### **Multimedia Indexing and Retrieval**

#### 3. Classical machine Learning for multimedia indexing

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### Learning

- Machine learning: learning from data.
- Unsupervised learning:
  - Without human intervention,
  - Simple data,
  - Automatic class extraction (clustering).
- Supervised learning:
  - With human intervention (annotation),
  - Labeled (or annotated) data
  - Classification (predefined classes),
  - Regression (continuous values).

### **Supervised learning**

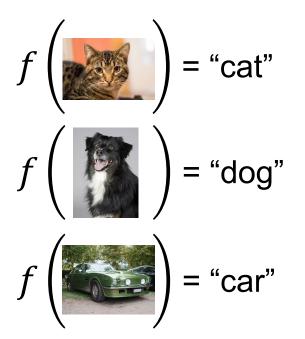
- A machine learning technique for creating a function from training data.
- The training data consist of pairs of input objects (typically vectors) and desired outputs.
- The output of the function can be a continuous value (regression) or a class label (classification) of the input object.
- The task of the supervised learner is to predict the value of the function for any valid input object after having seen a number of training examples (i.e. pairs of input and target output).
- To achieve this, the learner has to generalize from the presented data to unseen situations in a "reasonable" way.
- The parallel task in human and animal psychology is often referred to as concept learning (in the case of classification).
- Most commonly, supervised learning generates a global model that helps mapping input objects to desired outputs.

#### Learning a target function

• Target function:  $f: X \to Y$ 

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

- -x: input object, e.g., color image
- -y: desired output, e.g., class label or image tag
- X: set of valid input objects
- Y: set of possible output values



Set of possible color images:

$$X = \bigcup_{(w,h)\in\mathbb{N}^{*2}} [0,1]^{w\times h\times 3}$$

Set of possible image tags:

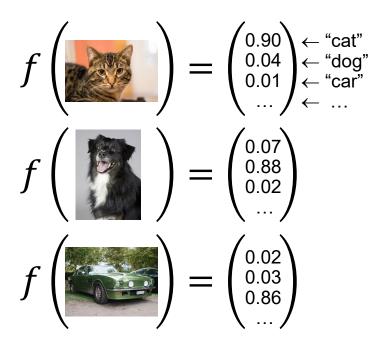
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#### Learning a target function

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Set of possible color images:

$$X = \bigcup_{(w,h)\in\mathbb{N}^{*2}} [0,1]^{w\times h\times 3}$$

Set of possible tag scores:

$$Y = \mathbb{R}^{|\{\text{``cat",``dog"}...\}|} = \mathbb{R}^{C}$$

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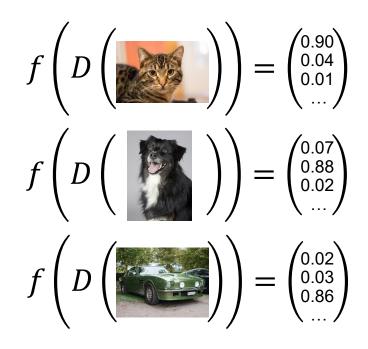
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#### Learning a target function

• Target function:  $f: X \to Y$ 

 $x \rightarrow y = f(x)$ 

- -x: input object, e.g., image descriptor
- -y: desired output, e.g., class label or image tag
- -X: set of valid input objects
- Y : set of possible output values



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Set of possible image descriptors:

 $X = \mathbb{R}^d$  (or subset of it)

Set of possible tag scores:

 $Y = \mathbb{R}^{c}$ 

D is a predefined and fixed function  $[0,1]^{w \times h \times 3}$  to  $\mathbb{R}^d$ from  $(w,h) \in \mathbb{N}^{*2}$ 2023-2024

### Learning from training data

- Training data:  $S = (x_i, y_i)_{(1 \le i \le I)}$ - *I*: number of training samples
- Learning algorithm:  $L: (X \times Y)^* \to Y^X$  $S \to f = L(S)$

$$((X \times Y)^* = \bigcup_{n \in N} (X \times Y)^n)$$
  
 $Y^X$ : set of all applications from  $X$  to  $Y$ 

• Regression or classification system: y = f(x) = [L(S)](x) = g(S, x)

