Lecture 1 Basic models: nearest neighbors and trees

Machine Learning
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Lecture plan

- Nearest neighbor classifier
- Generalized distance-based classifiers
- Non-parametric regression
- Decision trees
- Splitting criteria
- Regression trees

 The presentation is partly prepared with materials of the K.V. Vorontsov's course "Machine Leaning".

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Problem formulation

X is object set, Y is answer set, $y: X \to Y$ is unknown dependency, $|Y| \ll \infty$ $X^{\ell} = \{x_1, ..., x_n\}$ is training sample, $T^{\ell} = \{(x_1, y_1), ..., (x_{\ell}, y_{\ell})\}$ is set of examples.

Task: return an algorithm $a: X \rightarrow Y$.

What is this task?

Classification problem formulation

X is object set, Y is answer set, $y: X \to Y$ is unknown dependency, $|Y| \ll \infty$ $X^{\ell} = \{x_1, ..., x_n\}$ is training sample, $T^{\ell} = \{(x_1, y_1), ..., (x_{\ell}, y_{\ell})\}$ is set of examples.

Task: return an algorithm $a: X \rightarrow Y$.

What is this task? Classification, because $|Y| \ll \infty$.

Duck test

Duck test:

If it looks like a duck, swims like a duck, and quacks like a duck, then it probably is a duck.

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Looks	Swims	Quacks	A duck?
like a duck	like a duck	like a duck	Probably, a duck
totally not like a duck	can be a duck	not like a duck	Probably, not a duck

How is the classifier formalized?

What is the training sample?

Many ducks, many non-ducks (unducks).

What is classification procedure?

- 1. Ducks were described with **key features**.
- 2. Similarity concept was used.
- 3. Logical separator was used for classification.

Main idea

Key hypothesis: similar objects belong to same class.

Main idea: for an object we have to find a class, in which objects are the most similar to the given one.

- Reasoning by analogy (case-based)
- Lazy learning

Formalization of "similarity"

"Similarity" is a distance between objects. We will talk about **metrics**.

Distance: $\rho: X \times X \to [0; +\infty)$.

Metric space is a set with a metric $\rho(x, y)$, defined on it.

Commonly used metrics

Minkowski distance:

$$p(x,y) = \left(\sum_{i} |x_i - y_i|^p\right)^{\frac{1}{p}},$$

p = 2 is the Euclidian distance;

p = 1 is the Manhattan distance.

Mahalanobis distance:

$$p(x,y) = \sqrt{(x-y)^{\top} S^{-1}(x-y)},$$

where *S* is covariance matrix for *x* and *y*.

Entropic distance measure (in K^*)

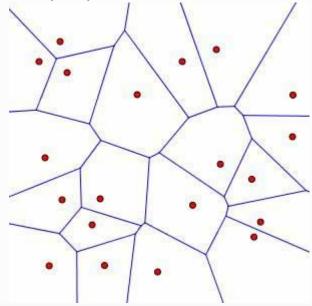
Nearest neighbor method (1NN)

 $x_{(u,1)}$ is **nearest** (1 = number one in sorted distances list) **neighbor of** classified object u:

$$x_{(u,1)} = \operatorname{argmin}_{x \in X^{\ell}} \rho(u, x).$$

Classifier: $a(u, T^{\ell}) = y(x_{(u,1)}) = y(u, 1)$.

Voronoi diagram:



1NN discussion

Advantages:

- simplicity;
- lucidity;
- interpretability.

Disadvantage:

- sensibility to noise;
- low efficacy;
- no parameters (explicitly);
- necessity to store all the examples.

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How can it be improved?

Very simple model

More complex model (more parameters)

How to choose "closeness"?

- Distance choosing
- Dimension reduction

Finding nearest object is too slow

- Usage of good structures for data storage
- Storage of only useful objects

kNN

Choose a distance ρ .

Sort objects:

$$\rho(u, x_{(u,1)}) \le \rho(u, x_{(u,2)}) \le \cdots \le \rho(u, x_{(u,\ell)}),$$

Algorithm *k*NN:

$$a(u; T^{\ell}) = \operatorname{argmax}_{y \in Y} \sum_{i=1}^{\ell} [y(u, i) = y][i \le k],$$

$$a(u; T^{\ell}) = \operatorname{argmax}_{y \in Y} \sum_{i=1}^{k} [y(u, i) = y].$$

Optimization of k

How to find a proper k?

k in this case is called **hyperparameter**.

We will work with this problem during the next lecture.

Generalized metric classifier

$$a(u; T^{\ell}) = \operatorname{argmax}_{y \in Y} \sum_{i=1}^{\ell} [y(u, i) = y] w(i, u),$$

where w(i, u) is a weight function representing importance of ith neighbor of u.

$$C_y(u) = \sum_{i=1}^{\ell} [y(u,i) = y]w(i,u)$$
 is estimation of object u closeness to class y .

$$a(u; T^{\ell}) = \operatorname{argmax}_{y \in Y} C_y(u)$$

What can be chosen as w?

