Embedded Methods** (more reliable feature estimates, reduce overfitting, robust to outliers, higher computational costs)
 - Lasso regularization.
 - Tree based random forest.
 - XGBoost
 - LightGBM
 - CatBoost...

Reading assignment 5

- An Introduction to Variable and Feature Selection, Journal of Machine Learning Research 2003
 - <https://www.jmlr.org/papers/volume3/guyon03a/guyon03a.pdf>
- Make sure you understand
 - mRMR
 - LASSO feature selection

Today...



Which Classifier is better?

Almost as many answers as there are performance measures!
(e.g., UCI Breast Cancer)

Algo	Acc	RMSE	TPR	FPR	Prec	Rec	F	AUC	Info S
NB	71.7	.4534	.44	.16	.53	.44	.48	.7	48.11
C4.5	75.5	.4324	.27	.04	.74	.27	.4	.59	34.28
3NN	72.4	.5101	.32	.1	.56	.32	.41	.63	43.37
Ripp	71	.4494	.37	.14	.52	.37	.43	.6	22.34
SVM	69.6	.5515	.33	.15	.48	.33	.39	.59	54.89
Bagg	67.8	.4518	.17	.1	.4	.17	.23	.63	11.30
Boost	70.3	.4329	.42	.18	.5	.42	.46	.7	34.48
RanF	69.23	.47	.33	.15	.48	.33	.39	.63	20.78

This and following slides courtesy of Nathalie Japkowicz, University of Ottawa

Which Classifier is better?

Ranking the results

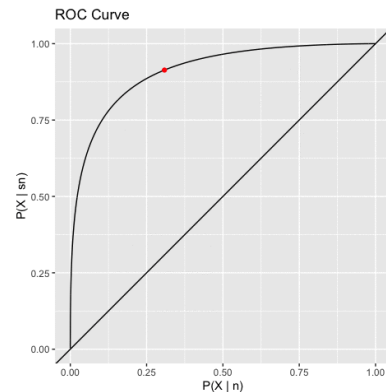
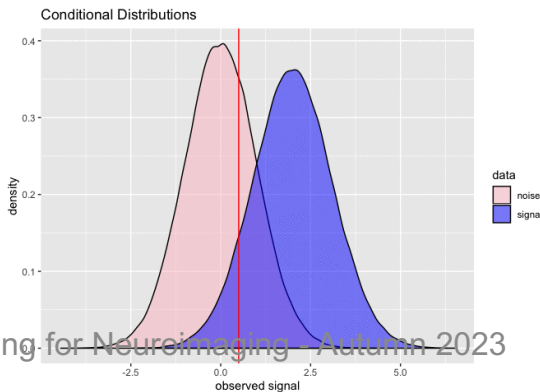
Algo	Acc	RMSE	TPR	FPR	Prec	Rec	F	AUC	Info S
NB	3	5	1	7	3	1	1	1	2
C4.5	1	1	7	1	1	7	5	7	5
3NN	2	7	6	2	2	6	4	3	3
Ripp	4	3	3	4	4	3	3	6	6
SVM	6	8	4	5	5	4	6	7	1
Bagg	8	4	8	2	8	8	8	3	8
Boost	5	2	2	8	7	2	2	1	4
RanF	7	6	4	5	5	4	7	3	7

A Few Confusion Matrix-Based Performance Measures

True class → Hypothesized class V	Pos	Neg
Yes	TP	FP
No	FN	TN
	P=TP+FN	N=FP+TN

A Confusion Matrix

- **Accuracy** = $(TP+TN)/(P+N)$
- **Precision** = $TP/(TP+FP)$
- **Recall/TP rate** = TP/P
- **FP Rate** = FP/N
- **ROC Analysis** moves the threshold between the positive and negative class from a small FP rate to a large one. It plots the value of the Recall against that of the FP Rate at each FP Rate considered.



Issues with Accuracy

True class →	Pos	Neg
Yes	200	100
No	300	400
	P=500	N=500

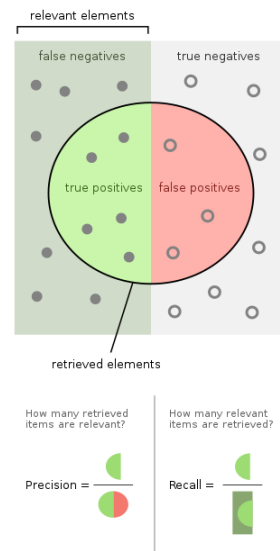
True class →	Pos	Neg
Yes	400	300
No	100	200
	P=500	N=500

- Both classifiers obtain 60% accuracy
- They exhibit very different behaviours:
 - On the left: weak positive recognition rate/strong negative recognition rate
 - On the right: strong positive recognition rate/weak negative recognition rate