### **Supervised learning**

• Target function:  $f: X \to Y$ 

 $x \to y = f(x)$ 

- -x: input object (typically vector)
- -y: desired output (continuous value or class label)
- X: set of valid input objects
- Y: set of possible output values
- Training data:  $S = (x_i, y_i)_{(1 \le i \le I)}$ 
  - *I* : number of training samples
- Learning algorithm:  $L : (X \times Y)^* \to Y^X$  $S \to f = L(S)$
- Regression or classification system:

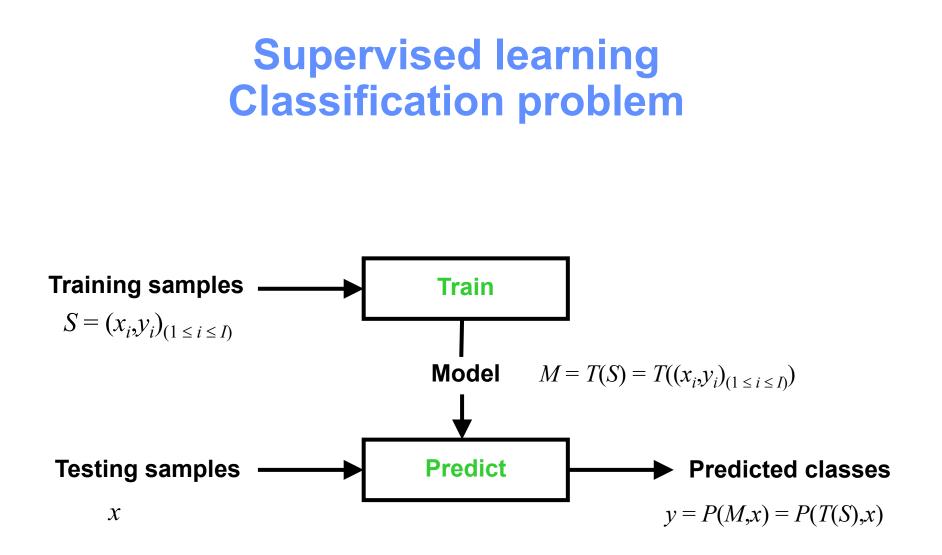
y = f(x) = [L(S)](x) = g(S, x)

### **Two types of functions**

- Target function:  $f: X \rightarrow Y$  $x \rightarrow y = f(x)$ 
  - maps input objects to desired outputs
  - often determined by a set of parameters
  - the function or its parameter are learnt from a training set
- Learning algorithm:  $L : (X \times Y)^* \to Y^X$  $S \to f = L(S)$ 
  - maps training sets to target functions
  - often controlled by a set of hyper-parameters
  - hyper-parameters may be tuned on a validation set

### Model based supervised learning

- Two functions, "train" and "predict", cooperating via a Model
- General regression or classification system:
  y = [L(S)](x) = g(S,x)
- Building of a model (train): M = T(S)
- Prediction using a model (predict): y = [L(S)](x) = g(S,x) = P(M,x) = P(T(S),x)



### **Classification methods**

- Gaussian Mixture Models
- Hidden Markov Models
- Decision trees
- Genetic algorithms
- Artificial neural networks
- K-nearest neighbor
- Linear discriminant analysis
- Support vector machines
- Minimum message length
- And many more.

#### k nearest neighbors (k-NN)

- No training : M = T(S) = S (T = identity)
- Compute the distances from the unknown sample *x* to all the training samples *x<sub>i</sub>*,
- Select the k closest  $x_i$ ,
- Compute the class of x from the classes of the closest x<sub>i</sub>'s:
  - -k = 1: the class of x is the class of the closest  $x_i$ ,
  - -k is odd and there are only two classes : majority vote.
- *k*-NN is a non linear classifier and can easily model classes with very irregular shapes,

