

DLM12022

Introduction to machine learning

By

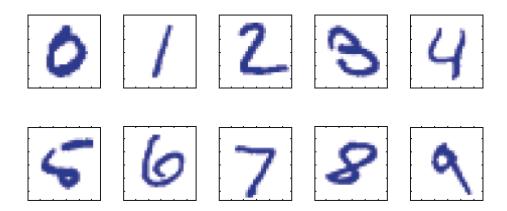
Pierre-Marc Jodoin (Hugo Larochelle)



Killer question...

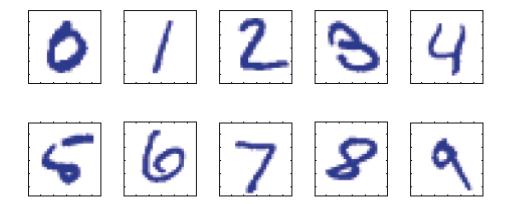
What is machine learning?





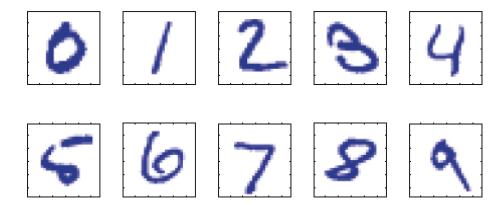
Answer: Design your own rules?

- > A series of aligned pixels => '1'
- > A circle of pixels => '0'
- > Etc.



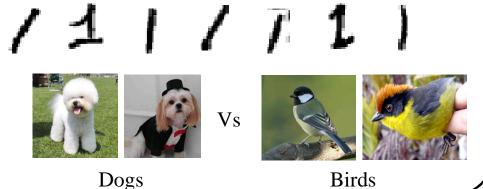
Answer: Design your own rules? Wrong

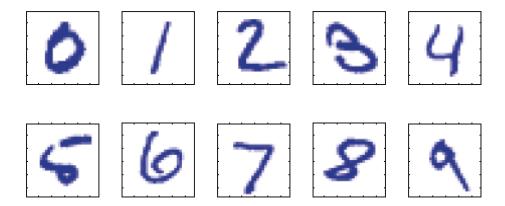
> Bad generalization / 1 / 7 1



Answer: Design your own rules? Wrong

- Bad generalization
- > Often difficult





Answer: Let the computer « **learn** » the rules

➤ Main goal of machine learning

Two large families

Supervised learning

Unsupervised learning

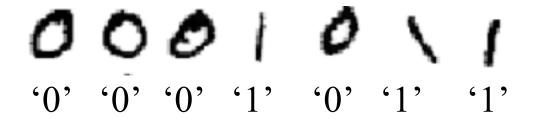
Two large families

Supervised learning

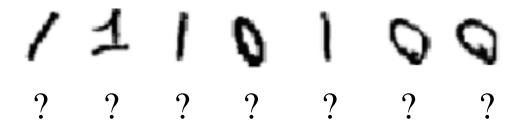
Unsupervised learning

Supervised learning

Provide the algorithm with annotated training data

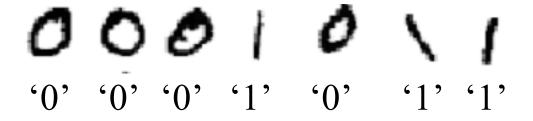


...and the algorithm returns a function capable of **generalizing** on new data



Supervised learning

Provide the algorithm with annotated training data



The training dataset

$$D = \{ (\vec{x}_1, t_1), (\vec{x}_2, t_2), \dots, (\vec{x}_N, t_N) \}$$

where $\vec{x}_i \in \mathbb{R}^d$ is an **input** and t_i is a **target**

Goal of a supervised machine learning method

From a **training dataset:**
$$D = \{(\vec{x}_1, t_1), (\vec{x}_2, t_2), ..., (\vec{x}_N, t_N)\}$$

$$\vec{x}_i \in \Re^d$$
input data t_i target associated to \vec{x}_i

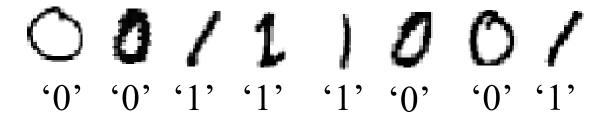
the goal is to learn a function that may predict t_i given \vec{x}_i

$$y_{\vec{w}}(\vec{x}_i) \rightarrow t_i$$

where \vec{w} are the **parameters** of the model.

Supervised learning

Once the model $y_{\vec{w}}(\vec{x})$ is trained, we use a **test set** D_{test} to gauge the **generalization** capabilities of the model.

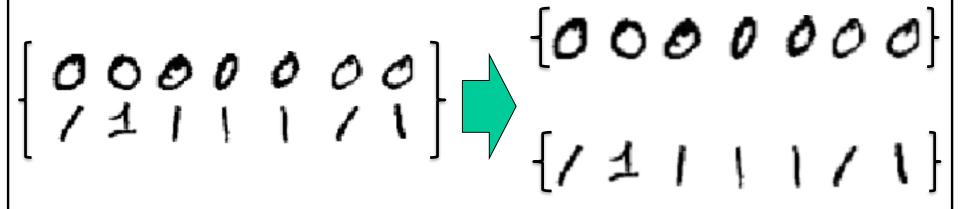


Two large families

Supervised learning

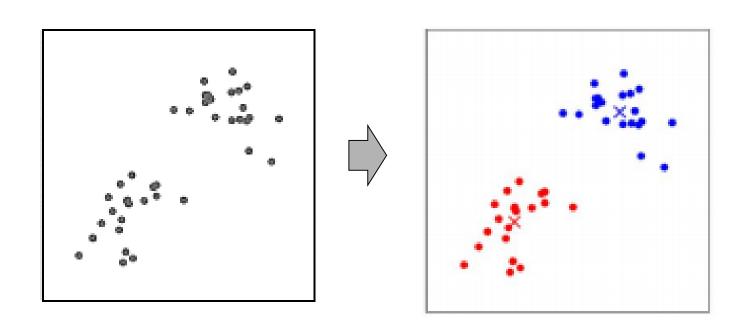
Unsupervised learning

When no target is explicitly provided > E.g. data *clustering*



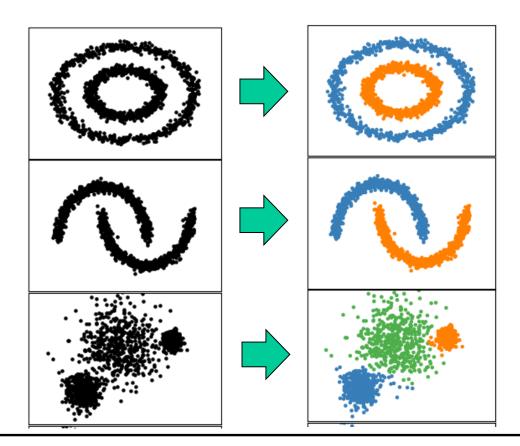
When no target is explicitly provided

E.g. data *clustering*

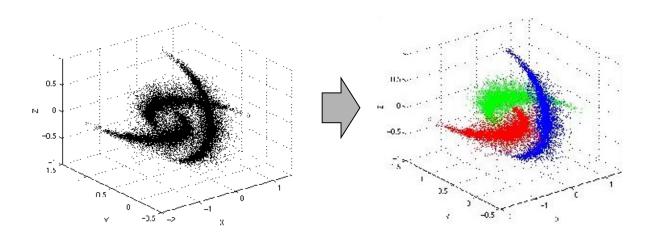


When no target is explicitly provided

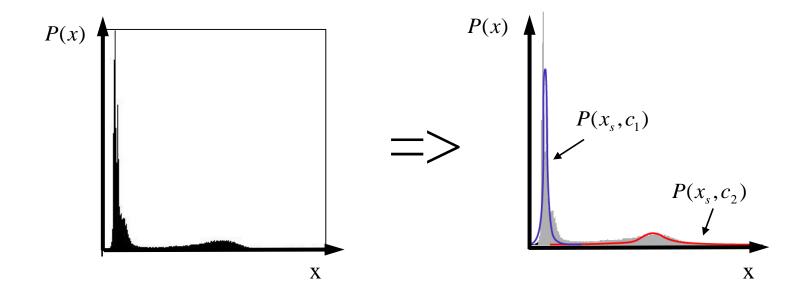
E.g. data *clustering*



No limit to dimensionality. Could be 3D, 4D,...100kD

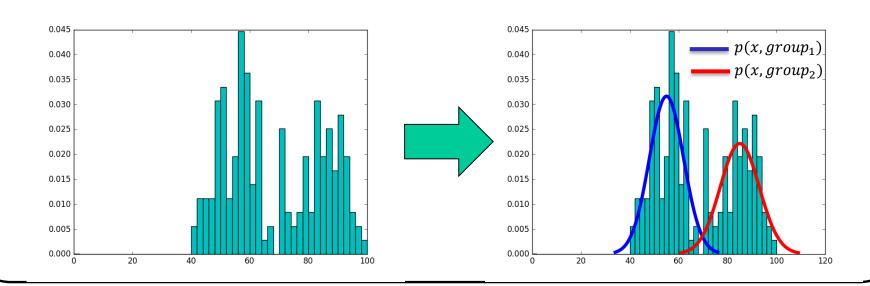


The goal is often to learn the distribution p(x) from which the data has been generated from



The goal is often to learn the distribution p(x) from which the data has been generated from

Example: find two groupes of patients following a memory test



The goal is often to learn the distribution p(x) from which the data has been generated from

Other applications

- > Data compression
- > Data visualisation
- > Population analysis
- > Image segmentation
- > etc.

