Classification fundamentals



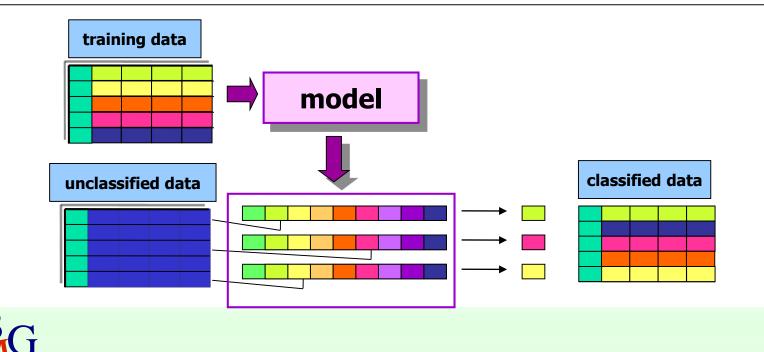
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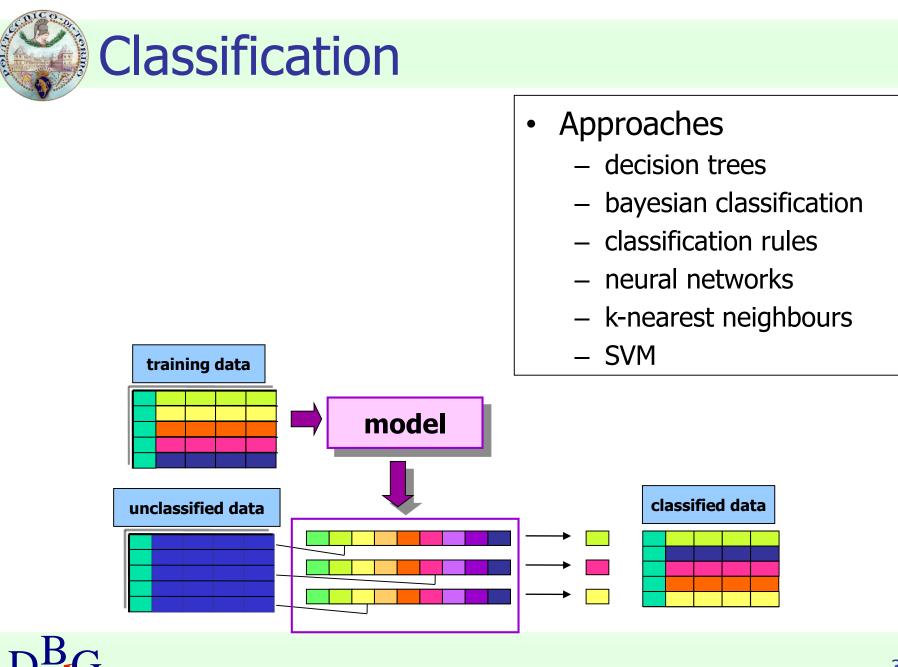


Classification

Objectives

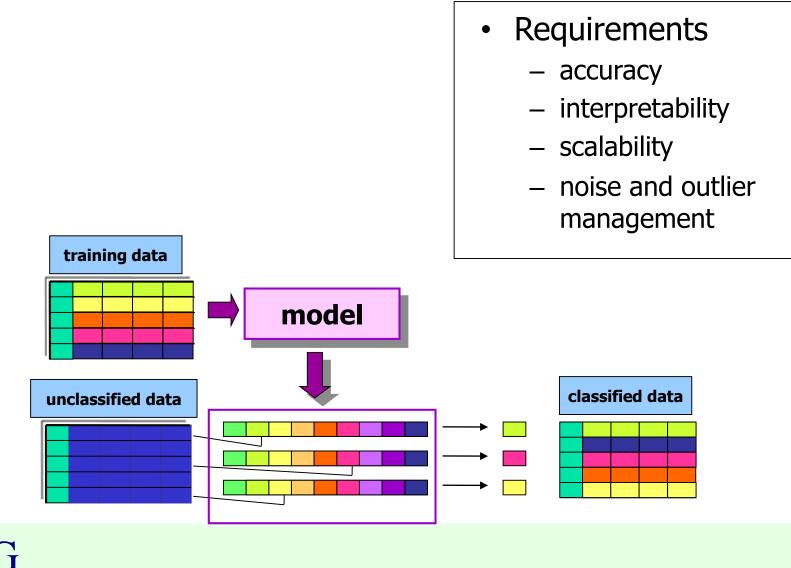
- prediction of a class label
- definition of an interpretable model of a given phenomenon









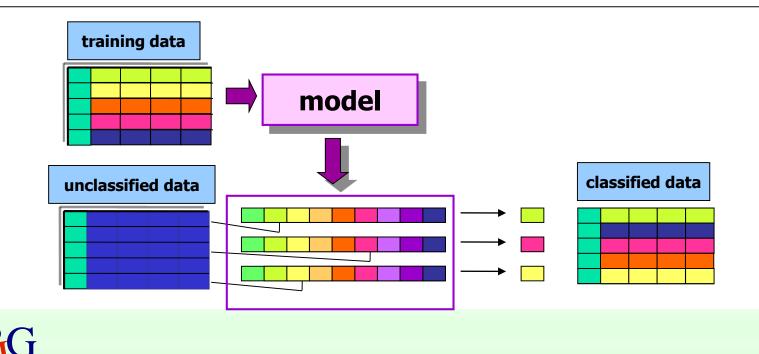




Classification

- Applications
 - detection of customer propension to leave a company (churn or attrition)
 - fraud detection
 - classification of different pathology types

...



Classification: definition

- Given
 - a collection of class labels
 - a collection of data objects labelled with a class label
- Find a descriptive profile of each class, which will allow the assignment of unlabeled objects to the appropriate class





Definitions

Training set

- Collection of labeled data objects used to learn the classification model
- Test set
 - Collection of labeled data objects used to validate the classification model





Classification techniques

- Decision trees
- Classification rules
- Association rules
- Neural Networks
- Naïve Bayes and Bayesian Networks
- k-Nearest Neighbours (k-NN)
- Support Vector Machines (SVM)



Evaluation of classification techniques

- Accuracy
 - quality of the prediction
- Efficiency
 - model building time
 - classification time
- Scalability
 - training set size
 - attribute number
- Robustness
 - noise, missing data
- Interpretability
 - model interpretability
 - model compactness



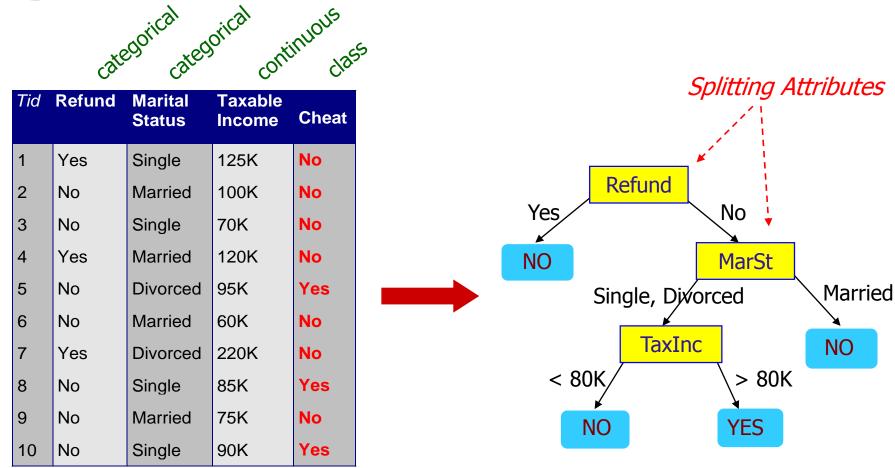
Decision trees



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Example of decision tree



Model: Decision Tree

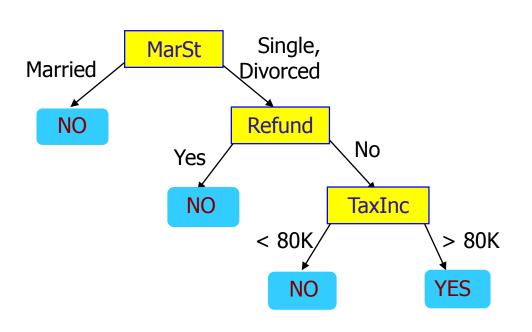
Training Data



Another example of decision tree



Tid	Refund	Marital Status	Taxable Income	Cheat
1	Yes	Single	125K	No
2	No	Married	100K	No
3	No	Single	70K	No
4	Yes	Married	120K	No
5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
8	No	Single	85K	Yes
9	No	Married	75K	No
10	No	Single	90K	Yes

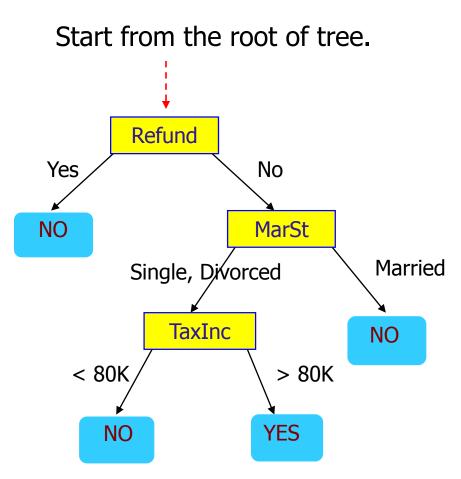


There could be more than one tree that fits the same data!





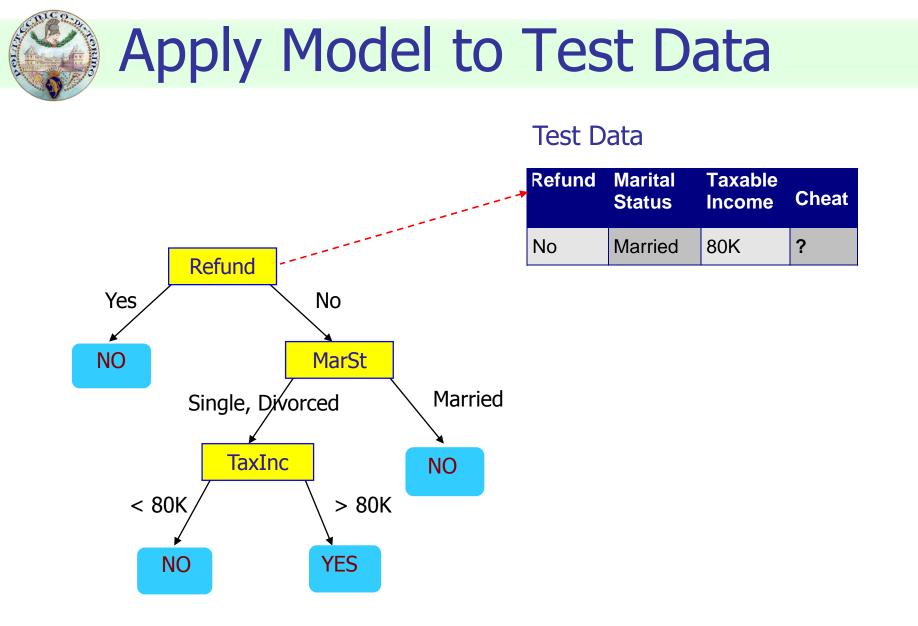
Apply Model to Test Data



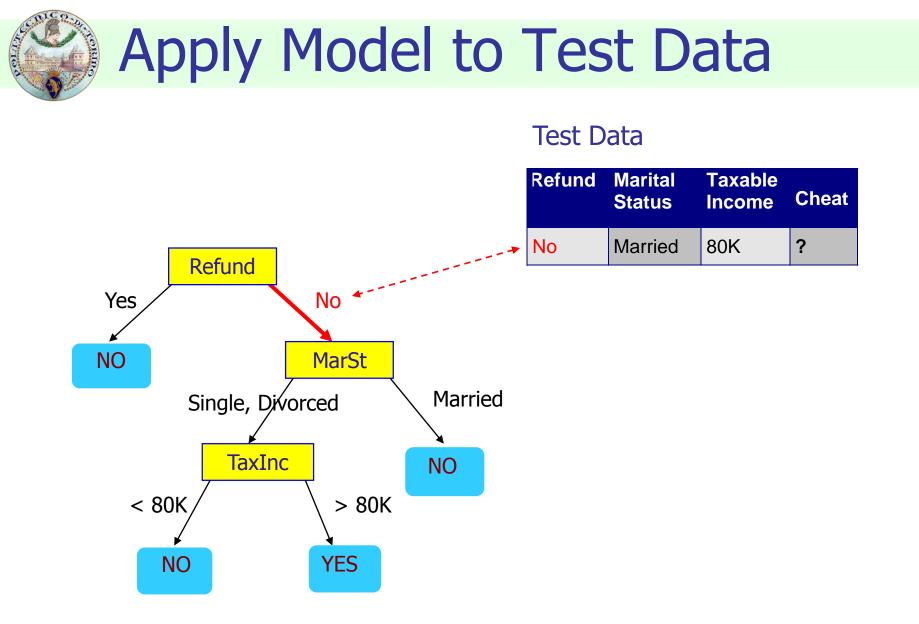
Test Data

Refund		Taxable Income	Cheat
No	Married	80K	?