#### k nearest neighbors (k-NN)

- 1-NN is a simple and quite often excellent classifier, it is often chosen as a baseline for comparison between systems,
- 3-NN is more robust against isolated outliers,
- Improvement: weight class values according to the (inverse) distance to the query point
- May be slow for classification because of the need to compute the distances with all the training samples
- However a single NN search may be performed for many classifications at once (multi-label problem)
- May be used for indexing (off-line) or for search (on-line, "similarity search")

### **Computation of distance for k-NN**

- Euclidian distance, angle between vectors,
- Comparison between a query vector to all the vectors in the database (no pre-selection),
- "Small" number of dimensions ( < 10) : clustering techniques, hierarchical search,
- "Medium" number of dimensions (~ 10+): methods based on space partitioning,
- "Large" number of dimensions( >> 10 ) : no known method faster that a full linear scan,
- Reduction of the number of dimensions by Principal Component Analysis.
- Approximate Nearest Neighbors: LSH

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## Locality Sensitive Hashing (LSH)

- Hashing: store many data samples into a table of fixed length; data placed into "buckets"
- "Regular" Hashing: avoid collision for faster access, polynomial and multiples XOR functions; any type of data
- Locality preserving hashing: favor collisions of "close" samples into the same buckets; data from highly dimensional Euclidean space, multiple projection functions

### **LSH: Multiple projection functions**

- Set of random directions
- Projection on the axes  $\rightarrow$  one component per direction
- Split values on axes according to a data distribution (two, four, eight ... intervals)
- One or more bits per direction (generally one)
- Concatenation for producing the bucket index
- Multiple projections: matrix vector multiplication

#### **LSH: Use of multiple tables**

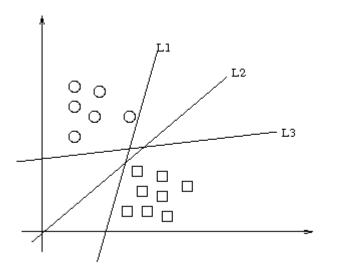
- Build many LSH tables
- For each table, select all the test samples that fall in the same bucket than the query sample
- Compute the Euclidean distance only for those samples
- Sort the test samples according to the Euclidean distances
- Euclidean distances are not approximate but some samples close to the query may not fall in the selected buckets
- The size and number of tables must be chosen so that enough and not too many samples are found for a query

### LSH: Use of hamming distance

- Build binary codes (bucket index) as for one LSH table
- Hamming distance: number of bit locations in which the binary values differ: bitwise XOR followed by a count on 1 bits; modern processors have this as a single instruction
- Compute the Hamming distance between the query and all test samples: much faster than Euclidean distance
- Select samples with closest Hamming distance
- Compute the Euclidean distance only for those samples
- Similar to multiple tables from there

# **Support Vector Machines (SVM)**

- Empirical risk minimization
- Linear classifier with maximum margin



• The "kernel trick" permits non linear classification also with maximum margin and minimum empirical risk

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# **SVM linear classification**

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- Maximum-margin hyperplanes for a SVM trained with samples from two classes.
- Samples along the hyperplanes are called the support vectors.
- The separated hyperplane is defined by:

 $w^T x - b = 0$ 

• The margin is 2/|w|

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### **SVM linear classification**

• If the data is linearly separable:

if  $y_i = -1$ :  $w^T x_i - b \le -1$  if  $y_i = +1$ :  $w^T x_i - b \ge 1$ 

• This can be rewritten as:

$$y_i \cdot (w^T \cdot x_i - b) \ge 1$$

• SVM problem primal form:

Minimize: 
$$\frac{1}{2} \|w\|^2$$
 subject to:  $y_i \cdot (w^T \cdot x_i - b) \ge 1$ ,  $1 \le i \le n$ .

• SVM problem dual form:  $w = \sum_{i=1}^{n} \alpha_i y_i x_i$ 

maximize: 
$$\sum_{i=1}^{n} \alpha_i - \sum_{i=1}^{n} \sum_{j=1}^{n} \alpha_i \alpha_j y_i y_j x_i^T x_j$$
 subject to  $\alpha_i \ge 0$ 

 $\alpha_i$  's are non zero only for the support vectors.