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w(i, u):
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- linearly decreasing functions;
- exponentially decreasing functions;

Parzen-Rosenblatt window

With fixed window width:

$$a(u; T^{\ell}; h; K) =$$

$$\sum_{i=1}^{\ell} \left(\rho(u, x_{(u,i)}) \right)$$

$$= \operatorname{argmax}_{y \in Y} \sum_{i=1}^{\ell} [y(u, i) = y] K\left(\frac{\rho(u, x_{(u, i)})}{h}\right),$$

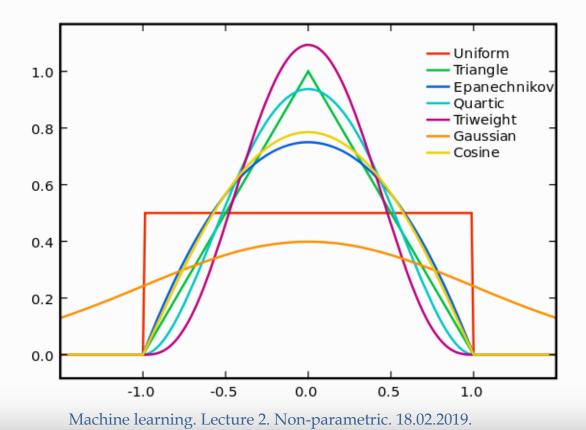
With variable window width:

$$a(u; T^{\ell}; \mathbf{k}; K) =$$

$$= \operatorname{argmax}_{y \in Y} \sum_{i=1}^{\ell} [y(u, i) = y] K\left(\frac{\rho(u, x_{(u,i)})}{\rho(u, x_{(u,k+1)})}\right).$$

Kernel function

Kernel function K(x) is symmetric non-negative function, $\int_{-\infty}^{+\infty} K(x) dx = 1$.



Distance selection (learning)

Distance can be learned (chosen).

Distance can be parametrized. Example (weighted Minkowski):

$$p(x,y) = \left(\sum_{i} \omega_{i} |x_{i} - y_{i}|^{p}\right)^{\frac{1}{p}}.$$

Now the problem is how to choose coefficients ω_i . When $\omega_i = 0$, the feature is thrown away (feature selection).

How to choose nearest neighbor?

Brute force (no data structure)

• py: bruteforce

weka: LinearNNSearch

K-D Tree

• py: kd_tree

weka: KDTree

Ball Tree

py: ball_tree

weka: BallTree

Cover Tree

weka: CoverTree

Nearest centroid classifier

Just create **centroids** for each class.

For a new object, choose the label of the nearest centroid.

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Main idea

Basic idea: lets think that $\theta(x) = \theta$ nearby $x \in X$:

$$Q(\theta, T^{\ell}) = \sum_{i=1}^{\ell} w_i(x)(\theta - y_i)^2 \to \min_{\theta \in \mathbb{R}}.$$

Main idea: use kernel smoothing:

$$w_i(x) = K\left(\frac{\rho(x_i, x)}{h}\right),\,$$

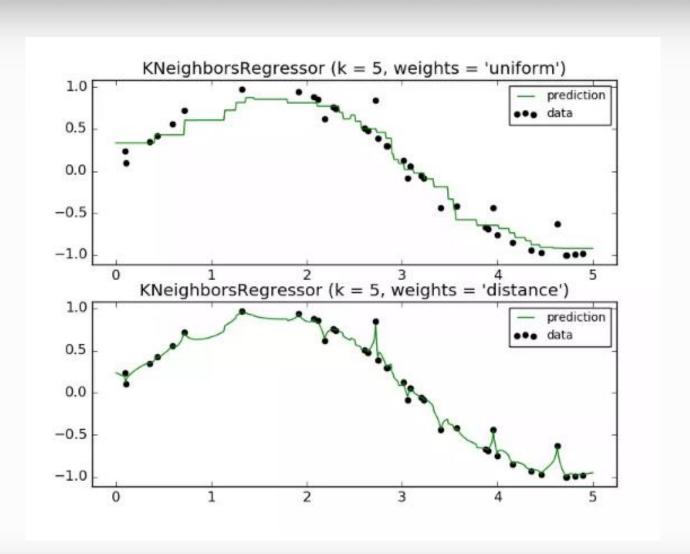
where *h* is window width.