Issues with Precision/Recall

True class →	Pos	Neg
Yes	200	100
No	300	400
	P=500	N=500

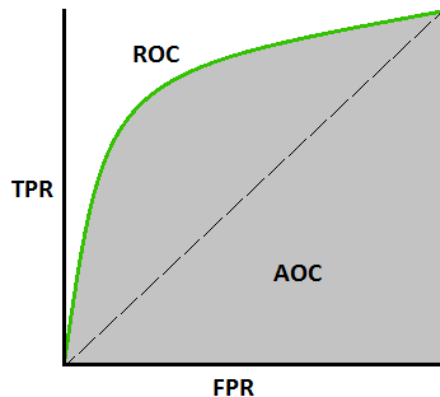
True class →	Pos	Neg
Yes	200	100
No	300	0
	P=500	N=100



- Both classifiers obtain the same precision and recall values of 66.7% and 40% (Note: the data sets are different)
- They exhibit very different behaviors:
 - Same positive recognition rate
 - Extremely different negative recognition rate: strong on the left / nil on the right
- Note: Accuracy has no problem catching this!

Is the AUC the answer?

- Many researchers have now adopted the AUC (the area under the ROC Curve).
- The principal advantage of the AUC is that it is **more robust** than Accuracy in class imbalanced situations.
- Indeed, given a 95% imbalance (in favour of the negative class, say), the accuracy of the default classifier that issues “negative” all the time will be 95%, whereas a more interesting classifier that actually deals with the issue, is likely to obtain a worse score.
- The AUC takes the **class distribution** into consideration.



RMSE

- The Root-Mean Squared Error (RMSE) is usually used for regression, but can also be used with probabilistic classifiers. The formula for the RMSE is:

$$\text{RMSE}(f) = \sqrt{\frac{1}{m} \sum_{i=1}^m (f(x_i) - y_i)^2}$$

where m is the number of test examples, $f(x_i)$, the classifier's probabilistic output on x_i and y_i the actual label.

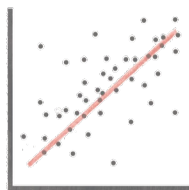
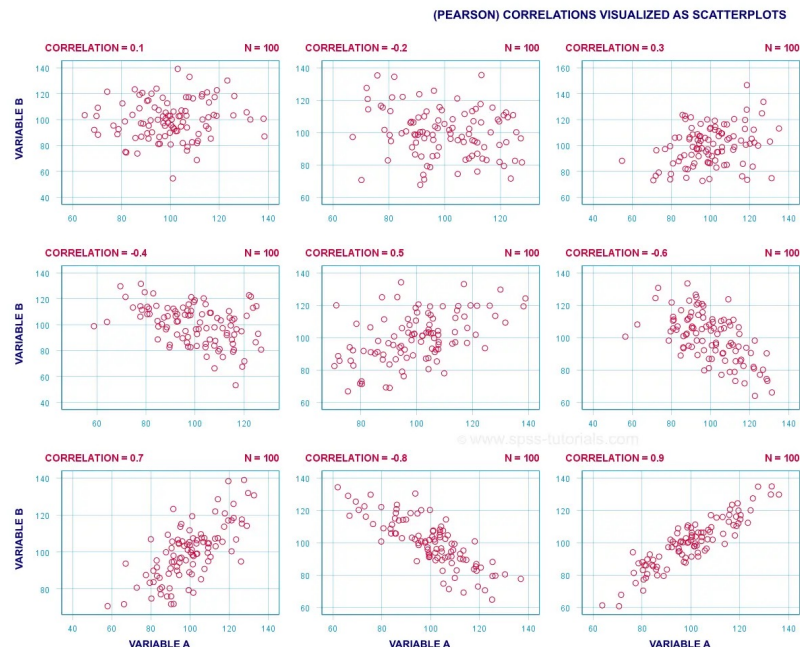
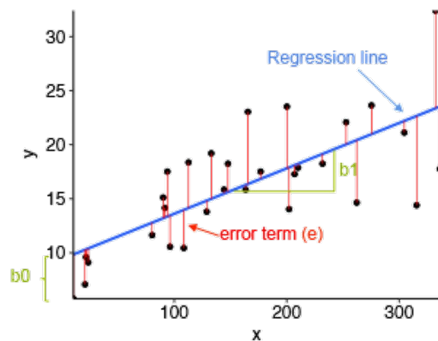
RMSE(f) RMSE(f)

ID	$f(x_i)$	y_i	$(f(x_i) - y_i)^2$
1	.95	1	.0025
2	.6	0	.36
3	.8	1	.04
4	.75	0	.5625
5	.9	1	.01

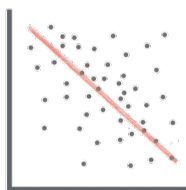
$$\begin{aligned}\text{RMSE}(f) &= \sqrt{\frac{1}{5} * (.0025 + .36 + .04 + .5625 + .01)} \\ &= \sqrt{0.975/5} = 0.4416\end{aligned}$$

Performance Metric for Regression Tasks

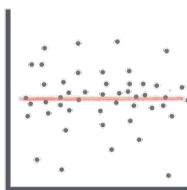
- Error
 - RMSE
 - MAE
- Correlation
 - Squared Correlation (R^2)