Supervised vs non-supervised

Supervised learning: there is a target

$$D = \{ (\vec{x}_1, t_1), (\vec{x}_2, t_2), \dots, (\vec{x}_N, t_N) \}$$

Unsupervised learning: unknown target

$$D = \left\{ \vec{x}_1, \vec{x}_2, \dots, \vec{x}_N \right\}$$

Supervised vs non-supervised

Supervised learning: there is a tar

$$D = \{ (\vec{x}_1, t_1), (\vec{x}_2, t_2) \}$$

Logistic regression
Perceptron
Multilayer perceptron
Convolutional neural networks
Recurrent neural networks
Graph Neural Nets
Etc.

Unsupervised learning: unknown target

$$D = \left\{ \vec{x}_1, \vec{x}_2, \dots, \vec{x}_N \right\}$$

Supervised vs non-supervised

Supervised learning: there is a target

$$D = \{ (\vec{x}_1, t_1), (\vec{x}_2, t_2), \dots, (\vec{x}_N, t_N) \}$$

Unsupervised learning: unknown

Autoencoders Variational autoencoders GANs

$$D = \left\{ \vec{x}_1, \vec{x}_2, \dots, \vec{x}_N \right\}$$

Back to supervised learning

Supervised learning

Two main applications

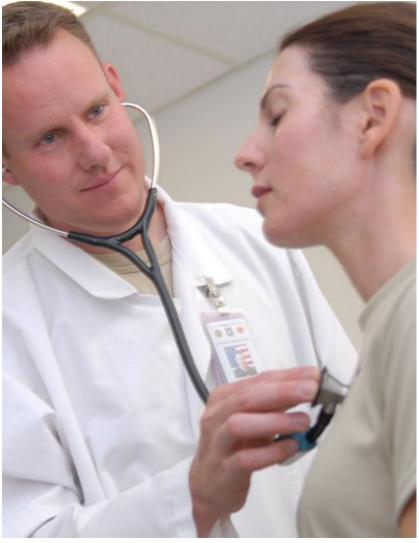
- \triangleright Classification: the target is a class label $t \in \{1, ..., K\}$
 - Exemple : disease recognition
 - \checkmark \vec{x} : vector of medical measures, age, sex, etc.
 - ✓ *t* : myocardial infarction, dilated cardiomyopathy, hypertrophic cardiomyopathy, normal
- **Régression :** the target is a real number $t \in \mathbb{R}$
 - Exemple : prediction of life expectancy
 - \checkmark \vec{x} : vector of medical measures, age, sex, etc.
 - \checkmark t: number of months before death.

Supervised learning

Two main applications

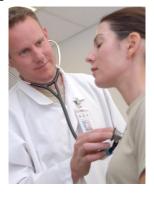
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Simple example of binary classification



From Wikimedia Commons the free media repository

Simple example of binary classification



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Patient 1	
Patient 2	

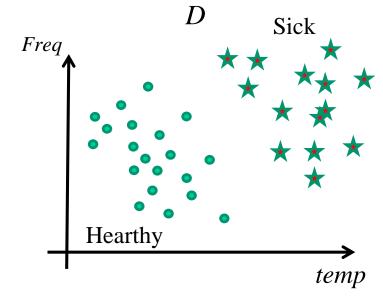
1 4110111 2

Patient 3

Patient N

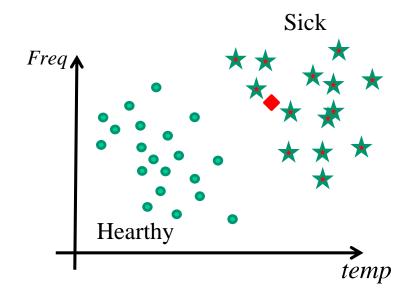
(temp, freq)	Diagnostic
(37.5, 72)	hearthy
(39.1, 103)	sick
(38.3, 100)	sick
()	•••
(36.7, 88)	hearthy
·	·





Simple example of binary classification

A new patient shows up at the hospital **How can we predict its state?**





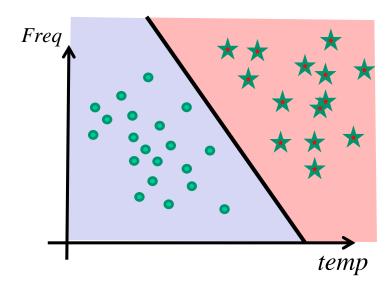
From Wikimedia Commons the free media repository

Solution



From Wikimedia Commons the free media repository

Divide the feature space in two regions : healthy and sick

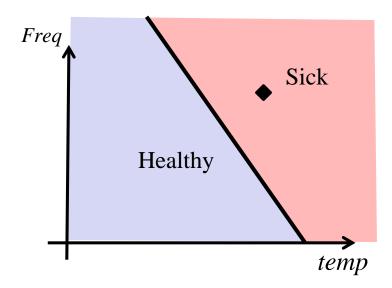


Solution



From Wikimedia Commons the free media repository

Divide the feature space in two regions : healthy and sick

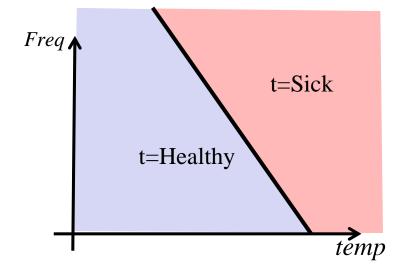


More formally

$$y_{\vec{w}}(\vec{x}) = \begin{cases} H \text{ ealthy if } \vec{x} \text{ is in the blue region} \\ Sick \text{ otherwise} \end{cases}$$

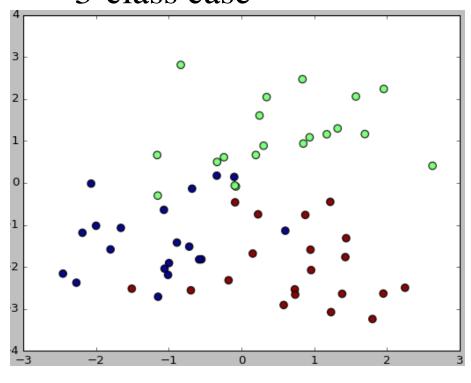


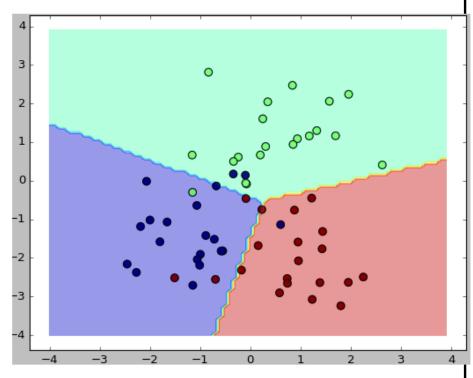
From Wikimedia Commons the free media repository