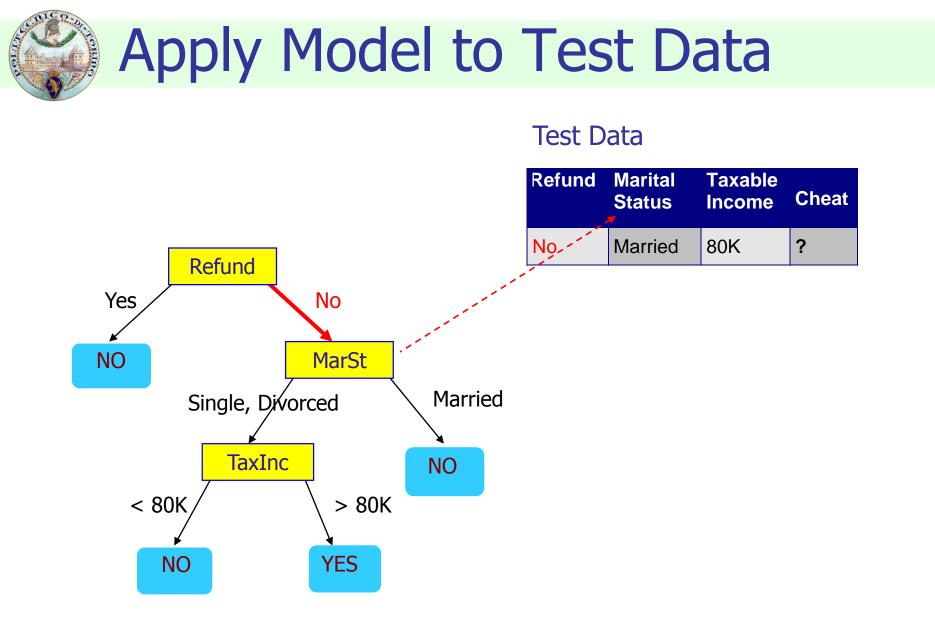


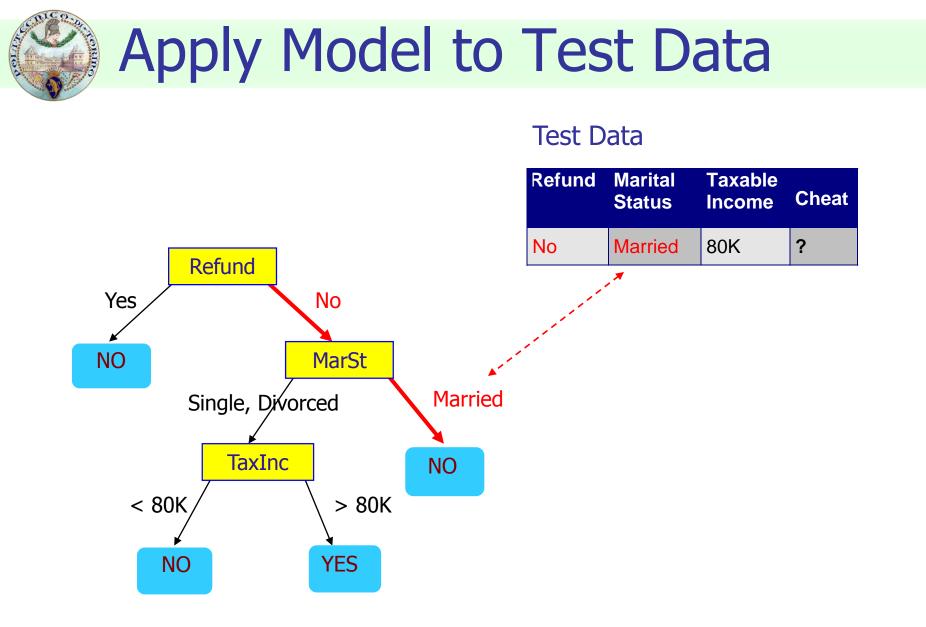


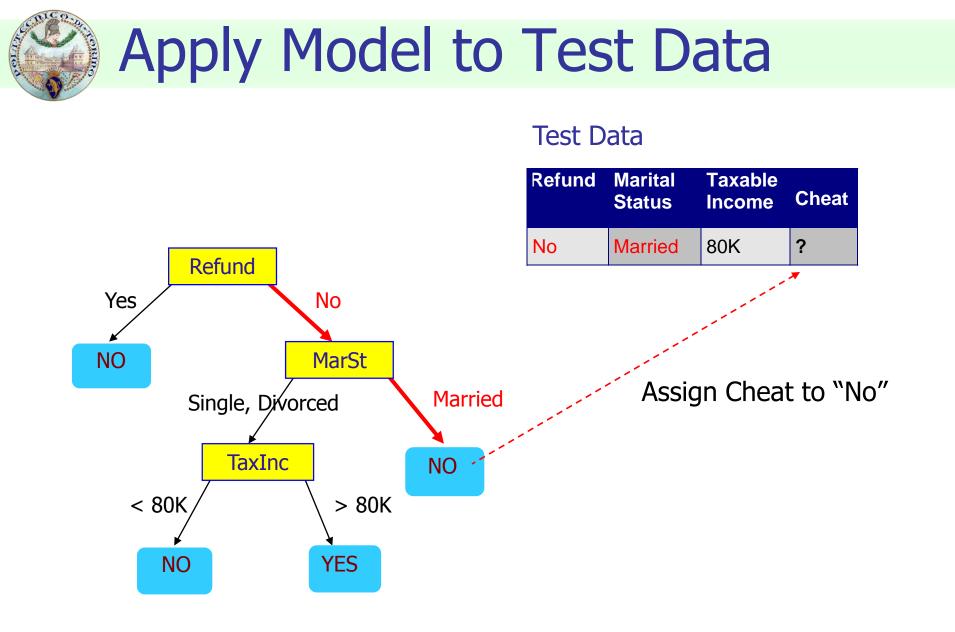
 $D_{M}^{B}G$











Decision tree induction

Many algorithms to build a decision tree

- Hunt's Algorithm (one of the earliest)
- CART
- ID3, C4.5, C5.0
- SLIQ, SPRINT

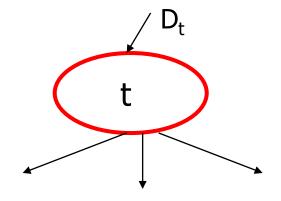


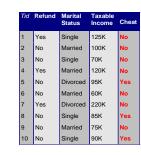
General structure of Hunt's algorithm

Basic steps

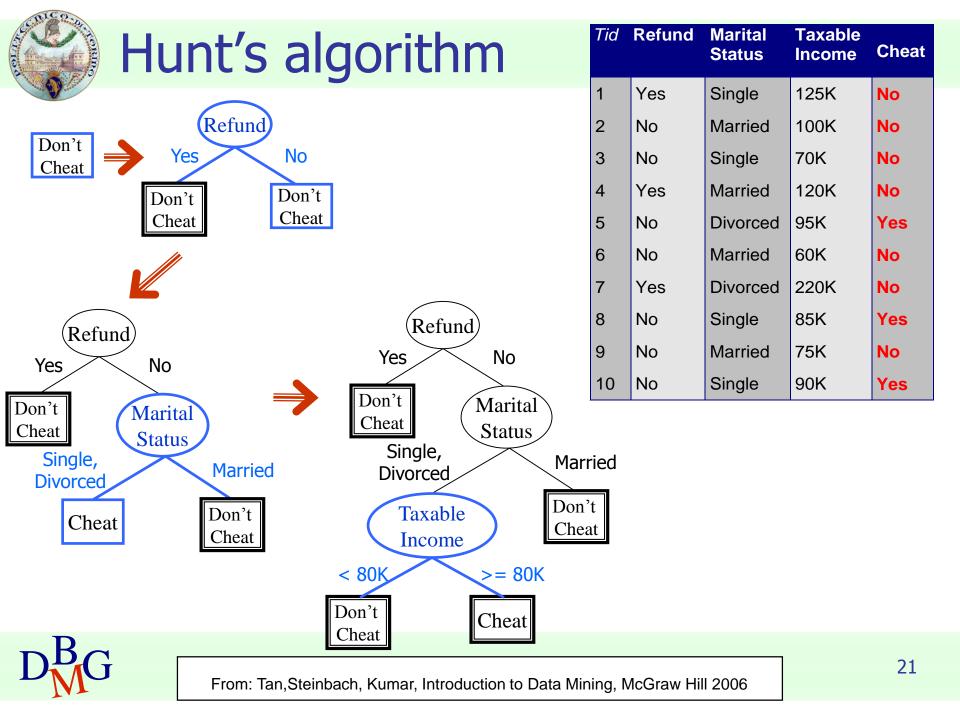
- If D_t contains records that belong to more than one class
 - select the "best" attribute A on which to split D_t and label node t as A
 - split D_t into smaller subsets and recursively apply the procedure to each subset
- If D_t contains records that belong to the same class y_t
 - then t is a leaf node labeled as y_t
- If D_t is an empty set
 - then t is a leaf node labeled as the default (majority) class, y_d

 $D_{t,r}$ set of training records that reach a node t











Decision tree induction

Adopts a greedy strategy

- "Best" attribute for the split is selected locally at each step
 - not a global optimum
- Issues
 - Structure of test condition
 - Binary split versus multiway split
 - Selection of the best attribute for the split
 - Stopping condition for the algorithm





Structure of test condition

Depends on attribute type

- nominal
- ordinal
- continuous

Depends on number of outgoing edges

- 2-way split
- multi-way split



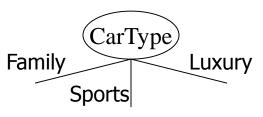


Splitting on nominal attributes

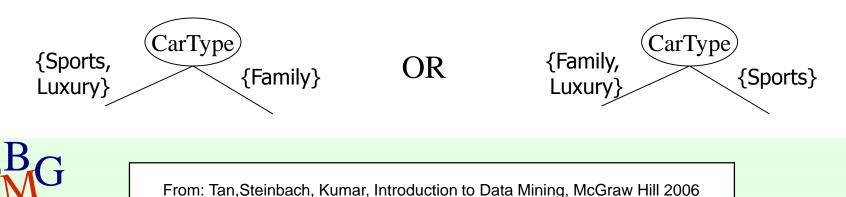
Multi-way split

use as many partitions as distinct values

Binary split



- Divides values into two subsets
- Need to find optimal partitioning





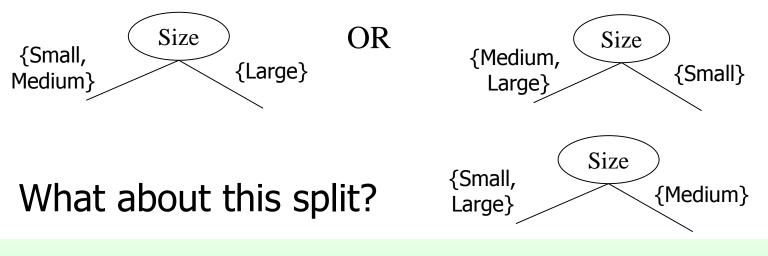
Splitting on ordinal attributes

Multi-way split

use as many partitions as distinct values

Binary split

- Divides values into two subsets
- Need to find optimal partitioning



Large

Size

Medium

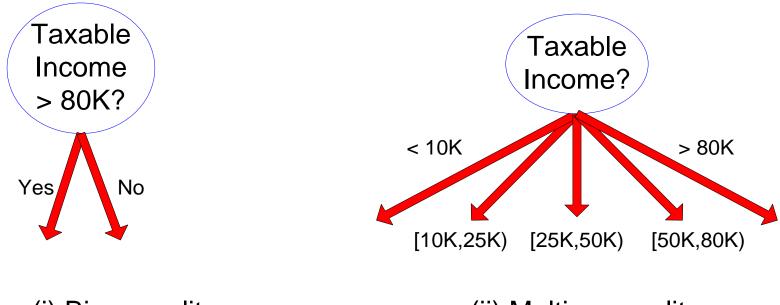
Small

Splitting on continuous attributes

Different techniques

- Discretization to form an ordinal categorical attribute
 - Static discretize once at the beginning
 - Dynamic discretize during tree induction
 - Ranges can be found by equal interval bucketing, equal frequency bucketing (percentiles), or clustering
- Binary decision (A < v) or (A \ge v)
 - consider all possible splits and find the best cut
 - more computationally intensive





(i) Binary split

(ii) Multi-way split

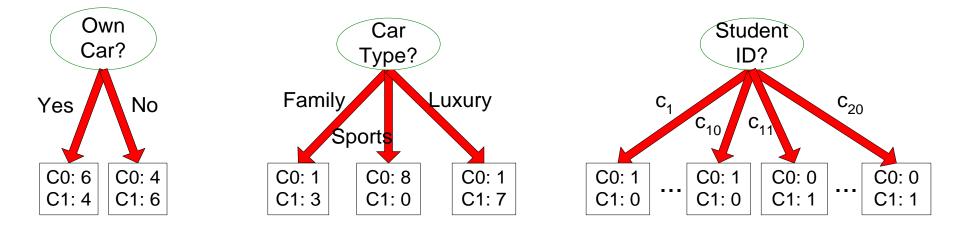




Selection of the best attribute

Before splitting:

10 records of class 0, 10 records of class 1



Which attribute (test condition) is the best?