Positive Correlation




Negative Correlation

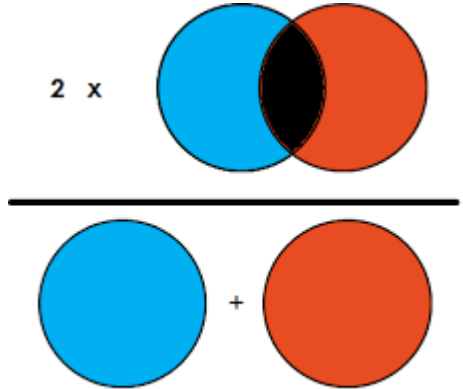


No Correlation

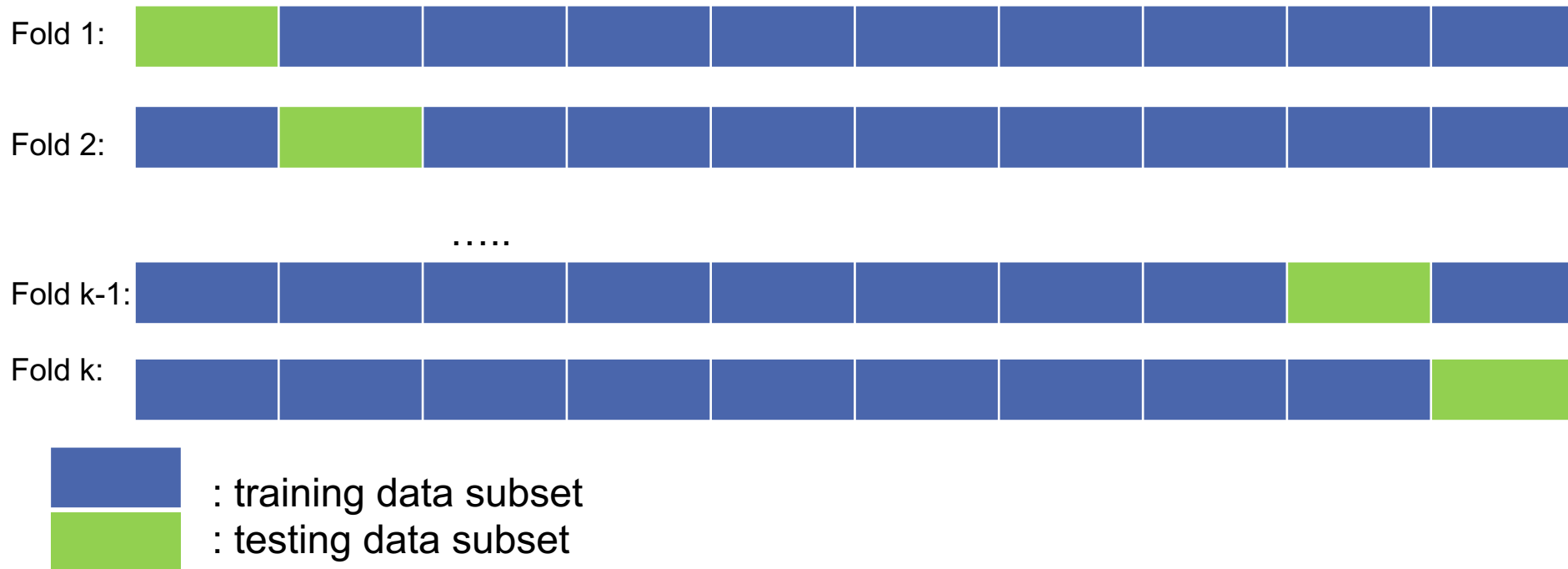
Question

In a segmentation task, what are the most popular performance measures?

$$\text{IoU} = \frac{\text{Area of Overlap}}{\text{Area of Union}}$$


$$\text{Dice Coefficient} = \frac{2 \times \text{Intersection}}{\text{Union} + \text{Intersection}} = \frac{2\text{TP}}{2\text{TP} + \text{FN} + \text{FP}}$$


k-fold Cross-Validation



In Cross-Validation, the data set is divided into k folds and at each iteration, a different fold is reserved for testing and all the others, used for training the classifiers.

Projects

- Start discussing your project ideas with us
 - Email us
 - Office hours and meetings?
- [Human Connectome Project \(HCP\)](#)
 - [Alzheimer's Disease Neuroimaging Initiative: ADNI](#)
 - [Parkinson's Progression Markers Initiative \(PPMI\)](#)
 - [OASIS Brains - Open Access Series of Imaging Studies](#)
 - [ABIDE - Autism Brain Imaging Data Exchange International Neuroimaging Data-sharing Initiative](#)
 - [Multimodal Brain Tumor Segmentation Challenge 2020 \(BraTS\)](#)
 - Dataset of your interest

Project Proposal

A **2-page document (including references)** with the following sections

1. Problem statement (clear input/output)
2. Motivation (why the problem is important)
3. Prior work
 - a) Key challenges of the problem
 - b) Challenges taken care of by the prior work, what is remained
4. Contribution (if any)
5. Technical Details
6. Datasets and Performance Metrics
7. Experiments Plan
8. Team Members