Classification

3-class case





3 classes •, •, • in a 2D feature space

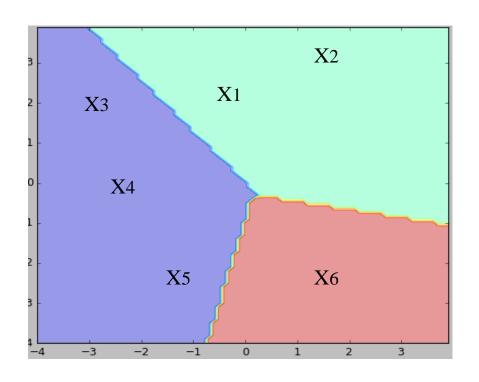
Once training is over

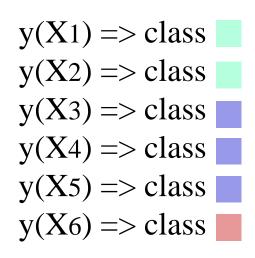
$$y(\bullet) = class 1$$

 $y(\bullet) = class 2$
 $y(\bullet) = class 3$

Classification

Once training is over, we have a function y(x) that convert a point x into a class label





Example of a classification dataset

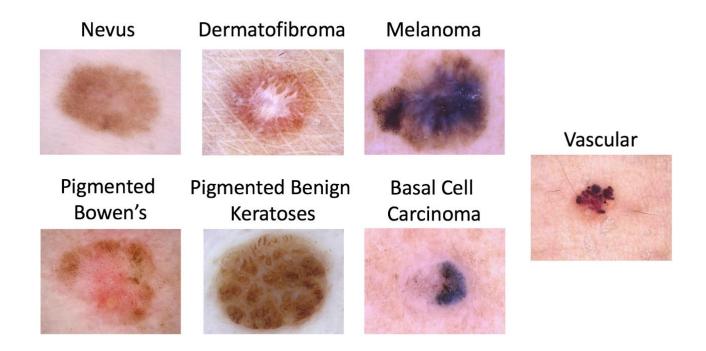
/ 1 | 1 / 1 / / / / 1 / / / / / ファチ17ァファファファファ

Example of a classification dataset

- 10 classes
- 70,000 images => 60,000 training
 - => 10,000 test
- Images are in grayscale => 28x28

We can **vectorize these images** and represent it by a vector of size 28x28 = 784 dimensions.

ISIC melanoma classification challenge.



https://challenge.isic-archive.com/landing/2018/

ISIC melanoma classification challenge dataset

- 7 classes
- 11,527 images,
 - => 10,015 training
 - => 1,512 test
- Each image is in RGB
 - => 628x417x3

We can **vectorize these images** and represent it by a vector of size 628x417x3 = 785,628 dimensions.

Chess X-Ray Pneumonia

Healthy





Pneumonia





https://www.kaggle.com/datasets/paultimothymooney/chest-xray-pneumonia

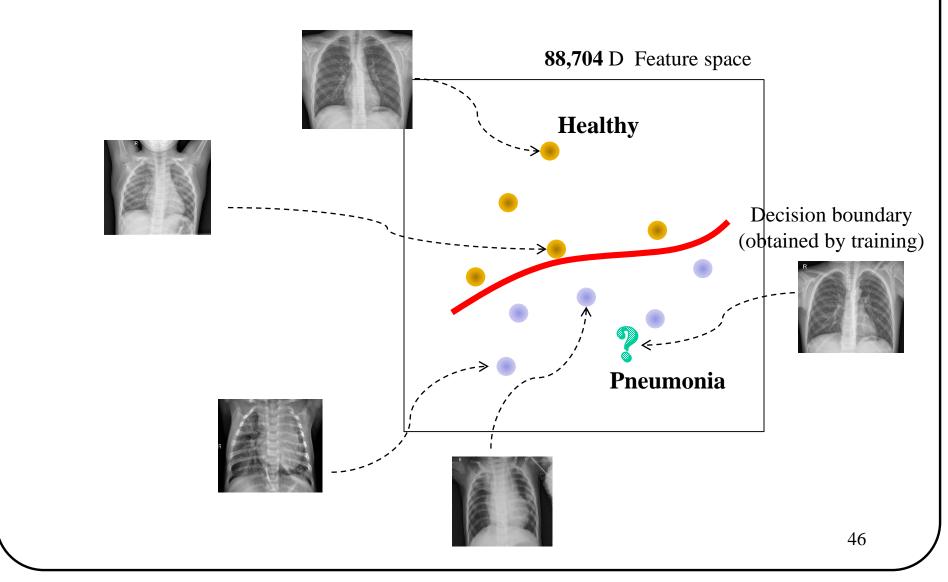
Chess X-Ray Pneumonia

- 2 classes
- 5,840 images,
 - => 5,216 training
 - => 624 test
- Each image is in grayscale

We can **vectorize these images** and represent it by a vector of size 336x264 = 88,704 dimensions.

Supervised learning

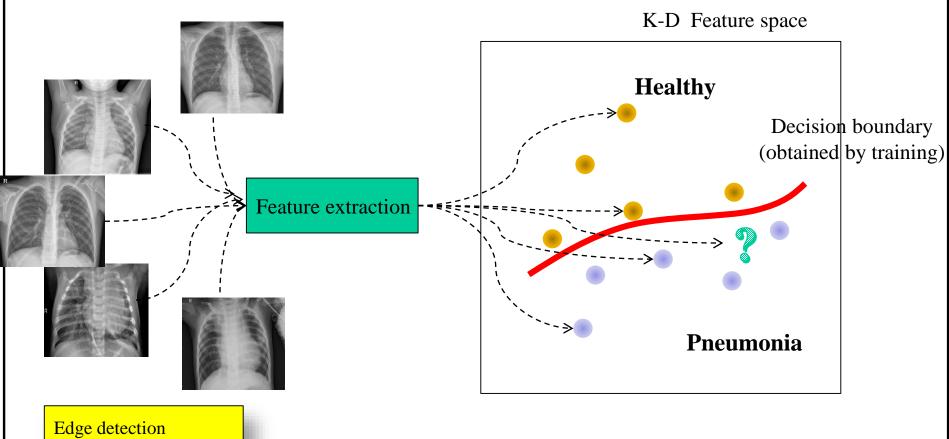
Chess X-Ray Pneumonia





Very large feature spaces (like 88,704 dim) are problematic.

Supervised learning

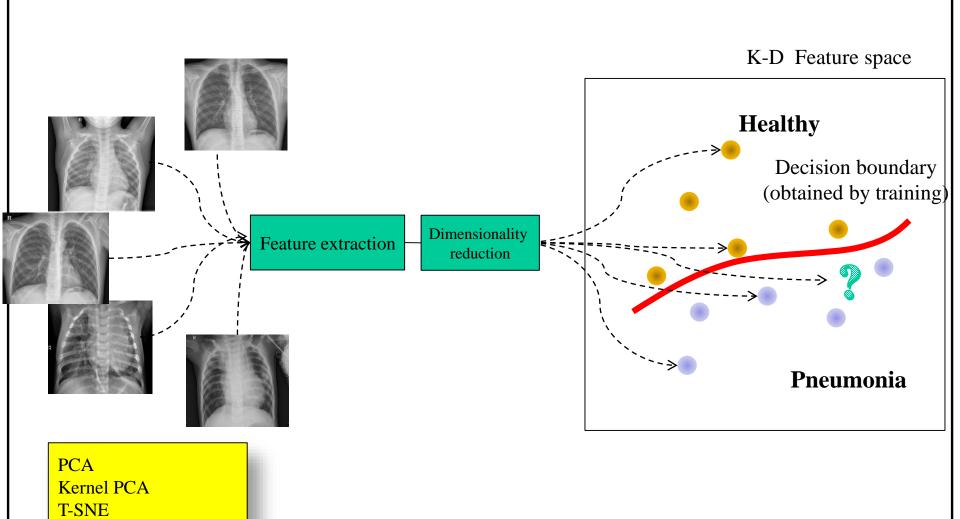


Edge detection
Histograms
Filters of all kinds
Moments
Etc.

Supervised learning

ISOMAPS

Etc.

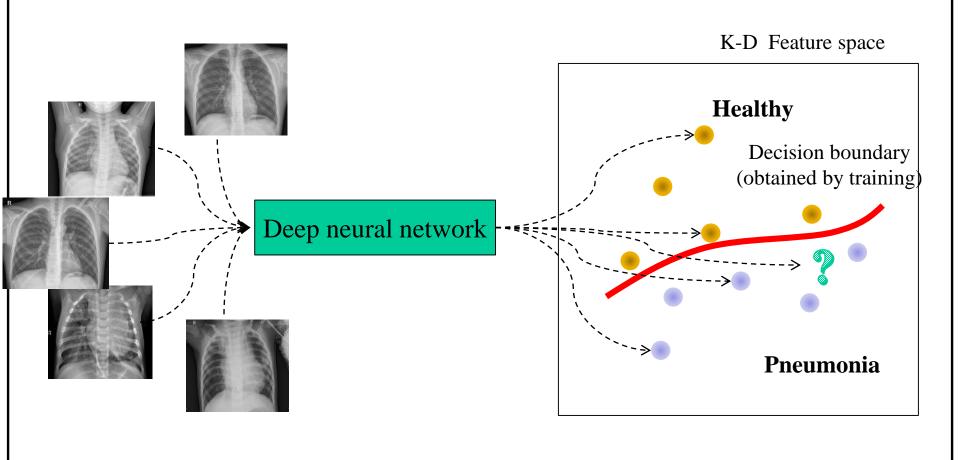


49



Spoiler alert

In 2022...