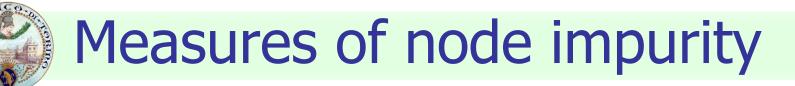
Selection of the best attribute

- Attributes with *homogeneous* class distribution are preferred
- Need a measure of node impurity

Non-homogeneous, high degree of impurity

Homogeneous, low degree of impurity

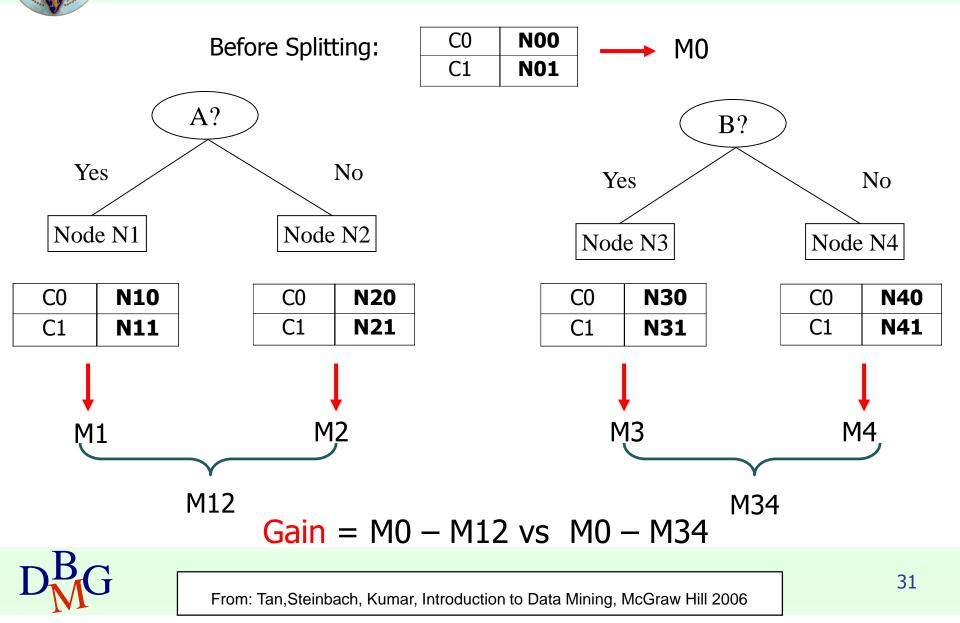




- Many different measures available
 - Gini index
 - Entropy
 - Misclassification error
- Different algorithms rely on different measures



How to find the best attribute





Gini Index for a given node t

$$GINI(t) = 1 - \sum_{j} [p(j | t)]^{2}$$

p(j | t) is the relative frequency of class j at node t

- Maximum (1 1/n_c) when records are equally distributed among all classes, implying higher impurity degree
- Minimum (0.0) when all records belong to one class, implying lower impurity degree

Gini=	0.000	Gini=	0.278	Gini=		Gin	i=0.500
C2	6	C2	5	C2	4	C2	3
C1	0	C1	1	C1	2	C1	3





Examples for computing GINI

$$GINI(t) = 1 - \sum_{j} [p(j | t)]^{2}$$

C1	0
C2	6

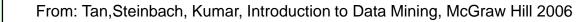
P(C1) = 0/6 = 0	P(C2) = 6/6 = 1
Gini = $1 - P(C1)^2$ -	$-P(C2)^2 = 1 - 0 - 1 = 0$

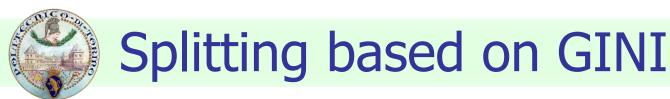
C1	1
C2	5

$$P(C1) = 1/6$$
 $P(C2) = 5/6$
Gini = 1 - (1/6)² - (5/6)² = 0.278

C1	2
C2	4

$$P(C1) = 2/6$$
 $P(C2) = 4/6$
Gini = 1 - (2/6)² - (4/6)² = 0.444





- Used in CART, SLIQ, SPRINT
- When a node p is split into k partitions (children), the quality of the split is computed as

$$GINI_{split} = \sum_{i=1}^{k} \frac{n_i}{n} GINI(i)$$

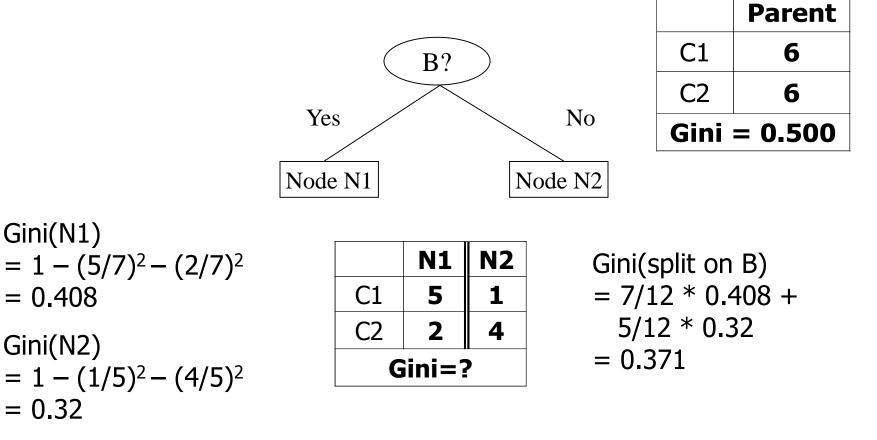
where

- n_i = number of records at child i
- n = number of records at node p



Computing GINI index: Boolean attribute

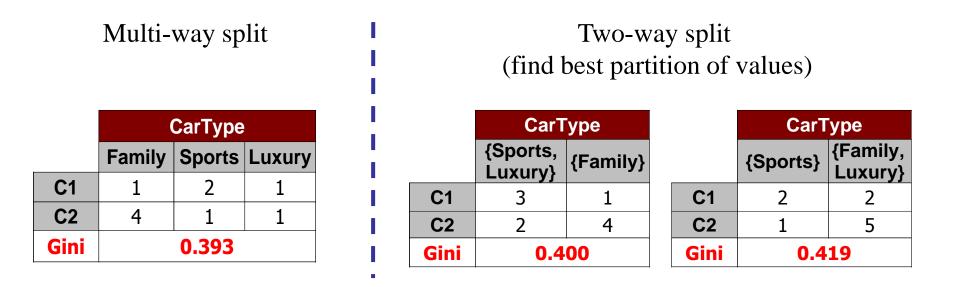
- Splits into two partitions
 - larger and purer partitions are sought for





Computing GINI index: Categorical attribute

- For each distinct value, gather counts for each class in the dataset
- Use the count matrix to make decisions



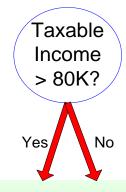




Computing GINI index: Continuous attribute

- Binary decision on one splitting value
 - Number of possible splitting values
 - = Number of distinct values
- Each splitting value v has a count matrix
 - class counts in the two partitions
 - A < v
 - A ≥ v

Tid	Refund	Marital Status	Taxable Income	Cheat
1	Yes	Single	125K	No
2	No	Married	100K	No
3	No	Single	70K	No
4	Yes	Married	120K	No
5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
8	No	Single	85K	Yes
9	No	Married	75K	No
10	No	Single	90K	Yes





Computing GINI index: Continuous attribute

For each attribute

- Sort the attribute on values
- Linearly scan these values, each time updating the count matrix and computing gini index
- Choose the split position that has the least gini index

	Cheat		No		No)	N	0	Ye	S	Ye	S	Ye	es	N	0	N	0	N	0		No	
Taxable								ble Income															
Sorted Values			60		70)	7	5	85	5	9()	9	5	10)0	12	20	1:	25		220	
Split Positions		5	5	6	5	7	2	8	0	8	7	9	2	9	7	11	10	1:	22	17	72	23	0
opine i obicionio		۳	>	<=	>	<=	>	<=	>	<=	>	<=	>	<=	>	<=	>	<=	>	<=	>	<=	>
	Yes	0	3	0	3	0	3	0	3	1	2	2	1	3	0	3	0	3	0	3	0	3	0
	No	0	7	1	6	2	5	3	4	3	4	3	4	3	4	4	3	5	2	6	1	7	0
Gini 0.420 0.400 0.375 0.343 0.417 0.400						100	<u>0.3</u>	<u>300</u>	0.3	343	0.3	575	0.4	00	0.4	20							





Entropy at a given node t

$$Entropy(t) = -\sum_{j} p(j | t) \log_{2} p(j | t)$$

p(j | t) is the relative frequency of class j at node t

- Maximum (log n_c) when records are equally distributed among all classes, implying higher impurity degree
- Minimum (0.0) when all records belong to one class, implying lower impurity degree
- Entropy based computations are similar to GINI index computations



Examples for computing entropy

$$Entropy(t) = -\sum_{j} p(j|t) \log_{2} p(j|t)$$

C1	0
C2	6

P(C1) = 0/6 = 0	P(C2) = 6/6 = 1
Entropy $= -0 \log$	$0 - 1 \log 1 = -0 - 0 = 0$

C1	1
C2	5

 $P(C1) = 1/6 \qquad P(C2) = 5/6$ Entropy = - (1/6) log₂ (1/6) - (5/6) log₂ (5/6) = 0.65

C1	2
C2	4

P(C1) = 2/6 P(C2) = 4/6 Entropy = $-(2/6) \log_2 (2/6) - (4/6) \log_2 (4/6) = 0.92$





Information Gain

$$GAIN_{split} = Entropy(p) - \left(\sum_{i=1}^{k} \frac{n_i}{n} Entropy(i)\right)$$

Parent Node, p is split into k partitions;

n_i is number of records in partition i

- Measures reduction in entropy achieved because of the split. Choose the split that achieves most reduction (maximizes GAIN)
- Used in ID3 and C4.5
- Disadvantage: Tends to prefer splits yielding a large number of partitions, each small but pure





Splitting Based on INFO

Gain Ratio

$$GainRATIO_{split} = \frac{GAIN_{split}}{SplitINFO}$$

$$SplitINFO = -\sum_{i=1}^{k} \frac{n_i}{n} \log \frac{n_i}{n}$$

Parent Node, p is split into k partitions n_i is the number of records in partition i

- Adjusts Information Gain by the entropy of the partitioning (SplitINFO). Higher entropy partitioning (large number of small partitions) is penalized
- Used in C4.5
- Designed to overcome the disadvantage of Information Gain





Classification error at a node t

$$Error(t) = 1 - \max_{i} P(i \mid t)$$

Measures misclassification error made by a node

- Maximum (1 1/n_c) when records are equally distributed among all classes, implying least interesting information
- Minimum (0.0) when all records belong to one class, implying most interesting information





Examples for computing error

$$Error(t) = 1 - \max_{i} P(i \mid t)$$

C1	0
C2	6

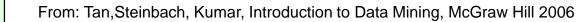
$$P(C1) = 0/6 = 0$$
 $P(C2) = 6/6 = 1$
Error = 1 - max (0, 1) = 1 - 1 = 0

C1	1
C2	5

$$P(C1) = 1/6$$
 $P(C2) = 5/6$
Error = 1 - max (1/6, 5/6) = 1 - 5/6 = 1/6

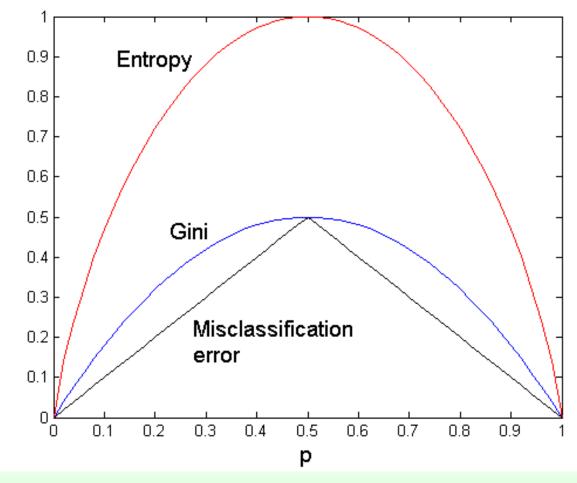
C1	2
C2	4

$$P(C1) = 2/6$$
 $P(C2) = 4/6$
Error = 1 - max (2/6, 4/6) = 1 - 4/6 = 1/3





For a 2-class problem



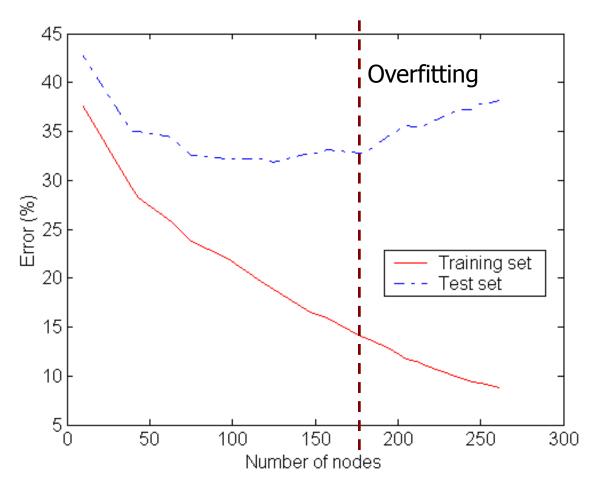




Stopping Criteria for Tree Induction

- Stop expanding a node when all the records belong to the same class
- Stop expanding a node when all the records have similar attribute values
- Early termination
 - Pre-pruning
 - Post-pruning

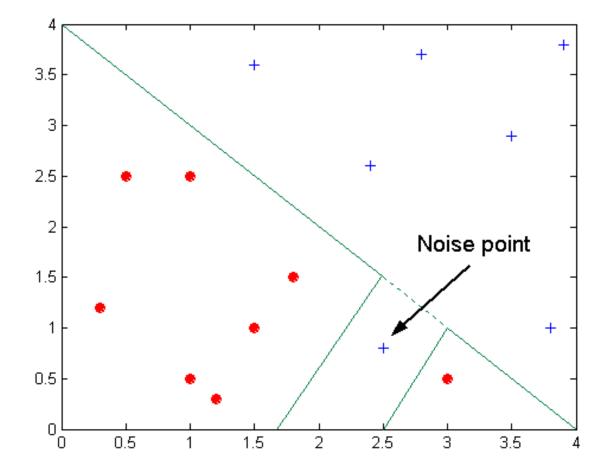
Underfitting and Overfitting



Underfitting: when model is too simple, both training and test errors are large



Overfitting due to Noise



Decision boundary is distorted by noise point



How to address overfitting

Pre-Pruning (Early Stopping Rule)

- Stop the algorithm before it becomes a fully-grown tree
- Typical stopping conditions for a node
 - Stop if all instances belong to the same class
 - Stop if all the attribute values are the same
- More restrictive conditions
 - Stop if number of instances is less than some user-specified threshold
 - Stop if class distribution of instances are independent of the available features (e.g., using χ^2 test)
 - Stop if expanding the current node does not improve impurity measures (e.g., Gini or information gain)



How to address overfitting

Post-pruning

- Grow decision tree to its entirety
- Trim the nodes of the decision tree in a bottomup fashion
- If generalization error improves after trimming, replace sub-tree by a leaf node.
- Class label of leaf node is determined from majority class of instances in the sub-tree





Data fragmentation

- Number of instances gets smaller as you traverse down the tree
- Number of instances at the leaf nodes could be too small to make any statistically significant decision



Handling missing attribute values

- Missing values affect decision tree construction in three different ways
 - Affect how impurity measures are computed
 - Affect how to distribute instance with missing value to child nodes
 - Affect how a test instance with missing value is classified





Other issues

- Data Fragmentation
- Search Strategy
- Expressiveness
- Tree Replication





Search strategy

Finding an optimal decision tree is NP-hard

- The algorithm presented so far uses a greedy, top-down, recursive partitioning strategy to induce a reasonable solution
- Other strategies?
 - Bottom-up
 - Bi-directional