Kernel smoothing

Nadaraya-Watson kernel smoothing:

$$a_h(x,T^{\ell}) = \frac{\sum_{i=1}^{\ell} y_i w_i(x)}{\sum_{i=1}^{\ell} w_i(x)} = \frac{\sum_{i=1}^{\ell} y_i K\left(\frac{\rho(x_i,x)}{h}\right)}{\sum_{i=1}^{\ell} K\left(\frac{\rho(x_i,x)}{h}\right)}.$$

Nearest neighbor regression



Method discussion

- kernel function has impact on smoothness;
- kernel function has small impact on approximation quality;
- *h* impacts on approximation quality;
- *k* can be tuned;
- sensitive to noise.

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Decision trees

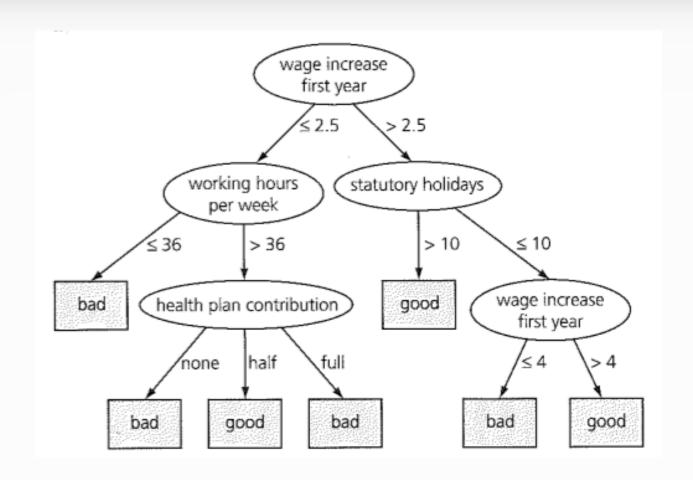
Decision tree is a classifier and even regression algorithm.

Nodes contain splitting rules (questions).

Each edge is a possible answer to its node question.

Leafs contain decisions (a class for classification problem and a number for regression problem).

Decision tree example



General scheme

With splitting rules space \mathcal{B} and split quality functional Φ .

- 1. Sent *S* to the root.
- 2. On each step process sample *S*.
- 2.1. If *S* contains objects from a single class *c*, create leaf of class *c* and stop.
- 2.2. Else choose splitting rule $\mathcal{E} \in \mathcal{B}$ which is the most informative with respect to Φ and split sample to S_1, \dots, S_k .
- 2.3. If stop-criterion, is true, then return the most popular class in S, otherwise create n children with samples S_i .
- 3. Prune the tree.

Stop-criteria

The most popular stop-criteria:

- one of classes is empty after splitting
- $\Phi(S)$ is lower than a certain threshold
- |*S*| is lower than a certain threshold
- tree height is higher than a certain threshold

Pruning

Premises: just first node impact on performance; decision trees tend to be overfitted.

Main idea: to cut lower branches.

Pruning is processing of created trees, when branches are delated consequently with a certain criterion (reduction number of errors, for example).

Pruning algorithm scheme

Split sample to train and test in proportion 2:1.

For each tree node apply operation, which is the best in terms of number of errors:

- 1) Don not change anything;
- 2) Replace node with its child (for each child);
- 3) Replace node with a leaf (for each class).

Trees discussion

Advantages:

- easily understandable and interpretable;
- learning is quick;
- can work with different data types.

Disadvantage:

- sensitive to noise;
- easily get overfitted.

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Selecting splitting rules family

Can be any family of classifiers.

• In most of cases it's single feature based rules like:

$$f_i(x) > m_i;$$

$$f_i(x) = d_i.$$

• Sometimes it can be a combination.

Selection of feature-based rules

There is $\ell - 1$ options to split sample.

- Check each and pick the most informative.
- Join diapasons of values .
- Skip small diapasons.

This is how you can synthesize a rule for each feature.

Sample splitting

- If sample is being split each time into 2 (k = 2), then \mathcal{B} is family of binary rules, tree is binary.
- If a feature is categorical, several edges can be added.
- If a feature is real, discretization / banalization is applied.

On each step number of edges can differ, but usually k is fixed.

Selecting split quality criterion

Split quality Φ sometimes can be represented as:

$$\Phi(S) = \Phi(S) - \sum_{i=1}^k \frac{|S_i|}{|S|} \Phi(S_i).$$

The most popular:

- $\phi_h(S)$ is an entropy, $\Phi_h(S)$ is IGain;
- $\phi_g(S) = 1 \sum_{i=1}^m p_i^2$, where p_i is probability (frequency) of ith class in sample S is **Gini index**. $\Phi_h(S)$ is GiniGain.

Many other are used

$$GainRatio = IGain(S) / Enthropy(S)$$
.

Split quality criterion usually does not matter.

Examples

ID3 (Quinlan, 1986):

IGain; only $\Phi(S) < 0$; no pruning.

C4.5 (Quinlan, 1993):

GainRatio; pruning.

CART (Breinman, 1984):

binary; GiniGain; pruning. Can solve regression (values in leafs).

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Regression trees