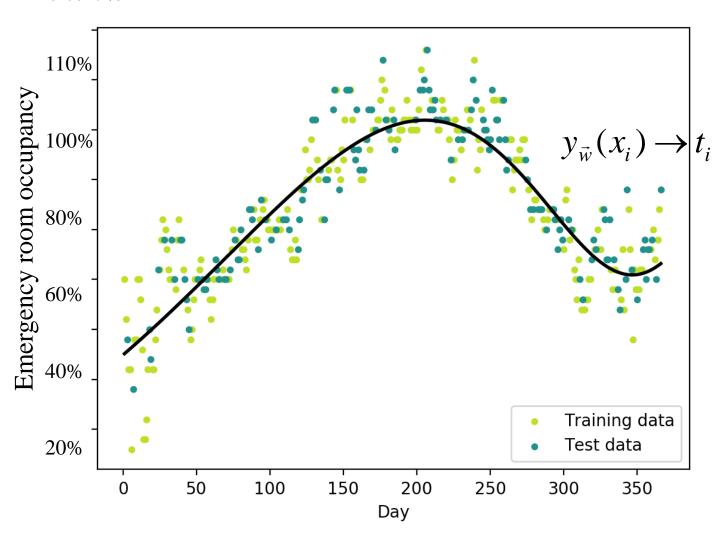
Supervised learning

Two main applications

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 - Exemple : disease recognition
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Example

Percent %



Formal example: **1D regression**

1D Regression

Simple example

> Data

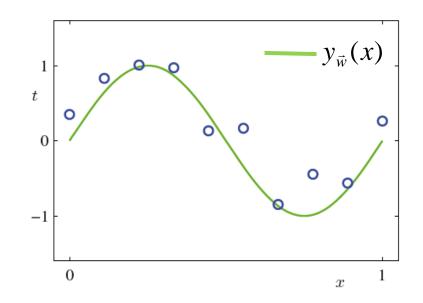
- ✓ input : scalar $x \in [0,1]$
- ✓ target : scalar $t \in [-1,1]$
- ➤ Training dataset *D* contains:

$$\checkmark X = (x_1, ..., x_N)^T$$

 $\checkmark T = (t_1, ..., t_N)^T$

≻ Goal:

 \checkmark Get a prediction \hat{t} for each new data \hat{x}



1D Regression

Simple example

> Data

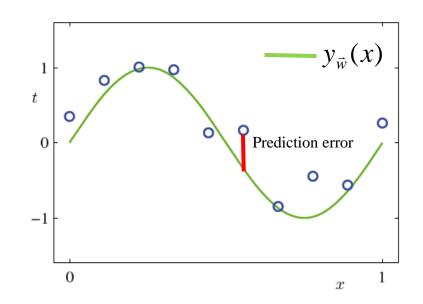
- ✓ input : scalar $x \in [0,1]$
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➤ Goal:

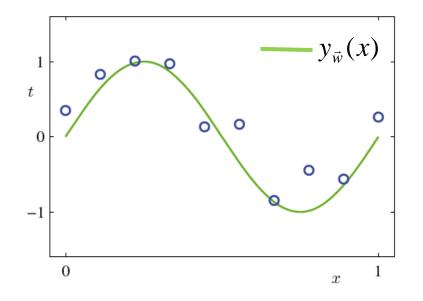
 \checkmark Get a prediction \hat{t} for each new data \hat{x}



Polynomial shape

Let's assume that our data has a polynomial shape

$$y_{\vec{w}}(x) = w_0 + w_1 x + w_2 x^2 + \dots + w_M x^M$$
$$= \sum_{i=0}^{M} w_i x^i$$



- $\triangleright y_{\vec{w}}(x)$ is our **model**
 - ✓ It implicitly includes our hypothesies on the problem at hand
 - \checkmark A model always has parameters that one needs to find (here \vec{w})

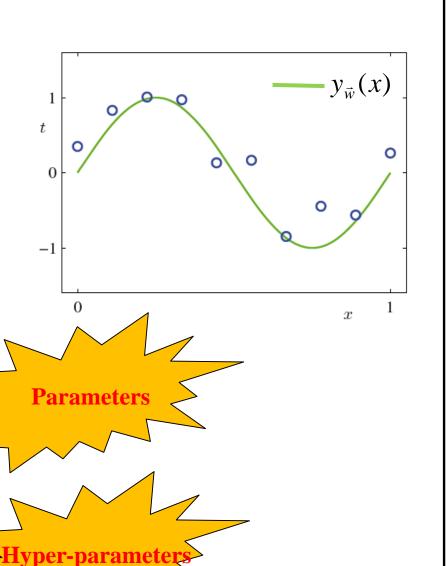
Unknowns

$$y_{\vec{w}}(x) = w_0 + w_1 x + w_2 x^2 + \dots + w_M x^M$$
$$= \sum_{i=0}^{M} w_i x^i$$

Two unknowns

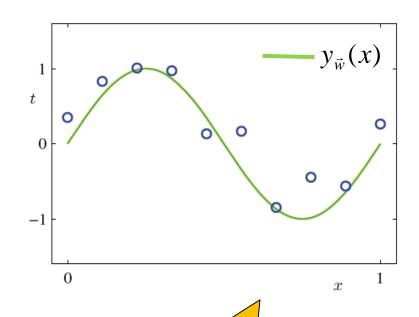
$$\vec{w} \in R^M$$

$$M \in \mathbb{N}^{\geq 0}$$



Training

$$y_{\vec{w}}(x) = w_0 + w_1 x + w_2 x^2 + \dots + w_M x^M$$
$$= \sum_{i=0}^{M} w_i x^i$$



Two unknowns

$$\vec{w} \in R^M$$

 $M \in \mathbb{N}^{\geq 0}$

Training = find the « Best » w (and somtimes M) from X and T

Polynomial regression

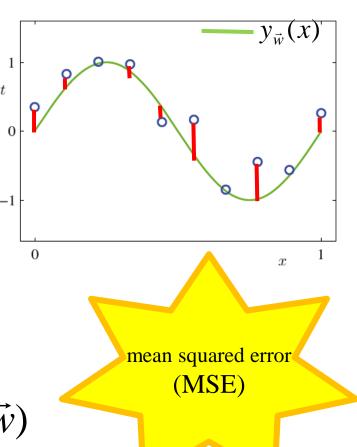
ightharpoonup If *M* is given, how can we find the best \vec{w} ?

The « best » \vec{w} is the one that minimizes the error (or the **loss**) on the training data

$$E_D(\vec{w}) = \frac{1}{N} \sum_{n=1}^{N} (y_{\vec{w}}(x_n) - t_n)^2$$

➤ More formally

$$\vec{w} = \arg\min_{\vec{w}} E_D(\vec{w})$$



Hyperparameters

 \triangleright How to find the best M?

Hyperparameters cannot be **estimated** with classical **optimization algorithms** (gradient descent, Newton method, Simplex method, etc.) as for \vec{w}

hyperparameters are often fixed manually.

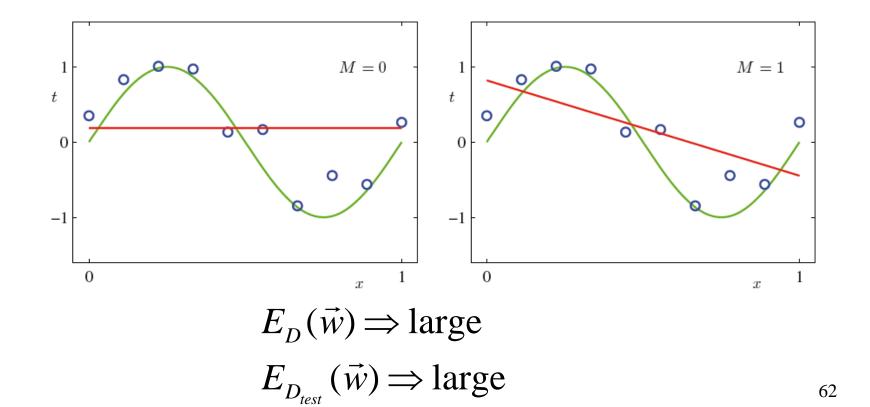
BUT BEWARE!!! Their value greatly influence the final results.