Expressiveness

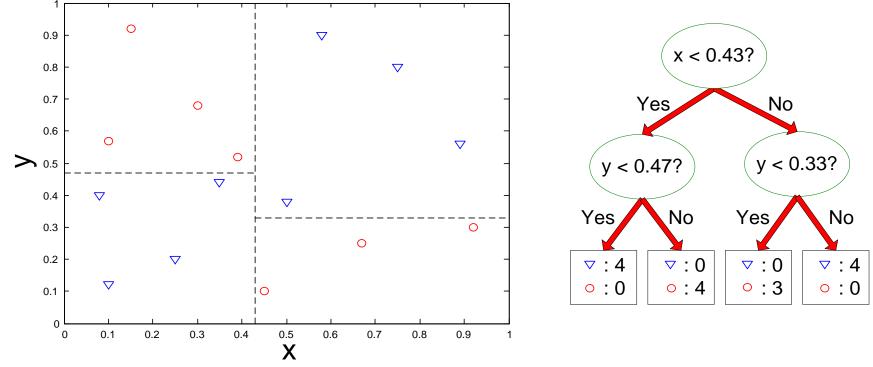
- Decision tree provides expressive representation for learning discrete-valued function
 - But they do not generalize well to certain types of Boolean functions
 - Example: parity function:
 - Class = 1 if there is an even number of Boolean attributes with truth value = True
 - Class = 0 if there is an odd number of Boolean attributes with truth value = True
 - For accurate modeling, must have a complete tree

Not expressive enough for modeling continuous variables

Particularly when test condition involves only a single attribute at-a-time



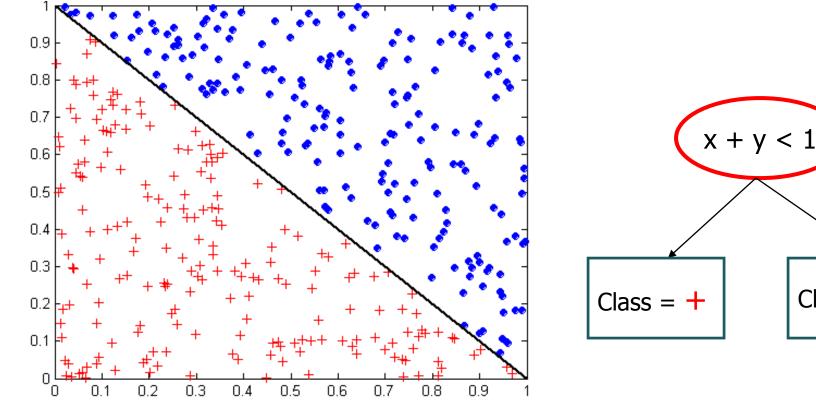




- Border line between two neighboring regions of different classes is known as decision boundary
- Decision boundary is parallel to axes because test condition involves a single attribute at-a-time



Oblique decision trees



- Test condition may involve multiple attributes
- More expressive representation
- Finding optimal test condition is computationally expensive

From: Tan, Steinbach, Kumar, Introduction to Data Mining, McGraw Hill 2006

Class =

Decision Tree Based Classification

Advantages

- Inexpensive to construct
- Extremely fast at classifying unknown records
- Easy to interpret for small-sized trees
- Accuracy is comparable to other classification techniques for many simple data sets

Disadvantages

accuracy may be affected by missing data





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Ensemble learning technique

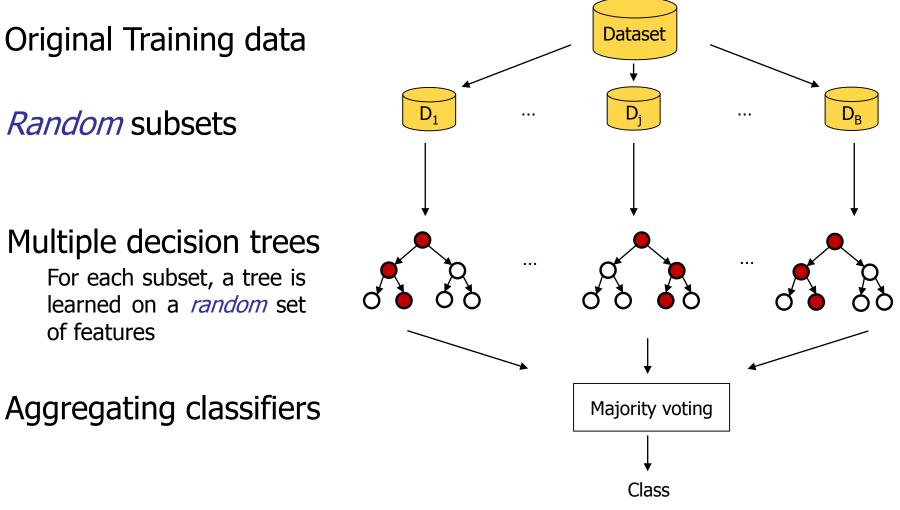
- multiple base models are combined
 - to improve accuracy and stability
 - to avoid overfitting

Random forest = set of decision trees

- a number of decision trees are built at training time
- the class is assigned by majority voting











Bootstrap aggregation

- Given a training set D of n instances, it selects B times a random sample with replacement from D and trains trees on these dataset samples
 - For b = 1, ..., B
 - Sample with replacement n' training examples, $n' \le n$
 - A dataset subset D_b is generated
 - Train a classification tree on D_b

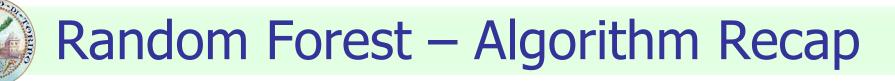




Feature Bagging

- Selects, for each candidate split in the learning process, a *random* subset of the features
 - being p the number of features, \sqrt{p} features are typically selected
- Trees are decorrelated
 - Feature subsets are sampled randomly, hence different features could be selected as best attribute for the split





- Given a training set *D* of *n* instances with p features
- For b = 1, ..., B
 - Sample randomly with replacement n' training examples. A subset D_b is generated
 - Train a classification tree on D_b
 - During the tree construction, for each candidate split
 - $m \ll p$ random features are selected (typically m $\approx \sqrt{p}$)
 - the best split is computed among these m features
- Class is assigned by majority voting among the B predictions





Strong points

- higher accuracy than decision trees
- fast training phase
- robust to noise and outliers
- provides global feature importance, i.e. an estimate of which features are important in the classification

Weak points

- results can be difficult to interpret
 - A prediction is given by hundreds of trees
 - but at least we have an indication through feature importance



Rule-based classification



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Model evaluation



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Model evaluation

- Methods for performance evaluation
 - Partitioning techniques for training and test sets
- Metrics for performance evaluation
 - Accuracy, other measures
- Techniques for model comparison
 - ROC curve

$$D_{M}^{B}G$$

Methods for performance evaluation

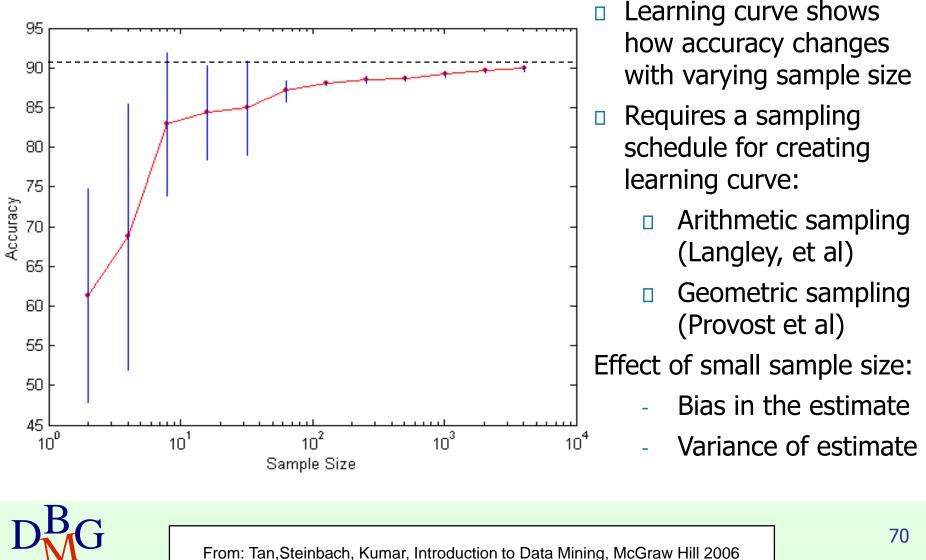
Objective

- reliable estimate of performance
- Performance of a model may depend on other factors besides the learning algorithm
 - Class distribution
 - Cost of misclassification
 - Size of training and test sets





Learning curve



Methods of estimation

- Partitioning labeled data in
 - training set for model building
 - test set for model evaluation
- Several partitioning techniques
 - holdout
 - cross validation
- Stratified sampling to generate partitions
 - without replacement
- Bootstrap
 - Sampling with replacement



Fixed partitioning

reserve 2/3 for training and 1/3 for testing

Appropriate for large datasets

- may be repeated several times
 - repeated holdout





Cross validation

Cross validation

- partition data into k disjoint subsets (i.e., folds)
- k-fold: train on k-1 partitions, test on the remaining one
 - repeat for all folds
- reliable accuracy estimation, not appropriate for very large datasets
- Leave-one-out
 - cross validation for k=n
 - only appropriate for very small datasets





Metrics for model evaluation

- Evaluate the predictive accuracy of a model
- Confusion matrix
 - binary classifier

	PRE			
		Class=Yes	Class=No	
ACTUAL	Class=Yes	а	b	a: TP (true positive) b: FN (false negative)
CLASS	Class=No	С	d	c: FP (false positive)
				d: TN (true negative)





Most widely-used metric for model evaluation

 $Accuracy = \frac{Number of correctly classified objects}{Number of classified objects}$

Not always a reliable metric





For a binary classifier

	PREDICTED CLASS				
		Class=Yes	Class=No		
ACTUAL	Class=Yes	a (TP)	b (FN)		
CLASS	Class=No	с (FP)	d (TN)		

Accuracy =
$$\frac{a+d}{a+b+c+d} = \frac{TP+TN}{TP+TN+FP+FN}$$





Limitations of accuracy

Consider a binary problem

- Cardinality of Class 0 = 9900
- Cardinality of Class 1 = 100

Model

() \rightarrow class 0

- Model predicts everything to be class 0
 accuracy is 9900/10000 = 99.0 %
- Accuracy is misleading because the model does not detect any class 1 object



Limitations of accuracy

- Classes may have different importance
 - Misclassification of objects of a given class is more important
 - e.g., ill patients erroneously assigned to the healthy patients class
- Accuracy is not appropriate for
 - unbalanced class label distribution
 - different class relevance





Evaluate separately for each class C

 $Recall(r) = \frac{Number of objects correctly assigned to C}{Number of objects belonging to C}$

 $Precision (p) = \frac{Number of objects correctly assigned to C}{Number of objects assigned to C}$

Maximize

F - measure (F) =
$$\frac{2rp}{r+p}$$





For a binary classification problem

on the confusion matrix, for the positive class

Precision (p) =
$$\frac{a}{a+c}$$

Recall (r) = $\frac{a}{a+b}$
F - measure (F) = $\frac{2rp}{r+p} = \frac{2a}{2a+b+c}$



ROC (Receiver Operating Characteristic)

- Developed in 1950s for signal detection theory to analyze noisy signals
 - characterizes the trade-off between positive hits and false alarms
- ROC curve plots
 - TPR, True Positive Rate (on the y-axis) TPR = TP/(TP+FN)

against

FPR, False Positive Rate (on the x-axis) FPR = FP/(FP + TN)

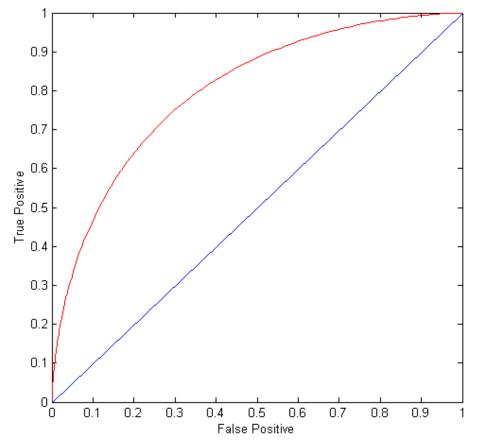




ROC curve

(FPR, TPR)

- (0,0): declare everything to be negative class
- (1,1): declare everything to be positive class
- (0,1): ideal
- Diagonal line
 - Random guessing
 - Below diagonal line
 - prediction is opposite of the true class





How to build a ROC curve

Inotonoo			
Instance	P(+ A)	True Class	
1	0.95	+	
2	0.93	+	
3	0.87	-	
4	0.85	-	
5	0.85	-	
6	0.85	+	
7	0.76	-	
8	0.53	+	
9	0.43	-	
10	0.25	+	

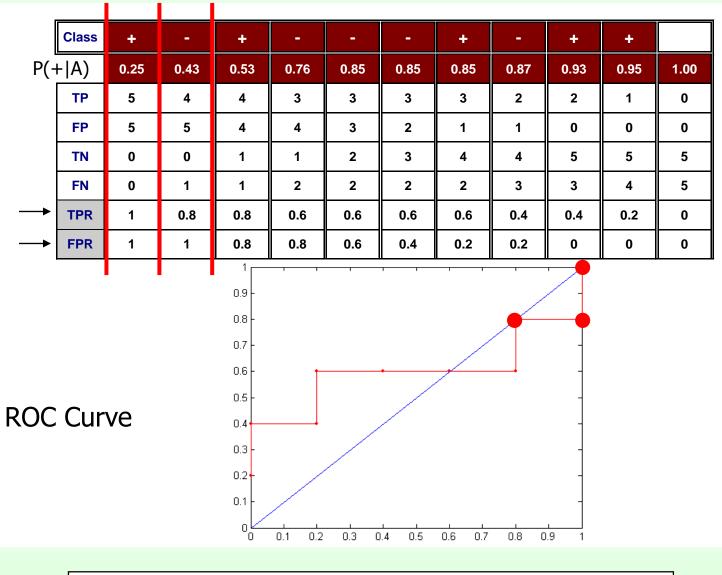
- Use classifier that produces posterior probability for each test instance P(+|A)
- Sort the instances according to P(+|A) in decreasing order
- Apply threshold at each unique value of P(+|A)
- Count the number of TP, FP, TN, FN at each threshold
 - TP rate
 - TPR = TP/(TP+FN)
 - FP rate