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## **SVM linear classification**

• Soft margin, primal form:

$$y_i.(w^T.x_i-b) \ge 1 \quad \rightarrow \quad y_i.(w^T.x_i-b) \ge 1-\xi_i$$
$$\min \frac{1}{2} \|w\|^2 \quad \rightarrow \quad \min \left(\frac{1}{2} \|w\|^2 + C\sum_{i=1}^n \xi_i\right)$$

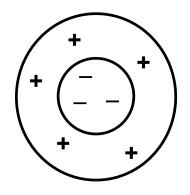
• Dual form:

$$\alpha_i \ge 0 \qquad \longrightarrow \quad 0 \le \alpha_i \le C$$

• Allows for "misclassified" samples.

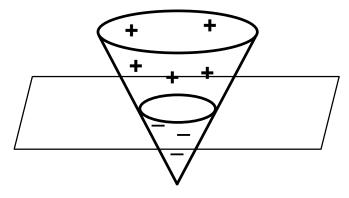
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#### **SVM non-linear classification**



No linear separation in original space

• Kernel trick: projection on a cone  $(2D \rightarrow 3D)$ :  $(x, y) \rightarrow \Phi(x, y) = (x, y, \sqrt{x^2 + y^2})$ 



Linear separation in  $im(\Phi)$  space

#### **SVM non-linear classification**

• Decision function:

$$f(x) = \langle w | x \rangle - b = \left\langle \sum_{i=1}^{n} \alpha_{i} y_{i} x_{i} | x \right\rangle - b = \left( \sum_{i=1}^{n} \alpha_{i} y_{i} \langle x_{i} | x \rangle \right) - b$$

• Quadratic form maximization:

$$\sum_{i=1}^{n} \alpha_{i} - \sum_{i=1}^{n} \sum_{j=1}^{n} \alpha_{i} \alpha_{j} y_{i} y_{j} \langle x_{i} | x_{j} \rangle$$

- Kernel trick:  $\langle x_i | x_j \rangle \rightarrow \langle \Phi(x_i) | \Phi(x_j) \rangle = K(x_i, x_j)$
- Φ : possibly non-linear function, does not need to be computed, implicitly defined via the kernel (K) definition, linear separation in the im(Φ) space, may be non linear in the original space.

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### **SVM non-linear classification**

- Mercer condition :  $K(x_i, x_j)$  must be definite positive.
- Common kernels:
  - Polynomial (homogeneous):  $K(x, y) = (x.y)^d$
  - Polynomial (inhomogeneous):  $K(x, y) = (x.y+1)^d$

- Radial Basis Function: 
$$K(x, y) = \exp\left(-\frac{\|x - y\|^2}{2\sigma^2}\right)$$

- Sigmoid:  $K(x, y) = \tanh(\kappa x. y + c)$ , for some (not every)  $\kappa > 0$  and c < 0

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## **SVM summary**

- Maximization of the margin for linearly separable data
- Use of a dual form for finding support vectors and coefficients (convex optimization)
- Use of soft margin for "almost" linearly separable data
- Use of the "kernel trick" for non-linearly separable data
- Most commonly used kernel:  $K(x, y) = e^{-\gamma ||x-y||^2} \rightarrow f(x) = \sum_{i=1}^{i=I} \propto_i y_i e^{-\gamma ||x-x_i||^2} + b$ : weighted sum of Gaussians centered on the support samples (vectors)

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#### **Hyper-parameter tuning**

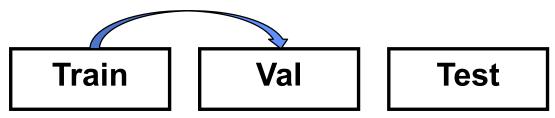
- Parameters:
  - Parameters of the model learnt from training data
  - e.g. values of the support vectors  $(\boldsymbol{x}_i)$  and Lagrange coefficients  $(\alpha_i)$  in SVMs
- "Hyper"-parameters:
  - Parameters that controls how the model (and "standard" parameters) are learnt
  - e.g. soft margin coefficient (C) in SVMs and the scale parameter in the RBF version ( $\gamma$ )
  - Possibly also class weights
  - Controls "how well" the classification algorithm generalizes from training data, especially the under fit versus over fit compromise