Underfitting

$$M = 0 \implies y_{\vec{w}}(x) = w_0$$

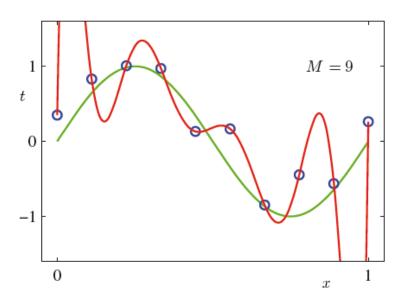
 $M = 1 \implies y_{\vec{w}}(x) = w_0 + w_1 x$

A small M gives a simplistic model that will **underfit** the data.



Overfitting

A large M gives a model that « learn by heart » and thus overfit training data



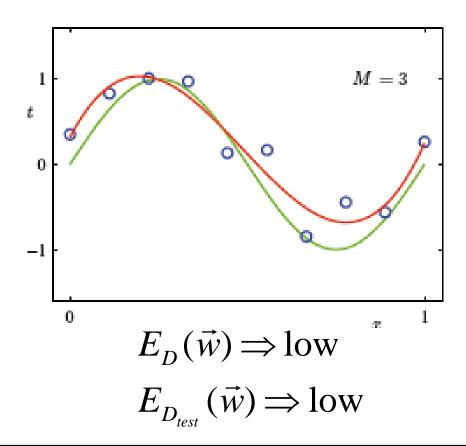
$$E_D(\vec{w}) \Rightarrow \text{VERY low}$$

 $E_{D_{test}}(\vec{w}) \Rightarrow \text{Large}$

$$E_{D_{tart}}(\vec{w}) \Longrightarrow \text{Large}$$

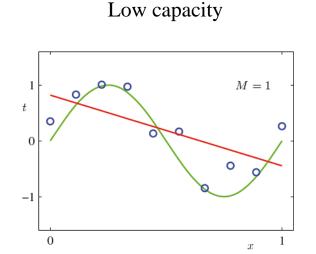
Over- and underfitting

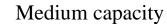
Need for an intermediate value for which the training and the testing errors are low

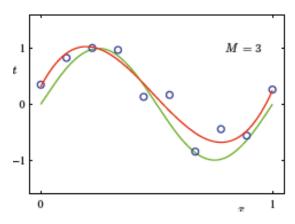


Hyperparameters often control the capacity of a model

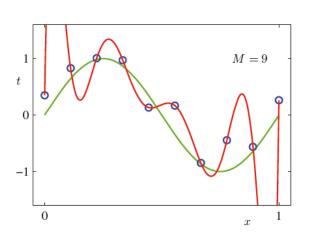
Capacity: ability of a model to fit the training data



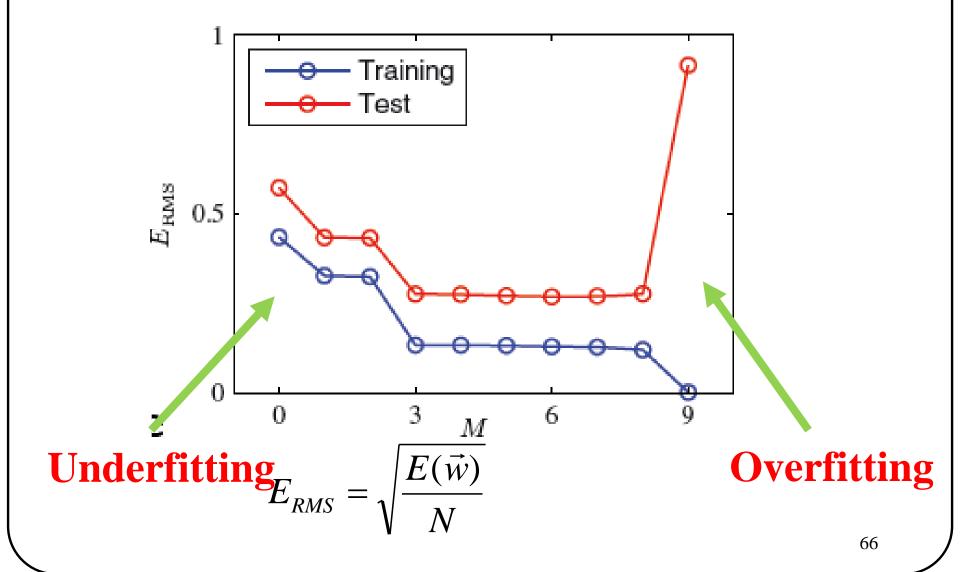




Large capacity

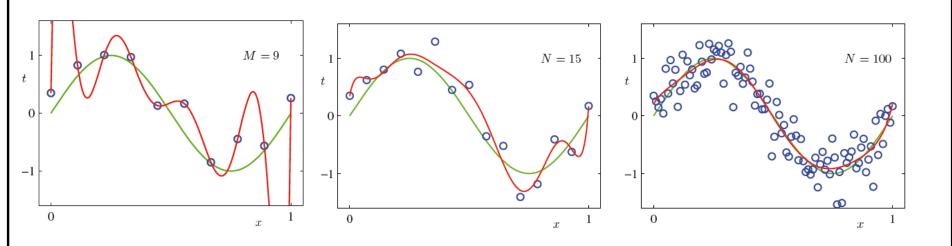


Over- and underfitting



Generalization

The more data you have, the better a high capacity model will generalize.



If we have a **large M** and **few training data**, how do we prevent our model from overfitting?



Parameter values \vec{w} for different M without regularization

	M=0	M = 1	M = 3	M = 9
w_0	0.19	0.82	0.31	0.35
w_1		-1.27	7.99	232.37
w_2			-25.43	-5321.83
w_3			17.37	48568.31
w_4				-231639.30
w_5				640042.26
w_6				-1061800.52
w_7				1042400.18
w_8				-557682.99
w_9				125201.43

To prevent over-fitting

- 1. Choose a small « M »
- 2. Reduce capacity by regularization

Exemple : penalyse the
$${\bf L2\ norm}$$

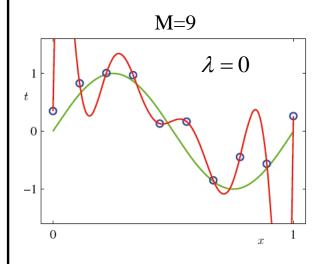
Constante that controls regularization

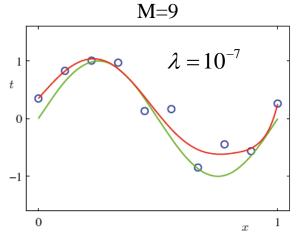
$$E_{D}(\vec{w}) = \frac{1}{N} \sum_{n=1}^{N} (t_{n} - y_{\vec{w}}(\vec{x}))^{2} + \lambda \|\vec{w}\|^{2}$$

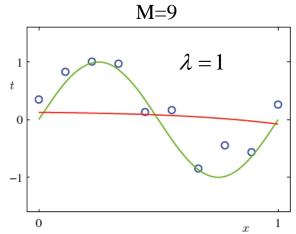
$$\|\vec{w}\|^2 = \vec{w}^T \vec{w} = w_0^2 + w_1^2 + \dots + w_M^2$$