FPR = FP/(FP + TN)

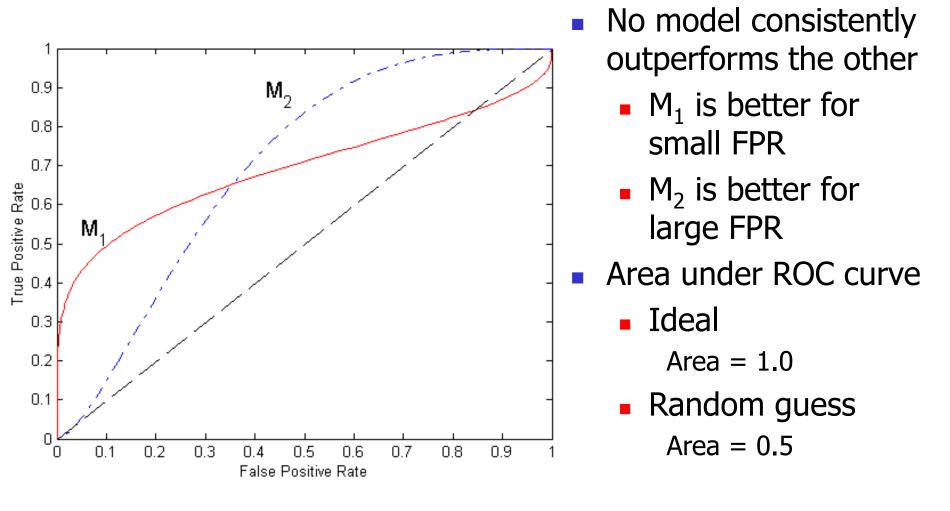




How to build a ROC curve



Using ROC for Model Comparison





Rule-based classification



Elena Baralis Politecnico di Torino

Rule-based classifier

- Classify records by using a collection of "if...then..." rules
- Rule: (*Condition*) $\rightarrow y$
 - where
 - *Condition* is a conjunction of attributes
 - y is the class label
 - LHS: rule antecedent or condition
 - *RHS*: rule consequent
- Examples of classification rules
 - (Blood Type=Warm) \land (Lay Eggs=Yes) \rightarrow Birds
 - (Taxable Income < 50K) \land (Refund=Yes) \rightarrow Cheat=No



Rule-based Classifier (Example)

Name	Blood Type	Give Birth	Can Fly	Live in Water	Class
human	warm	yes	no	no	mammals
python	cold	no	no	no	reptiles
salmon	cold	no	no	yes	fishes
whale	warm	yes	no	yes	mammals
frog	cold	no	no	sometimes	amphibians
komodo	cold	no	no	no	reptiles
bat	warm	yes	yes	no	mammals
pigeon	warm	no	yes	no	birds
cat	warm	yes	no	no	mammals
leopard shark	cold	yes	no	yes	fishes
turtle	cold	no	no	sometimes	reptiles
penguin	warm	no	no	sometimes	birds
porcupine	warm	yes	no	no	mammals
eel	cold	no	no	yes	fishes
salamander	cold	no	no	sometimes	amphibians
gila monster	cold	no	no	no	reptiles
platypus	warm	no	no	no	mammals
owl	warm	no	yes	no	birds
dolphin	warm	yes	no	yes	mammals
eagle	warm	no	yes	no	birds

R1: (Give Birth = no) \land (Can Fly = yes) \rightarrow Birds

- R2: (Give Birth = no) \land (Live in Water = yes) \rightarrow Fishes
- R3: (Give Birth = yes) \land (Blood Type = warm) \rightarrow Mammals
- R4: (Give Birth = no) \land (Can Fly = no) \rightarrow Reptiles
- R5: (Live in Water = sometimes) \rightarrow Amphibians



Rule-based classification

- A rule r covers an instance x if the attributes of the instance satisfy the condition of the rule
 - R1: (Give Birth = no) \land (Can Fly = yes) \rightarrow Birds
 - R2: (Give Birth = no) \land (Live in Water = yes) \rightarrow Fishes
 - R3: (Give Birth = yes) \land (Blood Type = warm) \rightarrow Mammals
 - R4: (Give Birth = no) \land (Can Fly = no) \rightarrow Reptiles

R5: (Live in Water = sometimes) \rightarrow Amphibians

Name	Blood Type	Give Birth	Can Fly	Live in Water	Class
hawk	warm	no	yes	no	?
grizzly bear	warm	yes	no	no	?

Rule R1 covers a hawk => Bird

Rule R3 covers the grizzly bear => Mammal





Rule-based classification

- R1: (Give Birth = no) \land (Can Fly = yes) \rightarrow Birds
- R2: (Give Birth = no) \land (Live in Water = yes) \rightarrow Fishes
- R3: (Give Birth = yes) \land (Blood Type = warm) \rightarrow Mammals
- R4: (Give Birth = no) \land (Can Fly = no) \rightarrow Reptiles
- R5: (Live in Water = sometimes) \rightarrow Amphibians

Name	Blood Type	Give Birth	Can Fly	Live in Water	Class
lemur	warm	yes	no	no	?
turtle	cold	no	no	sometimes	?
dogfish shark	cold	yes	no	yes	?

A lemur triggers (only) rule R3, so it is classified as a mammal

A turtle triggers both R4 and R5

A dogfish shark triggers none of the rules





Characteristics of rules

Mutually exclusive rules

- Two rule conditions can't be true at the same time
- Every record is covered by at most one rule

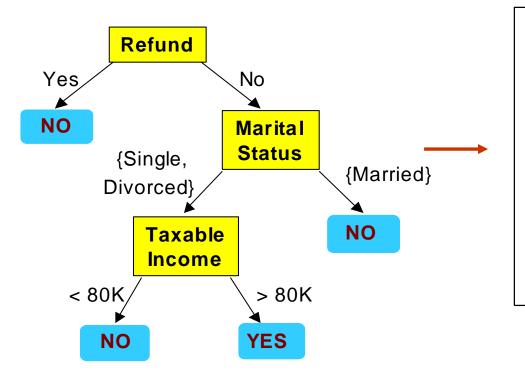
Exhaustive rules

- Classifier rules account for every possible combination of attribute values
- Each record is covered by at least one rule





From decision trees to rules



Classification Rules

(Refund=Yes) ==> No

(Refund=No, Marital Status={Single,Divorced}, Taxable Income<80K) ==> No

(Refund=No, Marital Status={Single,Divorced}, Taxable Income>80K) ==> Yes

(Refund=No, Marital Status={Married}) ==> No

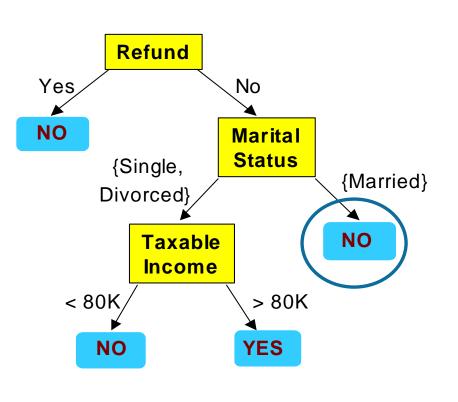
Rules are mutually exclusive and exhaustive

Rule set contains as much information as the tree





Rules can be simplified



Tid	Refund	Marital Status	Taxable Income	Cheat
1	Yes	Single	125K	No
2	No	Married	100K	No
3	No	Single	70K	No
4	Yes	Married	120K	No
5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
8	No	Single	85K	Yes
9	No	Married	75K	No
10	No	Single	90K	Yes

Initial Rule: (Refund=No) \land (Status=Married) \rightarrow No

Simplified Rule: (Status=Married) \rightarrow No

Effect of rule simplification

Rules are no longer mutually exclusive

- A record may trigger more than one rule
- Solution?
 - Ordered rule set
 - Unordered rule set use voting schemes
- Rules are no longer exhaustive
 - A record may not trigger any rules
 - Solution?
 - Use a default class

Ordered rule set

- Rules are rank ordered according to their priority
 - An ordered rule set is known as a decision list
- When a test record is presented to the classifier
 - It is assigned to the class label of the highest ranked rule it has triggered
 - If none of the rules fired, it is assigned to the default class

R1: (Give Birth = no)
$$\land$$
 (Can Fly = yes) \rightarrow Birds
R2: (Give Birth = no) \land (Live in Water = yes) \rightarrow Fishes
R3: (Give Birth = yes) \land (Blood Type = warm) \rightarrow Mammals
R4: (Give Birth = no) \land (Can Fly = no) \rightarrow Reptiles
R5: (Live in Water = sometimes) \rightarrow Amphibians

. * *					
Name	Blood Type	Give Birth	Can Fly	Live in Water	Class
turtle	cold	no	no	sometimes	?





Building classification rules

Direct Method

- Extract rules directly from data
- e.g.: RIPPER, CN2, Holte's 1R

Indirect Method

- Extract rules from other classification models (e.g. decision trees, neural networks, etc).
- e.g: C4.5rules





Advantages of rule-based classifiers

- As highly expressive as decision trees
- Easy to interpret
- Easy to generate
- Can classify new instances rapidly
- Performance comparable to decision trees



Associative classification



Elena Baralis Politecnico di Torino

Associative classification

The classification model is defined by means of association rules

(Condition) $\rightarrow y$

- rule body is an itemset
- Model generation
 - Rule selection & sorting
 - based on support, confidence and correlation thresholds
 - Rule pruning

Database coverage: the training set is covered by

selecting topmost rules according to previous sort



Associative classification

Strong points

- interpretable model
- higher accuracy than decision trees
 - correlation among attributes is considered
- efficient classification
- unaffected by missing data
- good scalability in the training set size

Weak points

- rule generation may be slow
 - it depends on support threshold
- reduced scalability in the number of attributes
 - rule generation may become unfeasible



Bayesian Classification



Elena Baralis Politecnico di Torino



Let C and X be random variables P(C,X) = P(C|X) P(X)P(C,X) = P(X|C) P(C)Hence P(C|X) P(X) = P(X|C) P(C)and also P(C|X) = P(X|C) P(C) / P(X)





Bayesian classification

- Let the class attribute and all data attributes be random variables
 - C = any class label
 - $X = \langle x_1, ..., x_k \rangle$ record to be classified
- Bayesian classification
 - compute P(C|X) for all classes
 - probability that record X belongs to C
 - assign X to the class with *maximal* P(C|X)
- Applying Bayes theorem

 $P(C|X) = P(X|C) \cdot P(C) / P(X)$

- P(X) constant for all C, disregarded for maximum computation
- P(C) a priori probability of C

$$P(C) = N_c/N$$



Bayesian classification

- How to estimate P(X|C), i.e. $P(x_1,...,x_k|C)$?
- Naïve hypothesis

 $P(x_1,...,x_k|C) = P(x_1|C) P(x_2|C) ... P(x_k|C)$

- statistical independence of attributes x₁,...,x_k
- not always true
 - model quality may be affected
- Computing $P(x_k|C)$
 - for discrete attributes

 $P(x_k|C) = |x_{kC}| / N_c$

- where $|\boldsymbol{x}_{kC}|$ is number of instances having value \boldsymbol{x}_k for attribute k and belonging to class C
- for continuous attributes, use probability distribution
- Bayesian networks
 - allow specifying a subset of dependencies among attributes



Bayesian classification: Example

Outlook	Temperature	Humidity	Windy	Class
sunny	hot	high	false	Ν
sunny	hot	high	true	Ν
overcast	hot	high	false	Ρ
rain	mild	high	false	Ρ
rain	cool	normal	false	Ρ
rain	cool	normal	true	Ν
overcast	cool	normal	true	Ρ
sunny	mild	high	false	Ν
sunny	cool	normal	false	Ρ
rain	mild	normal	false	Ρ
sunny	mild	normal	true	Ρ
overcast	mild	high	true	Ρ
overcast	hot	normal	false	Ρ
rain	mild	high	true	Ν



From: Han, Kamber,"Data mining; Concepts and Techniques", Morgan Kaufmann 2006



Bayesian classification: Example

outlook	
P(sunny p) = 2/9	P(sunny n) = 3/5
P(overcast p) = 4/9	P(overcast n) = 0
P(rain p) = 3/9	P(rain n) = 2/5
temperature	
P(hot p) = 2/9	P(hot n) = 2/5
P(mild p) = 4/9	P(mild n) = 2/5
P(cool p) = 3/9	P(cool n) = 1/5
humidity	
P(high p) = 3/9	P(high n) = 4/5
P(normal p) = 6/9	P(normal n) = 2/5
windy	
P(true p) = 3/9	P(true n) = 3/5
P(false p) = 6/9	P(false n) = 2/5

 $D_{M}^{B}G$

From: Han, Kamber,"Data mining; Concepts and Techniques", Morgan Kaufmann 2006



Data to be labeled

X = <rain, hot, high, false>

- For class p
 - $P(X|p) \cdot P(p) =$
 - = P(rain|p) P(hot|p) P(high|p) P(false|p) P(p)
 - $= 3/9 \cdot 2/9 \cdot 3/9 \cdot 6/9 \cdot 9/14 = 0.010582$
- For class n

 $P(X|n) \cdot P(n) =$

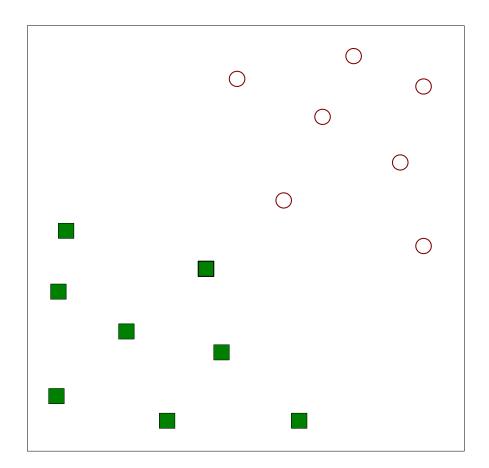
= P(rain|n) P(hot|n) P(high|n) P(false|n) P(n)= 2/5 2/5 4/5 2/5 5/14 = 0.018286