#### Hyper-parameter tuning, validation set

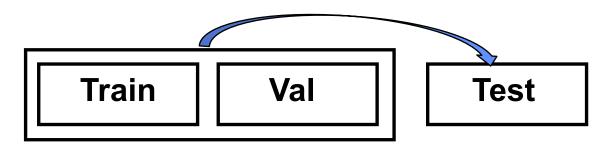
- A dataset used for training cannot be used for evaluation (over-fitting)
- Standard method: use different datasets for training and performance evaluation, each with annotated samples.
- Tuning of hyper-parameters on the test set is bad (over-fitting again)
- Good solution: use three datasets: train, val and test, all with annotated samples
- Train and evaluate several hyper-parameter values between train and val and then apply to test.

#### Hyper-parameter tuning, validation set

 Parameter tuning: selection of the optimal hyperparameter combination by training on train and evaluating on val.

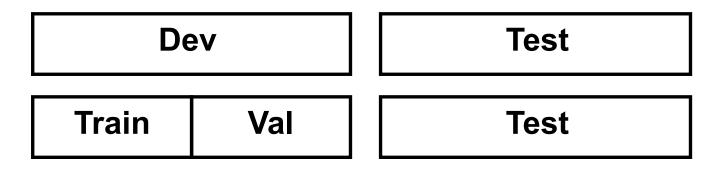


 Actual evaluation: keep the optimal hyper-parameter values, train on train+val and evaluate on test.

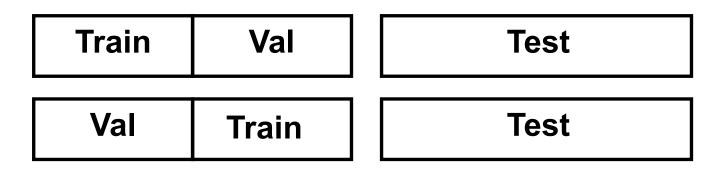


#### No validation set: split the training set

• Split into two equal parts, use first part as train and second part for validation ("one-fold" cross-validation)



Two-fold cross-validation



#### **Two-fold cross-validation**

• Use two parts alternatively for training and validation

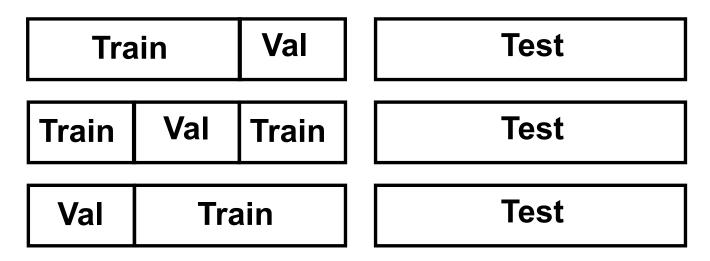
Train	Val	Test
Val	Train	Test

- The whole development set is used both for training and for evaluation during hyper-parameter tuning
- Tuning is done on MAP (hyper-parameters)
  - Either average the MAP on the two validations
  - Or compute a global MAP on the concatenated scores
- Training is done on half of the development set each time

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#### **N-fold cross-validation**

 Use N parts of 1/N od the development set alternatively for validation and the complement ((N-1)/N) for training



- The whole development set is used both for training and for evaluation during hyper-parameter tuning
- Training is done on (N-1)/N of the development set each time, the greater N, the better.

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#### **Probabilized output**

- SVM scores possibly ranges from  $-\infty$  to  $+\infty$
- Probabilities are expected to range from 0 to 1
- Sigmoid transform: p(score) = 1/(1+e<sup>(A\*score+B)</sup>)
- Additional hint: among the samples within a small interval around p, a fraction of about p would have positive labels
- Platt's (1999) method: learn A and B by cross-validation to optimally satisfy the above hint
- Probability normalized outputs better for late fusion
- Linear SVM + sigmoid normalization ~ logistic regression