Ridge model

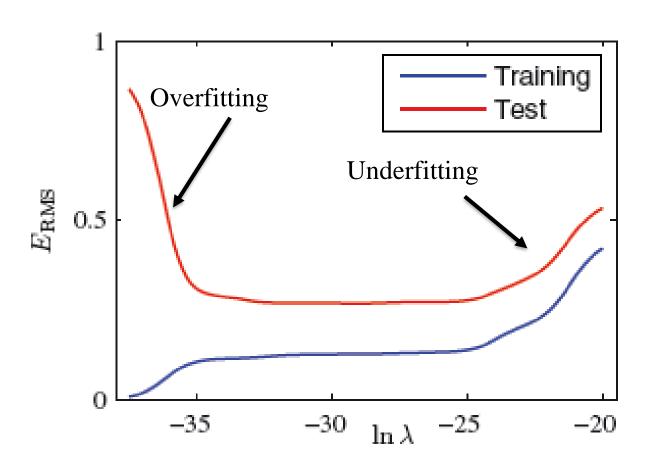
Strong regularization = less capacity





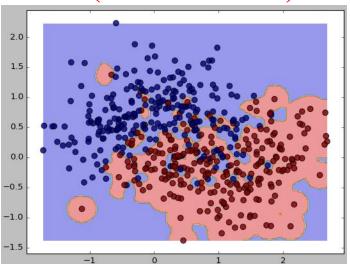


Influence on the training and testing error

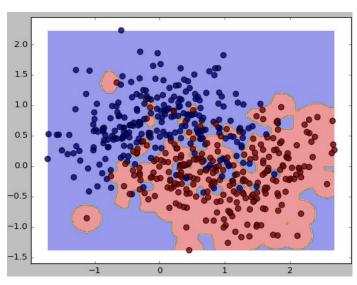


Over- and under-fitting also influence classification

Overfitting (Classification)

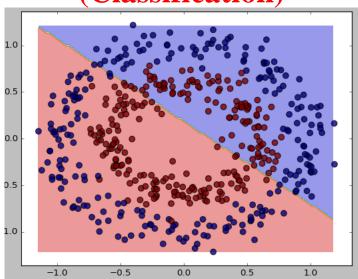


Training accuracy = 99.6%

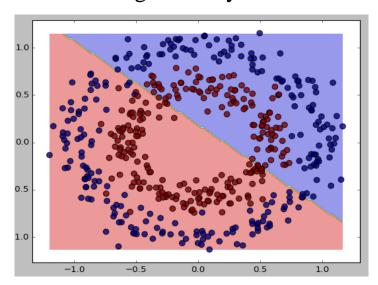


Testing accuracy = 78%

Underfitting (Classification)

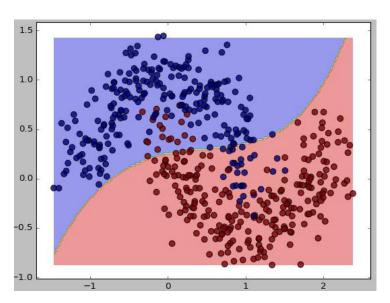


Training accuracy =52.2%

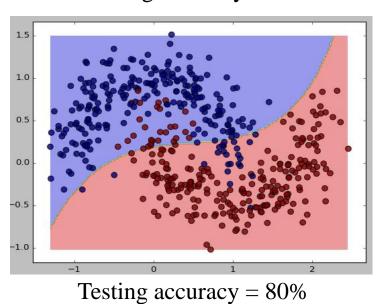


Testing accuracy = 51.2%

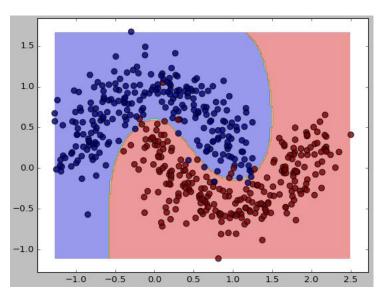
Could be better...



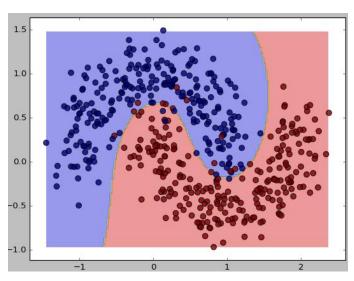
Training accuracy =82%



Wonderful !!!



Training accuracy =97.8%



Testing accuracy = 96.2%

$$E_D(\vec{w}) = \sum_{n=1}^{N} (y_{\vec{w}}(x_n) - t_n)^2 + \lambda ||\vec{w}||^2$$
$$||\vec{w}||^2 = \vec{w}^T \vec{w} = w_0^2 + w_1^2 + \dots + w_M^2$$

Model selection

How to find the right hyper-parameters?

M and λ

How to find the right M and the right λ ?

- Very bad idea : choose randomly
- Bad idea: take many (M, λ) and keep the one with the lowest training error
 - > overfitting
- Bad idea: take many (M, λ) and keep the one with the lowest tesintg error
 - $\triangleright_{D_{test}}$ should NEVER be used to train a model
- Good solution: take many (M, λ) and keep the one with the lower validation error

Cross-validation

1- Randomly devide data in 2 groups



2- FOR M from M_{\min} to M_{\max} FPR λ from λ_{\min} to λ_{\max}

Train the model on $D_{\it train}$ Compute error on $D_{\it valid}$

3- Keep (M, λ) with the lowest **validation error**

k-fold cross-validation

FOR M from M_{\min} to M_{\max} FOR λ from λ_{\min} to λ_{\max} FOR j from 0 à K

Divide the labeled data in 2 groups => $D_{train}D_{valid}$

Train the model on $D_{\it train}$ Compute error on $D_{\it valid}$

Keep (M, λ) with the lowest **mean validation error**

K-fold cross-validation with K = 10

Mean validation error

STD

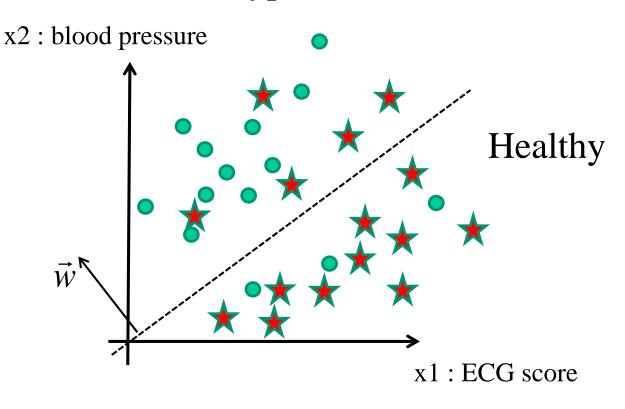
```
2.832 (+/-0.116) for {'regression': 'poly', 'M': 3, 'lambda': 0.01}
1.854 (+/-0.072) for {'regression': 'poly', 'M': 3, 'lambda': 0.1}
1.910 (+/-0.065) for {'regression': 'poly', 'M': 3, 'lambda': 1}
1.902 (+/-0.077) for { 'regression': 'poly', 'M': 3, 'lambda': 10}
2.844 \ (+/-0.101)  for {'regression': 'poly', 'M': 4, 'lambda': 0.01}
2.864 \ (+/-0.089)  for {'regression': 'poly', 'M': 4, 'lambda': 0.1}
1.910 (+/-0.065) for {'regression': 'poly', 'M': 4, 'lambda': 1}
1.894 (+/-0.086) for {'regression': 'poly', 'M': 4, 'lambda': 10}
2.848 \ (+/-0.080)  for {'regression': 'poly', 'M': 5, 'lambda': 0.01}
                                                                      BEST!
1.904 (+/-0.064) for { 'regression': 'poly', 'M': 5, 'lambda': 0.1}
0.916 \ (+/-0.069)  for {'regression': 'poly', 'M': 5, 'lambda': 1}
                                                                      M=5,
1.870 (+/-0.072) for { 'regression': 'poly', 'M': 5, 'lambda': 10}
                                                                       \lambda = 1
2.846 \ (+/-0.090)  for {'regression': 'poly', 'M': 6, 'lambda': 0.01}
2.906 (+/-0.062) for {'regression': 'poly', 'M': 6, 'lambda': 0.1}
1.904 (+/-0.075) for {'regression': 'poly', 'M': 6, 'lambda': 1}
2.858 (+/-0.112) for {'regression': 'poly', 'M': 6, 'lambda': 10}
```

In short

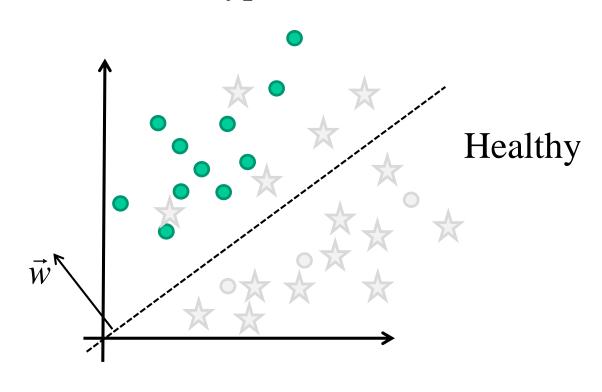
- ✓ The goal is to train a model on a training dataset with good generalization capabilites
- ✓ Has hyper-parameters that control the capacity of the model, choisis à l'aide d'une procédure de sélection de modèle
- ✓ mesure sa performance de **généralisation** sur un **ensemble de test**
- ✓ Aura une meilleure performance de généralisation si la quantité de données d'entraînement augmente
- ✓ Peut souffrir de sous-apprentissage (pas assez de capacité) ou de sur-apprentissage (trop de capacité)