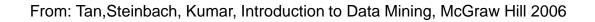
Support Vector Machines

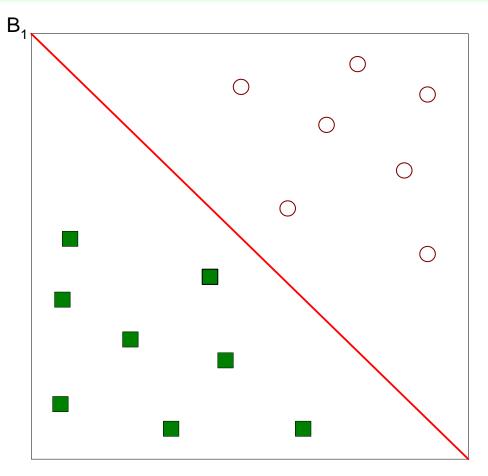


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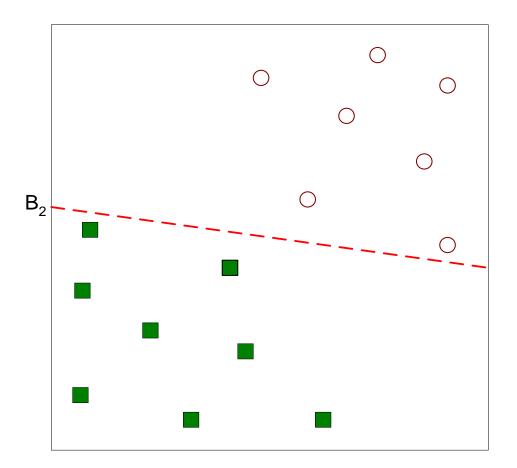
Find a linear hyperplane (decision boundary) that will separate the data





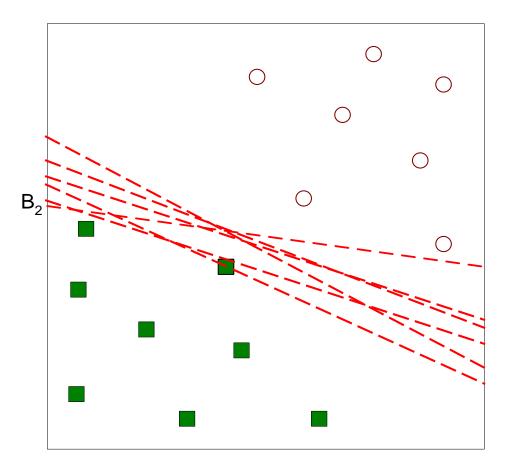
One Possible Solution





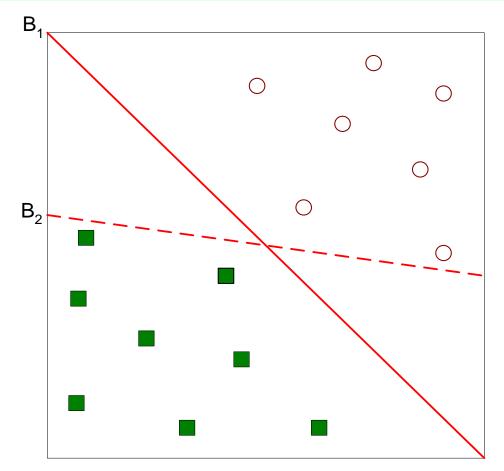
Another possible solution



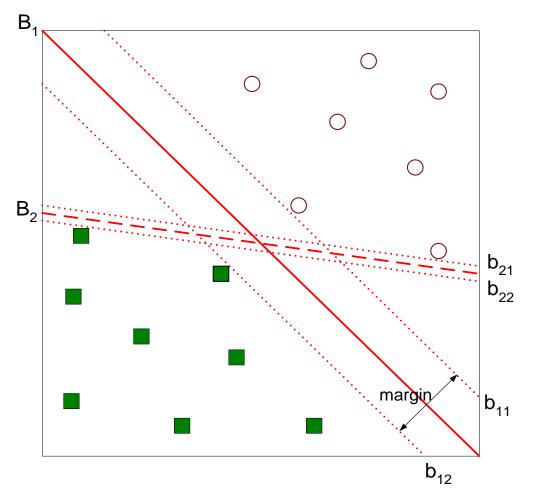


Other possible solutions





- Which one is better? B1 or B2?
- How do you define better?

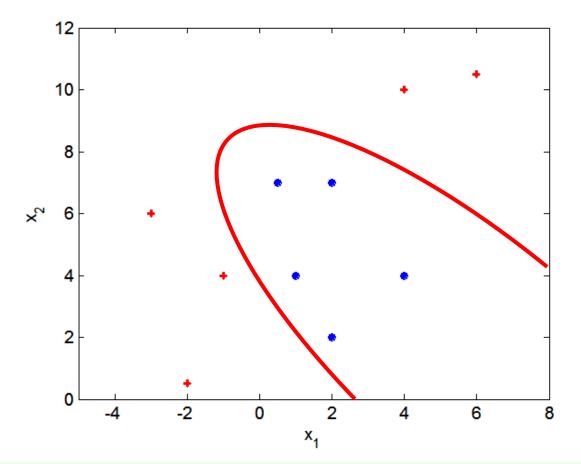


• Find hyperplane maximizes the margin => B1 is better than B2



Nonlinear Support Vector Machines

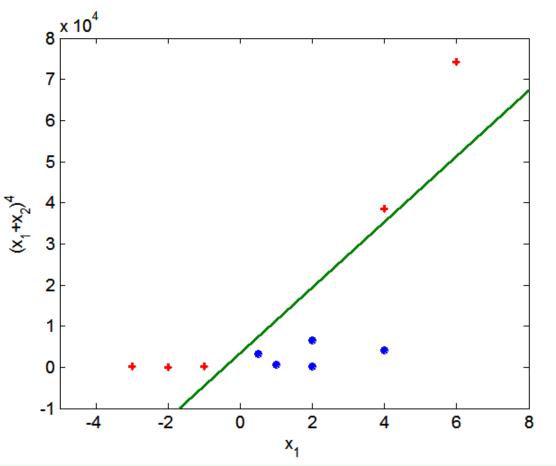
What if decision boundary is not linear?





Nonlinear Support Vector Machines

Transform data into higher dimensional space





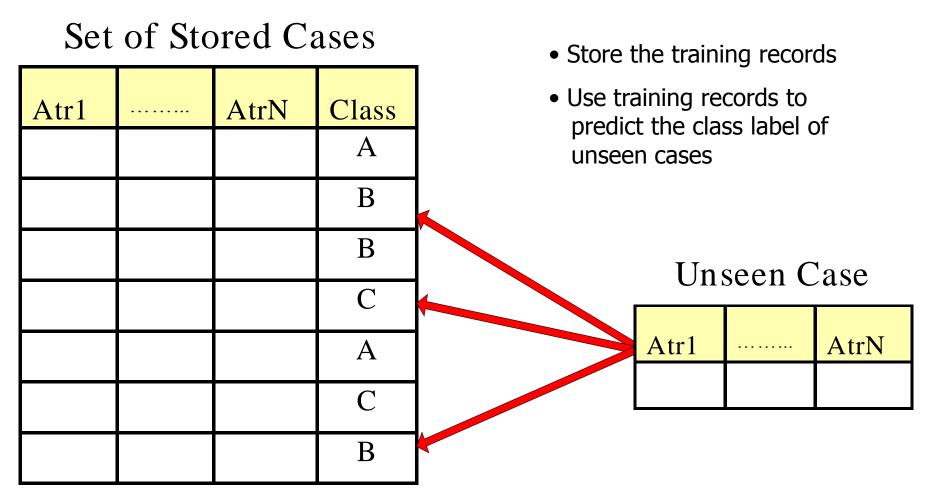
K-Nearest Neighbor



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Instance-Based Classifiers







Instance Based Classifiers

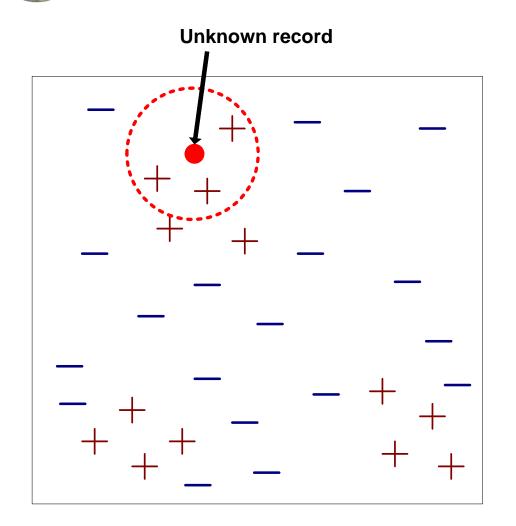
Examples

Rote-learner

- Memorizes entire training data and performs classification only if attributes of record match one of the training examples exactly
- Nearest neighbor
 - Uses k "closest" points (nearest neighbors) for performing classification



Nearest-Neighbor Classifiers

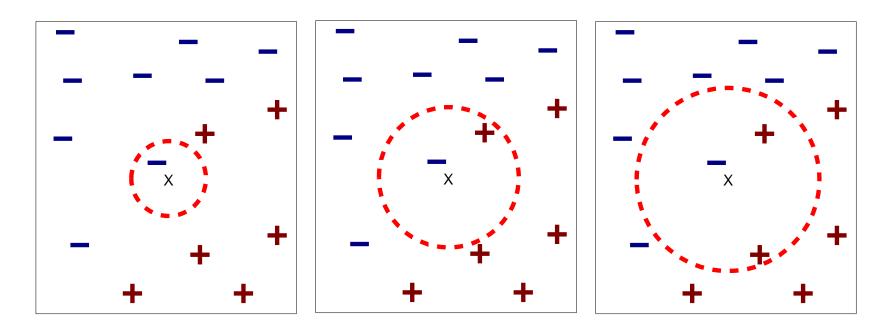


- Requires
 - The set of stored records
 - Distance Metric to compute distance between records
 - The value of *k*, the number of nearest neighbors to retrieve
- To classify an unknown record
 - Compute distance to other training records
 - Identify *k* nearest neighbors
 - Use class labels of nearest neighbors to determine the class label of unknown record (e.g., by taking majority vote)





Definition of Nearest Neighbor



(a) 1-nearest neighbor

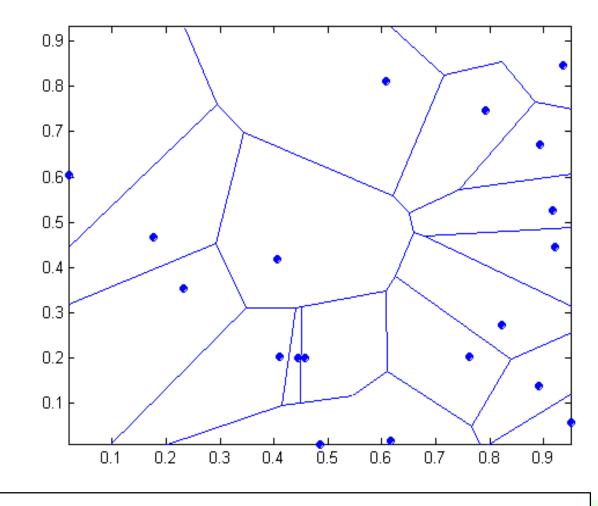
(b) 2-nearest neighbor

(c) 3-nearest neighbor

K-nearest neighbors of a record x are data points that have the k smallest distance to x



Voronoi Diagram





Nearest Neighbor Classification

Compute distance between two points

Euclidean distance

$$d(p,q) = \sqrt{\sum_{i} (p_i - q_i)^2}$$

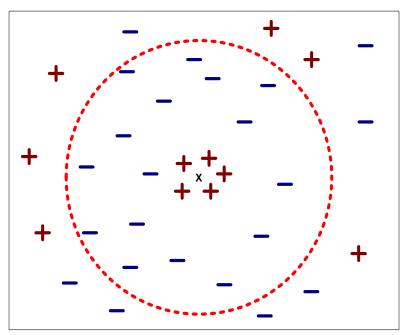
- Determine the class from nearest neighbor list
 - take the majority vote of class labels among the k-nearest neighbors
 - Weigh the vote according to distance
 - weight factor, $w = 1/d^2$



Nearest Neighbor Classification

Choosing the value of k:

- If k is too small, sensitive to noise points
- If k is too large, neighborhood may include points from other classes





Nearest Neighbor Classification

Scaling issues

- Attribute domain should be normalized to prevent distance measures from being dominated by one of the attributes
- Example: height [1.5m to 2.0m] vs. income [\$10K to \$1M]
- Problem with distance measures
 - High dimensional data
 - curse of dimensionality



Artificial Neural Networks



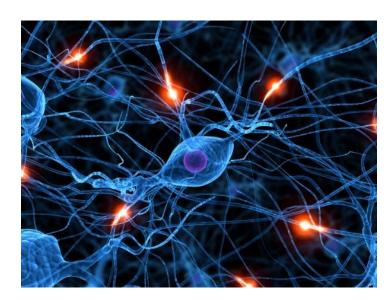
Data Base and Data Mining Group of Politecnico di Torino

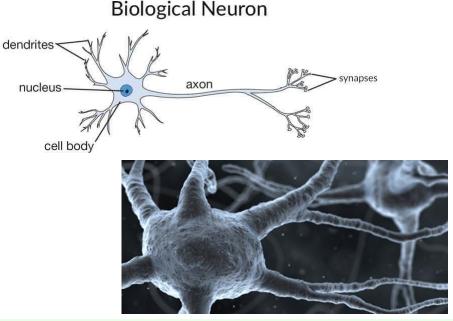
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Artificial Neural Networks

- Inspired to the structure of the human brain
 - Neurons as elaboration units
 - Synapses as connection network

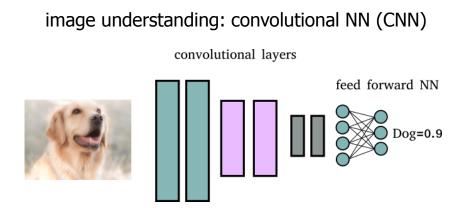




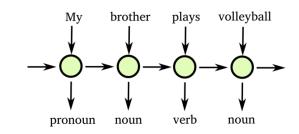


Artificial Neural Networks

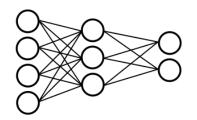
Different tasks, different architectures