Evaluation metrics

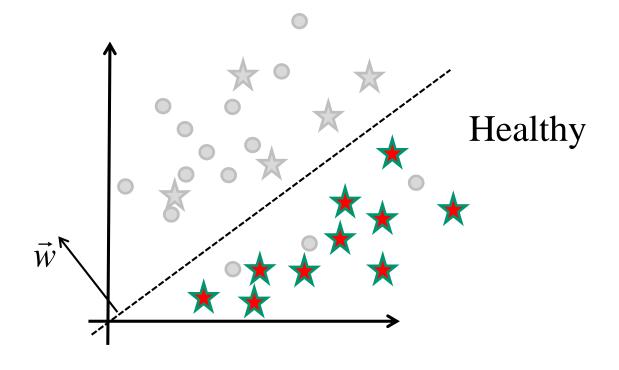
How to evaluate a model?



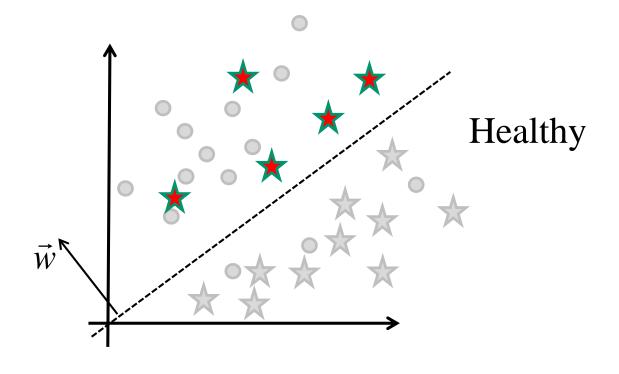
True positive (11)



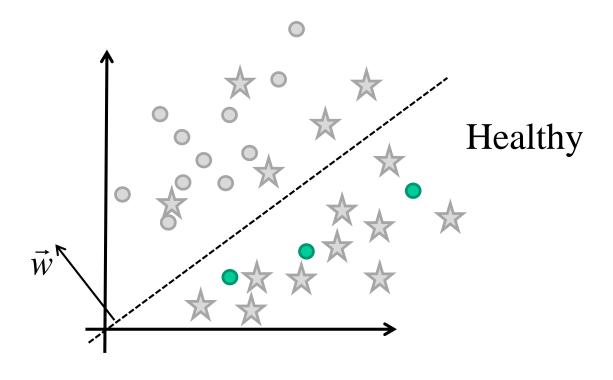
True negative (10)



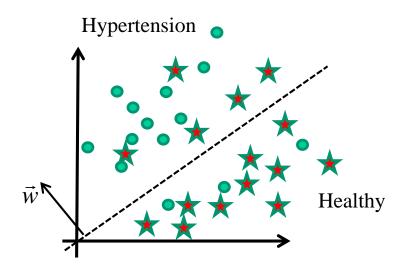
False positive (5)



False negative (3)



Confusion matrix



Ground truth

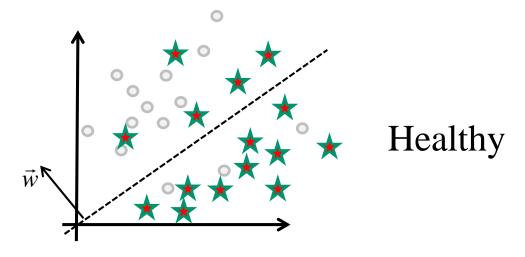
Model prediction Negative Positive

	Positive	Negative
1 OSITIVO	TP = 11	FP=5
	FN=3	TN=10

$$TP + FP = 16 = TOTAL \# of patients classified +1$$

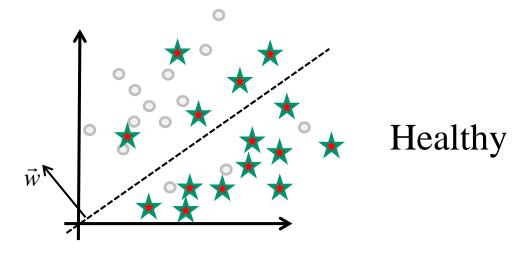
 $FN + TN = 13 = TOTAL \# of patients classified -1$

False positive rate



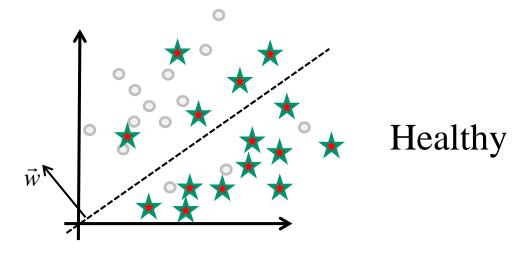
$$FPR = FP/(FP + TN) = 5/15$$

Specificity



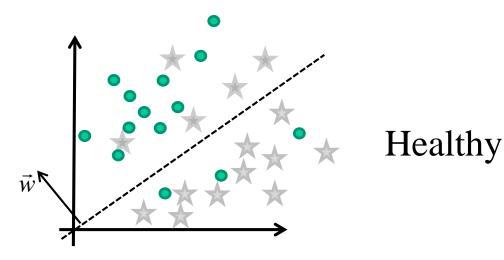
$$Sp = TN/(FP+TN)=11/15$$

Specificity



$$Sp = TN/(FP+TN)=1-FNR=11/15$$

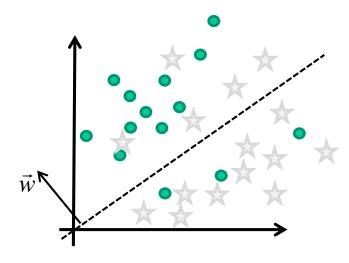
False negative rate



$$FNR = FN/(FN+TP) = 3/14$$

Recall

Hypertension

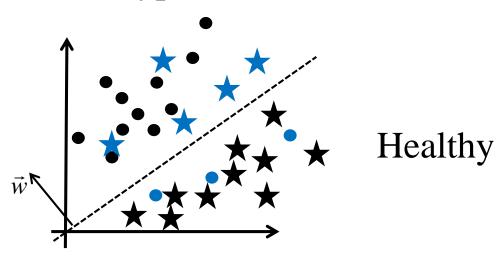


Healthy

$$Re = TP/(FN+TP)=1-FNR=11/14$$

Accuracy

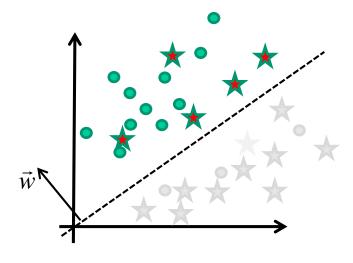
Hypertension



Rate of good classification = (TP+TN)/(FP+FN+TP+TN) = 21/29

Precision

Hypertension



Healthy

$$Pr = TP/(TP+FP) = 11/16$$

In short

Ground truth

	Positive	Negative
Model prediction egative Positive	TP = 11	FP=5
Model p	FN=3	TN=10

$$TN + FP = 15 = TOTAL \# negative$$

 $TP + FN = 14 = TOTAL \# positive$

$$TP + FP = 16 = TOTAL \# of patients classified +1$$

 $FN + TN = 13 = TOTAL \# of patients classified -1$

False positive rate =
$$FP/(FP+TN) = 5/15$$

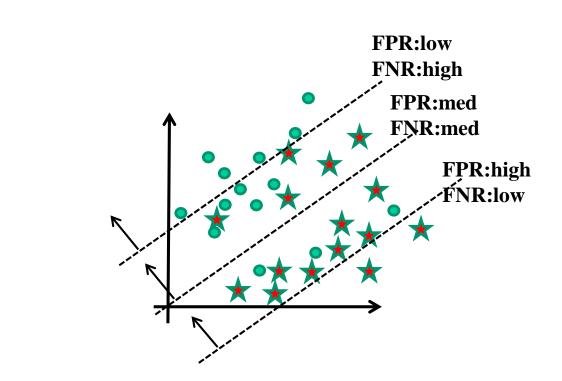
False negative rate = $FN/(FN+TP) = 3/14$