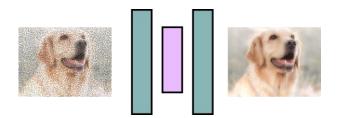
time series analysis: recurrent NN (RNN)



numerical vectors classification: feed forward NN (FFNN)

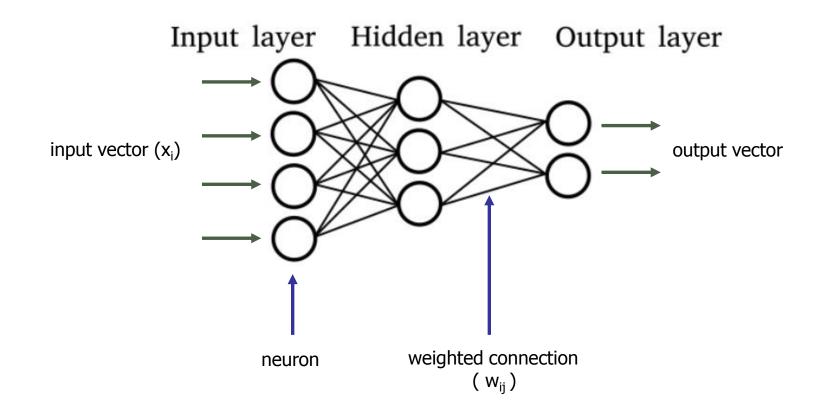


denoising: auto-encoders

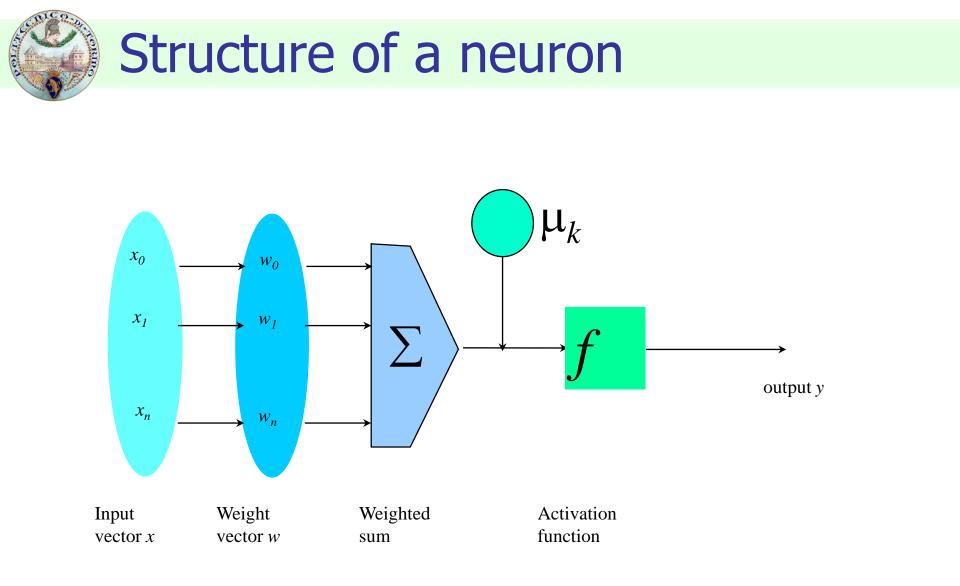












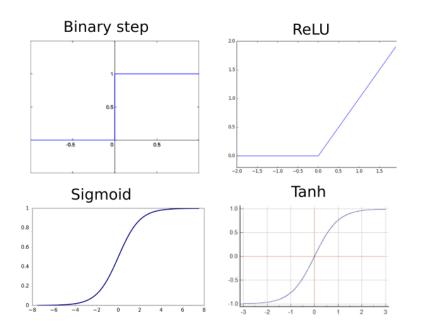


From: Han, Kamber,"Data mining; Concepts and Techniques", Morgan Kaufmann 2006



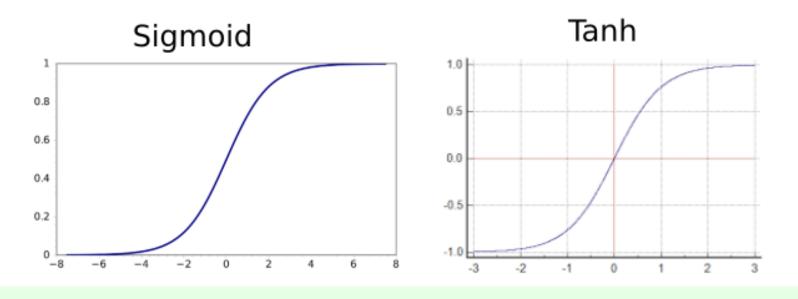
Activation

- simulates biological activation to input stymula
- provides non-linearity to the computation
- may help to saturate neuron outputs in fixed ranges



Sigmoid, tanh

- saturate input value in a fixed range
- non linear for all the input scale
- typically used by FFNNs for both hidden and output layers
 - E.g. *sigmoid* in output layers allows generating values between 0 and 1 (useful when output must be interpreted as likelihood)





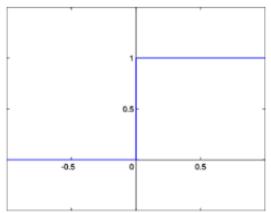
Binary step

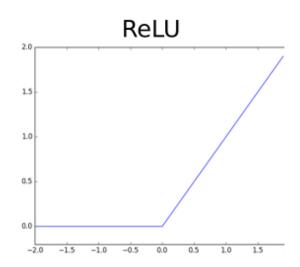
Binary Step

- outputs 1 when input is non-zero
- useful for binary outputs
- issues: not appropriate for gradient descent
 - derivative not defined in x=0
 - derivative equal to 0 in every other position

ReLU (Rectified Linear Unit)

- used in deep networks (e.g. CNNs)
 - avoids vanishing gradient
 - does not saturate
- neurons activate linearly only for positive input









Softmax

- differently to other activation functions
 - it is applied only to the **output** layer
 - works by considering all the neurons in the layer
- after softmax, the output vector can be interpreted as a discrete distribution of probabilities
 - e.g. the probabilities for the input pattern of belonging to each class

$$softmax(z_j) = \frac{e^{z_j}}{\sum_{i=0}^{N-1} e^{z_i}}$$

$$softmax = \frac{z_0 \\ z_1 \\ z_2 \\ z_2 \\ z_2 \\ z_3 \\ z_4 \\ z_2 \\ z_5 \\ z_5 \\ z_6 \\ z_1 \\ z_2 \\ z_6 \\ z_1 \\ z_2 \\ z_5 \\ z_6 \\ z_1 \\ z_2 \\ z_6 \\ z_6 \\ z_1 \\ z_6 \\ z_1 \\ z_2 \\ z_6 \\ z_1 \\ z_1 \\ z_2 \\ z_6 \\ z_1 \\ z_1 \\ z_2 \\ z_1 \\ z_2 \\ z_1 \\ z_2 \\ z_1 \\ z_1 \\ z_1 \\ z_2 \\ z_1 \\ z_1 \\ z_2 \\ z_1 \\ z_1 \\ z_1 \\ z_2 \\ z_1 \\ z$$





Building a FFNN

For each node, definition of

- set of weights
- offset value

providing the highest accuracy on the training data

 Iterative approach on training data instances





Building a FFNN

Base algorithm

Initially assign random values to weights and offsets

Process instances in the training set one at a time

- For each neuron, compute the result when applying weights, offset and activation function for the instance
- Forward propagation until the output is computed
- Compare the computed output with the expected output, and evaluate error
- Backpropagation of the error, by updating weights and offset for each neuron
- The process ends when
 - % of accuracy above a given threshold
 - % of parameter variation (error) below a given threshold
 - The maximum number of epochs is reached





Feed Forward Neural Networks

Strong points

- High accuracy
- Robust to noise and outliers
- Supports both discrete and continuous output
- Efficient during classification

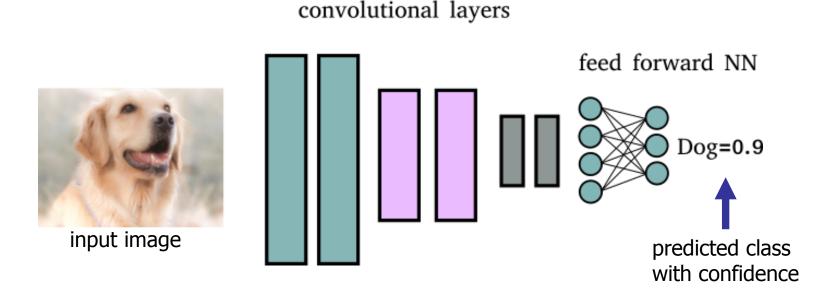
Weak points

- Long training time
 - weakly scalable in training data size
 - complex configuration
- Not interpretable model
 - application domain knowledge cannot be exploited in the model





Allow automatically extracting **features** from images and performing **classification**

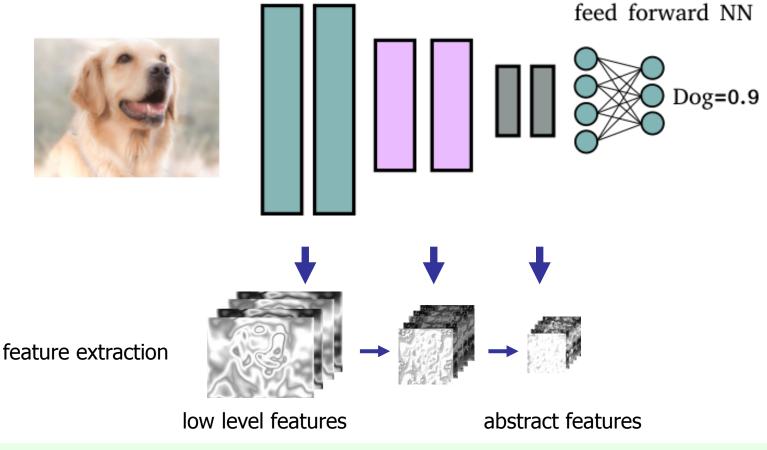


Convolutional Neural Network (CNN) Architecture



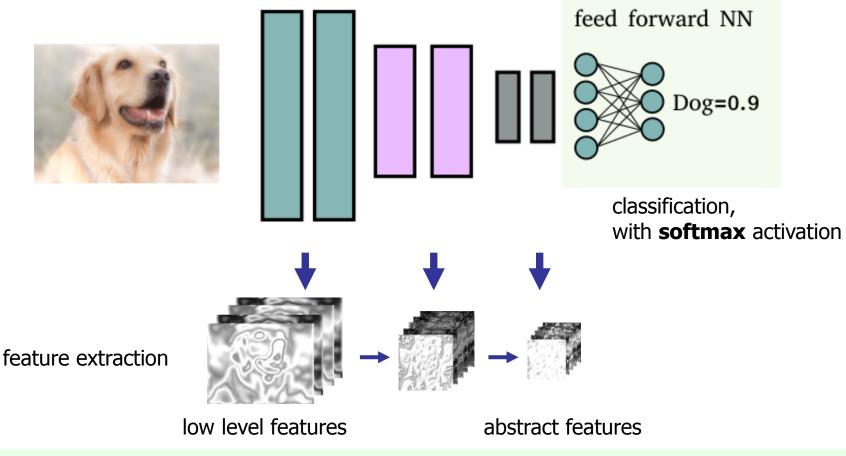


convolutional layers





convolutional layers

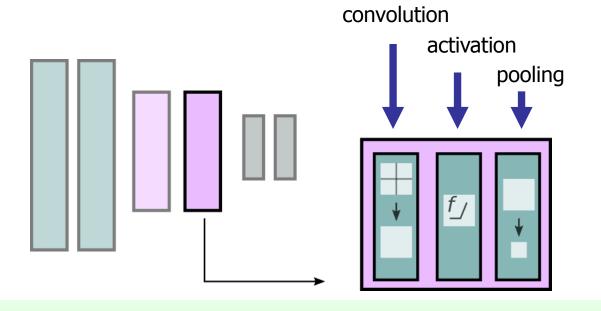






Typical convolutional layer

- *convolution* stage: feature extraction by means of (hundreds to thousands) sliding filters
- sliding filters *activation*: apply activation functions to input tensor
- *pooling*: tensor downsampling

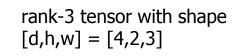


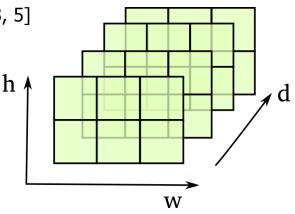




Tensors

- data flowing through CNN layers is represented in the form of *tensors*
- Tensor = N-dimensional vector
- *Rank* = number of dimensions
 - scalar: rank 0
 - 1-D vector: rank 1
 - 2-D matrix: rank 2
- Shape = number of elements for each dimension
 - e.g. a vector of length 5 has shape [5]
 - e.g. a matrix w x h, w=5, h=3 has shape [h, w] = [3, 5]



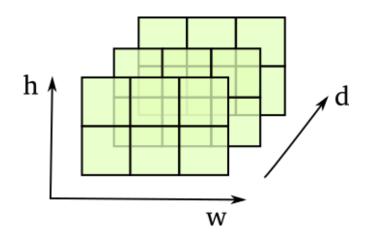






Images

- rank-3 tensors with shape [d,h,w]
- where h=height, w=width, d=image depth (1 for grayscale, 3 for RGB colors)

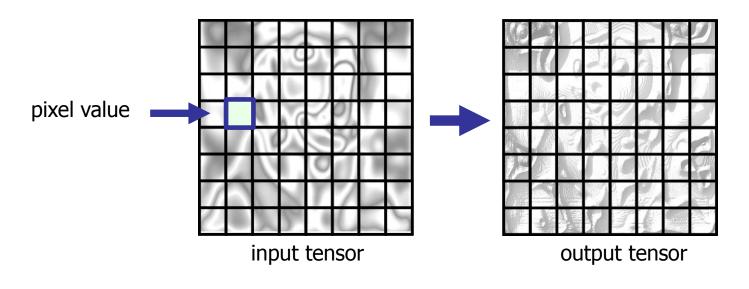






Convolution

- processes data in form of *tensors* (multi-dimensional matrices)
- input: input image or intermediate features (tensor)
- **output**: a tensor with the extracted features

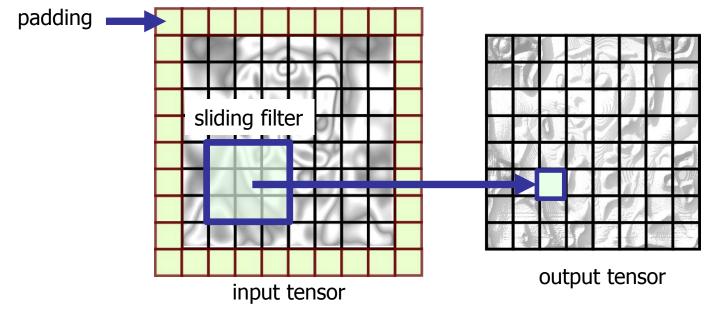






Convolution

- a *sliding filter* produces the values of the output tensor
- sliding filters contain the trainable weights of the neural network
- each convolutional layer contains many (hundreds) filters

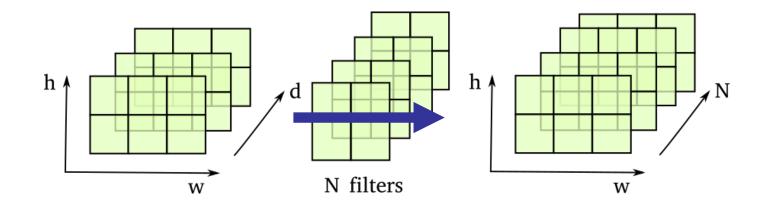






Convolution

- images are transformed into features by convolutional filters
- after convolving a tensor [d,h,w] with *N filters* we obtain
 - a rank-3 tensor with shape [N,h,w]
 - hence, each filter generates a layer in the depth of the output tensor

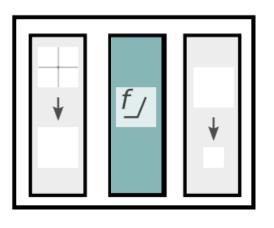


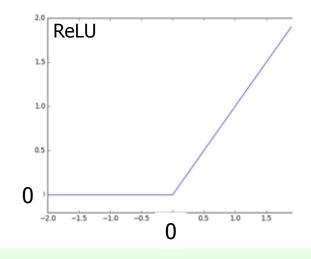




Activation

- symulates biological activation to input stymula
- provides non-linearity to the computation
- ReLU is typically used for CNNs
 - faster training (no vanishing gradients)
 - does not saturate
 - faster computation of derivatives for backpropagation



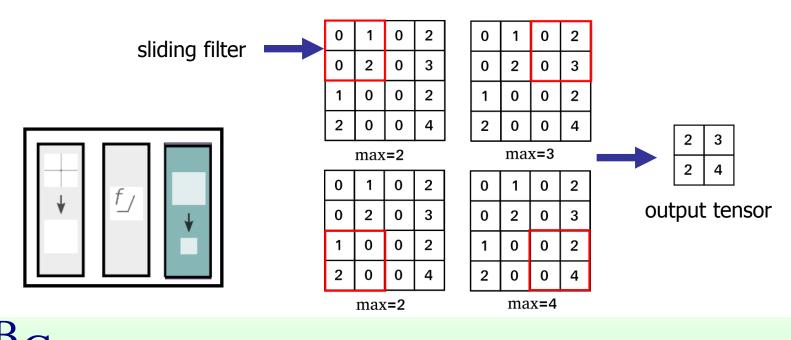






Pooling

- performs tensor *downsampling*
- *sliding filter* which replaces tensor values with a *summary* statistic of the nearby outputs
- *maxpool* is the most common: computes the maximum value as statistic



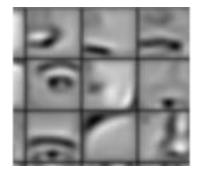


Convolutional layers training

- during training each sliding filter learns to recognize a particular pattern in the input tensor
- filters in *shallow layers* recognize textures and edges
- filters in *deeper layers* can recognize objects and parts (e.g. eye, ear or even faces)

shallow filters





deeper filters

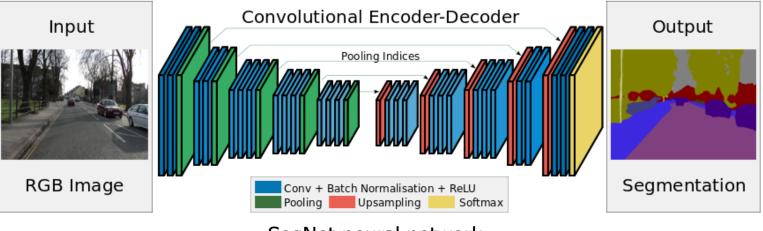






Semantic segmentation CNNs

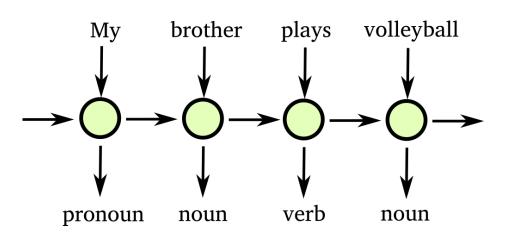
- allow assigning a class to each pixel of the input image
- composed of 2 parts
 - encoder network: convolutional layers to extract abstract features
 - decoder network: deconvolutional layers to obtain the output image from the extracted features



SegNet neural network



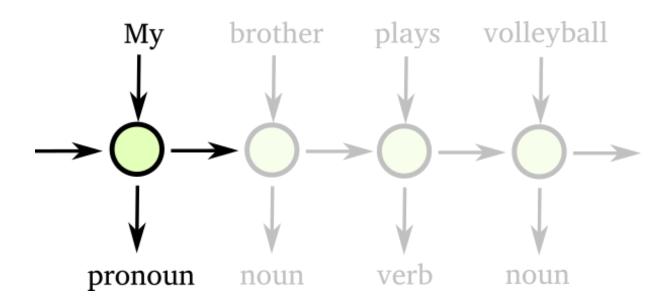
- Allow processing *sequential* data x(t)
- Differently from normal FFNN they are able to keep a *state* which evolves during time
- Applications
 - machine translation
 - time series prediction
 - speech recognition
 - part of speech (POS) tagging







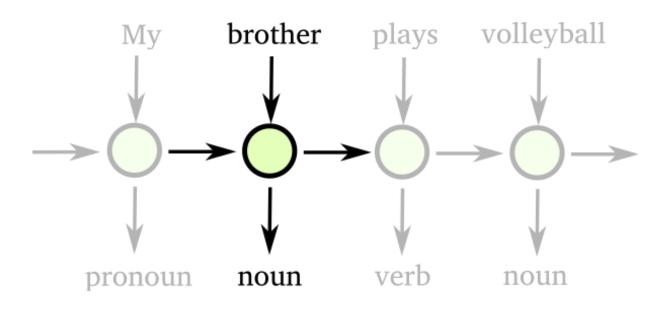
instance of the RNN at time t1



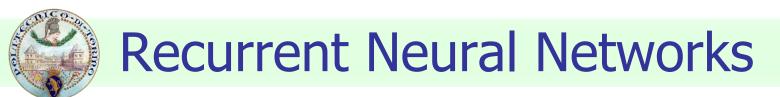


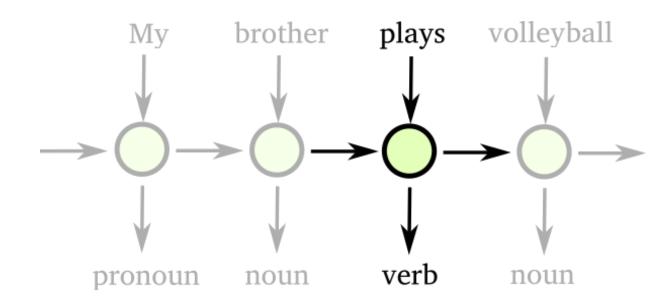








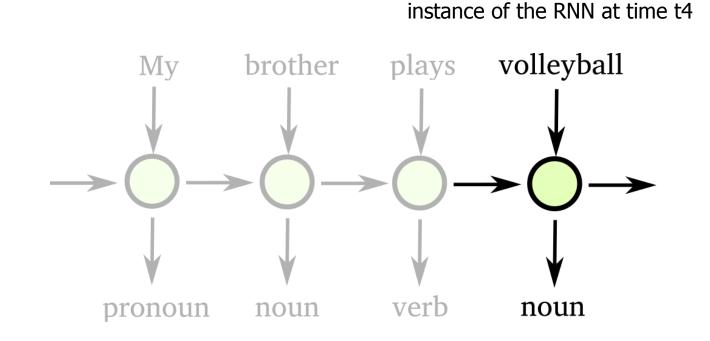




instance of the RNN at time t3

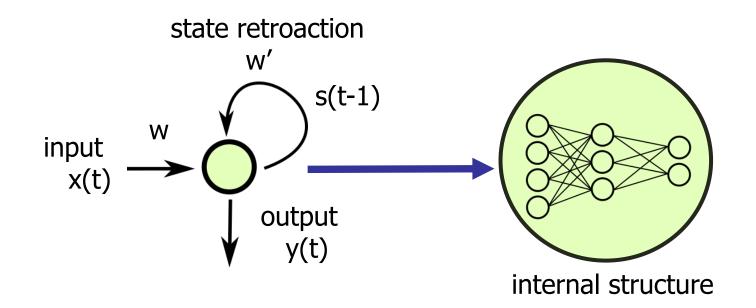






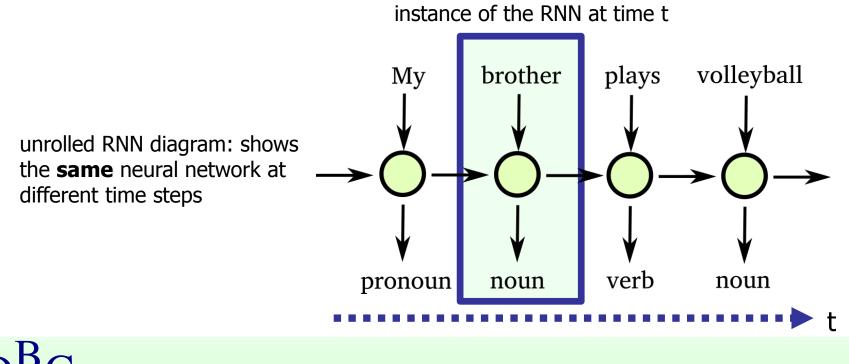


- A RNN receives as input a vector x(t) and the state at previous time step s(t-1)
- A RNN typically contains many *neurons organized in different layers*



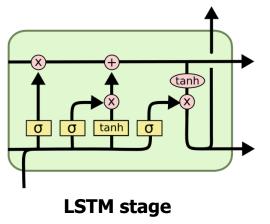


- Training is performed with *Backpropagation Through Time*
- Given a pair training sequence x(t) and expected output y(t)
 - error is propagated through time
 - weights are updated to minimize the error across all the time steps



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- Issues
 - vanishing gradient: error gradient decreases rapidly over time, weights are not properly updated
 - this makes harder having RNN with *long-term* memories
- Solution: *LSTM* (Long Short Term Memories)
 - RNN with "gates" which encourage the state information to flow through long time intervals

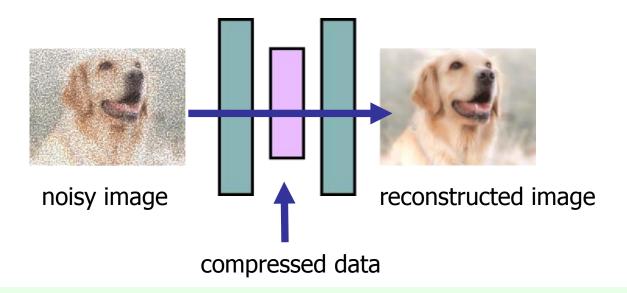






Autoencoders

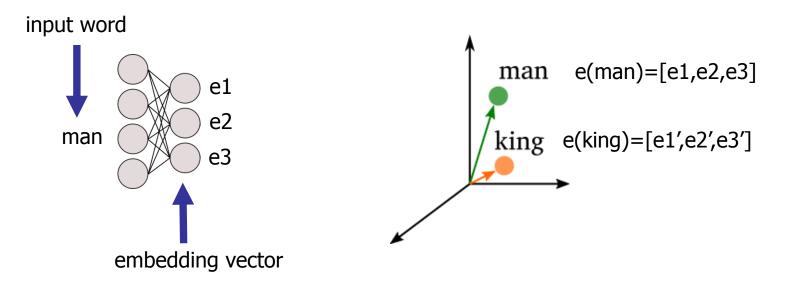
- Autoencoders allow *compressing* input data by means of compact representations and from them *reconstruct* the initial input
 - for feature extraction: the compressed representation can be used as significant set of features representing input data
 - for image (or signal) *denoising*: the image reconstructed from the abstract representation is denoised with respect to the original one





Word Embeddings (Word2Vec)

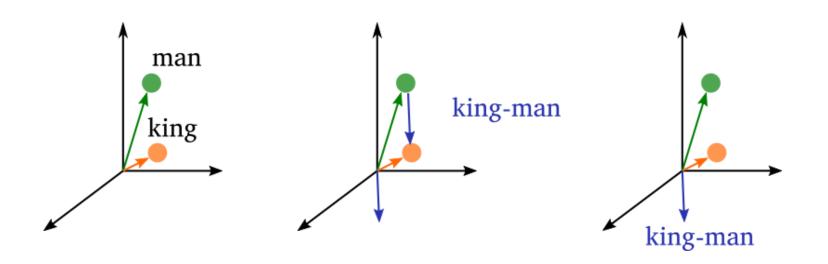
- Word *embeddings* associate words to n-dimensional vectors
 - trained on big text collections to model the word distributions in different sentences and contexts
 - able to capture the *semantic* information of each word
 - words with similar *meaning* share vectors with similar characteristics







 Since each word is represented with a vector, operations among words (e.g. difference, addition) are allowed







Semantic relationiships among words are captured by vector positions