Specificity (
$$\mathbf{Sp}$$
) = TN/(FP+TN)=1-FPR=10/15
Recall (\mathbf{Re}) = TP/(TP+TN)=11/14
Precision (\mathbf{Pr}) = TP/(TP+FP) = 11/16

Accuracy =
$$(TP+TN)/(FP+FN+TP+TN) = 21/29$$

F-measure = $2* (Re*Pr)/(Pr+Re)=0.73$

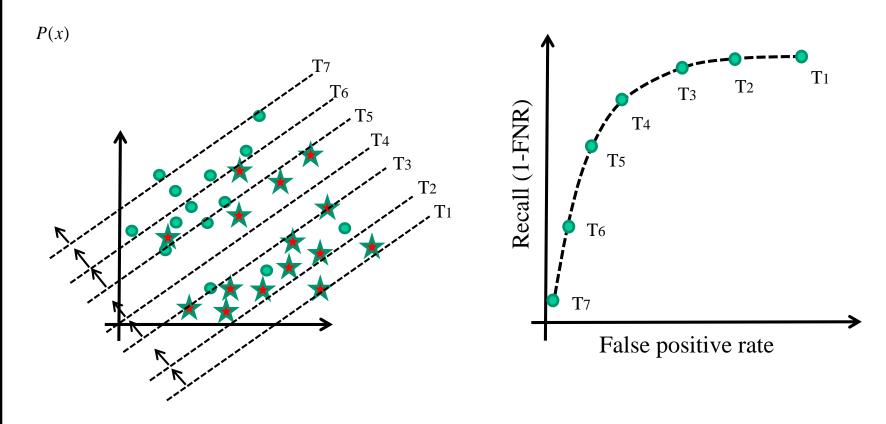
Different thresholds, different results



P(x)

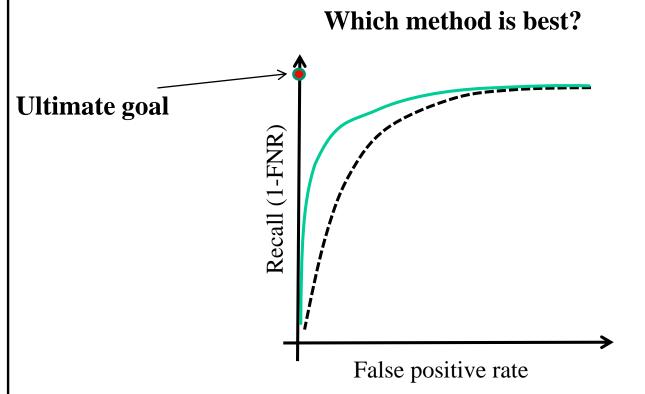
ROC curves

Compute Recall and FPR for **different thresholds**



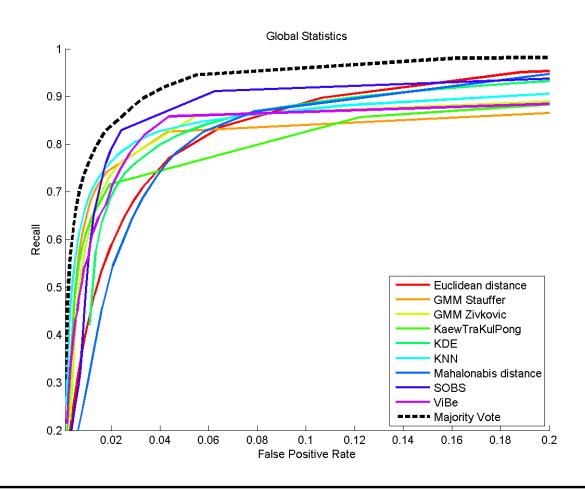
ROC curves

Good way for comparing methods



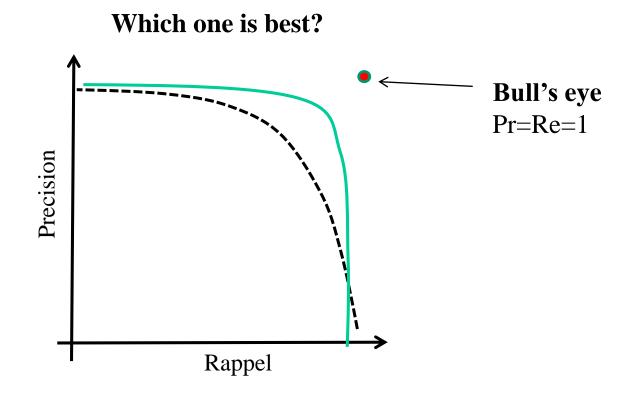
ROC curves

Example: 10 motion detection methods



Precision recall curve

Sampe spirit that the ROC curve



Model ensembling

Why use only one model?

Does combining several model works well?

In practice ... **Yes!**

Combining several models is often called ensembling

Why use only one model?

Combining what?

- Several different models
- The same model trained with different hyperparameters
- The same model trained on different data.

Typically 2 ways of combining models

- > Bagging: good for models with a large capacity
- > **Boosting**: good for models with a **low capacity**

Typically 2 ways of combining models

- > Bagging: good for models with a large capacity
- **Boosting**: good for models with a **low capacity**

Bagging

- A simple way of combining several models
 - FOR i = 1 to mo Train model $y_{W,i}(\vec{x})$
- **Ensemble** the *m* models
 - > For regression

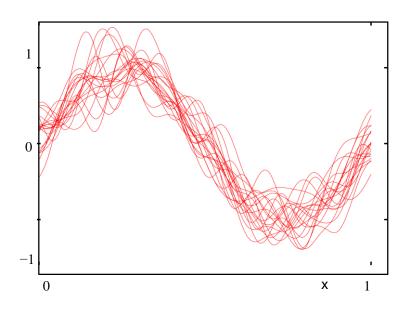
$$\circ \quad y_{COM}\left(\vec{x}\right) = \frac{1}{m} \sum_{i=1}^{m} y_{W,i}\left(\vec{x}\right)$$

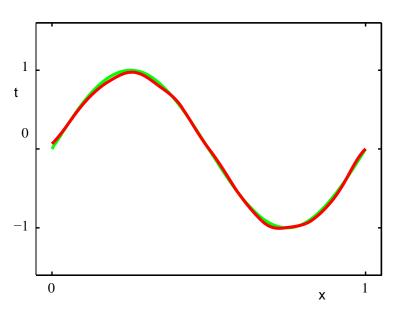
- > For a classification
 - o majority vote

Ex: polynomial regression M=25

100 models trained on 100 different training sets

Ensemble of 100 models
Vs
True model

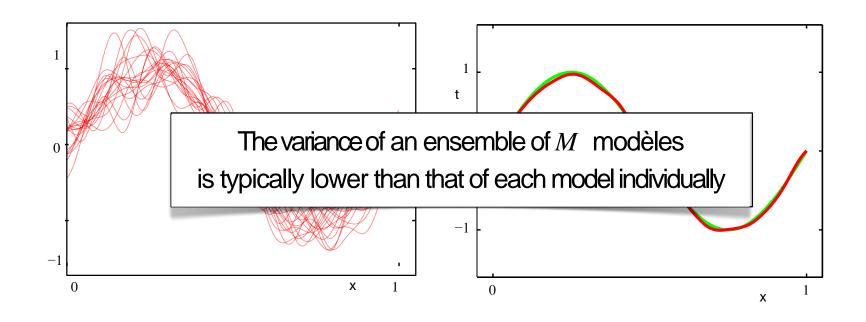




Ex: polynomial regression M=25

100 models trained on 100 different training sets

Ensemble of 100 models
Vs
True model



Bootsrap

What if you do not have enough data for building 100 trainingsets? What can you do???

Solution 1 : **Data augmentation**

Solution 2: **Bootsraping**.

FOR j from 1 to 100 DO

$$D_{bootstrap} = \{ \ \}$$

FOR N iterations

- Choose randonly a natural number between I and N

-
$$D_{bootstrap} = D_{bootstrap} \cup \{(\vec{x}_n, t_n)\}$$

$$D_{train,j} = D_{bootstrap}$$

Train j-th model

In short we have seen

- Supervised vs unsupervised learning
- Regression vs Classification
- Linear vs non-linear models
- Parameters vs hyper-parameters
- Over vs Underfitting
- Cross-validation
- Metrics
- Ensembling Bootsrapping

Thank you!