



Machine Learning: a methodology survey with practical examples

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- What is learning?
- Parametric methods
- Non-parametric methods
- Stochastic methods
- Non-metric methods (skip)
- Universal principles
- Unsupervised learning
- Examples





- the process of acquiring knowledge from experience
- focus on observations that can be described in terms of measurable quantities
 - ullet an observation corresponds to a point $\mathbf{x} \in \mathbb{R}^d$
- given a set of observations $\mathcal{D} = \{\mathbf{x}_i\}$ say something about its structure or about a new observation \mathbf{x}



Supervised learning



Classification





Supervised learning



Classification





Supervised learning



Classification

Regression





Unsupervised learning



Clustering



X X



Unsupervised learning



Clustering







Unsupervised learning



Clustering

Classification







The theory behind



parametric methods

- probabilistic assumption on the generation of the data $\mathcal{D} = \{\mathbf{x}_i\}$
- known functional shape of probability distributions, but unknown parameters

non parametric

- the shape of the distribution is not known
- no probabilistic assumption at all (heuristics)





- Nature assumes one of c states ω_j with a priori probability $P(\omega_j)$
- \blacksquare When in state $\omega_j,$ nature emits observations \mathbf{x} with distribution $p(\mathbf{x}|\omega_j)$





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Bayes decision theory





 $P(\omega_j | \mathbf{x}) = \frac{p(\mathbf{x} | \omega_j) P(\omega_j)}{n(\mathbf{x})}$



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posterior probabilities









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Parameter estimation





• ideally: $p(\mathbf{x}|\omega_j)$ i.e. $p(\mathbf{x}|\theta_j)$ in reality: $p(\mathbf{x}|\hat{\theta}_j)$ or $p(\mathbf{x}|\mathcal{D})$



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Assumptions:

- \bullet samples from class ω_i do not influence estimate for class $\omega_j, \ i \neq j$
- samples from the same class are independent and identically distributed (i.i.d.)



Parameter estimation (cont.)



class independence assumption:





Parameter estimation (cont.)



class independence assumption:



- Maximum likelihood estimation
- Bayesian estimation





- Find parameter vector $\hat{\theta}$ that maximises $p(\mathcal{D}|\theta)$ with $\mathcal{D} = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}$
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• Consider θ as a random variable

- characterise θ with the posterior distribution $p(\theta | D)$ given the data
- using Bayes formula, the posterior can be computed from the likelihood $p(\mathcal{D}|\theta)$ and the prior $p(\theta)$

$$p(\theta|\mathcal{D}) = \frac{p(\mathcal{D}|\theta)p(\theta)}{\int p(\mathcal{D}|\theta)p(\theta)d\theta}$$

■ ML: $\mathcal{D} \rightarrow \hat{\theta}$ Bayes: $\mathcal{D}, p(\theta) \rightarrow p(\theta|\mathcal{D})$



 $p(\mathbf{x}|\hat{\theta})$



- we can compute $p(\mathbf{x}|\mathcal{D})$ instead of $p(\mathbf{x}|\hat{\theta})$
 - integrate the join density $p(\mathbf{x}, \theta | \mathcal{D}) = p(\mathbf{x} | \theta) p(\theta | \mathcal{D})$





Bayesian estimation



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Bayesian estimation (cont.)



Pros:

- better use of the data
- makes a priori assumptions explicit
- easily implemented recursively
 - use posterior $p(\theta|\mathcal{D})$ as new prior



Bayesian estimation (cont.)



Pros:

- better use of the data
- makes a priori assumptions explicit
- easily implemented recursively
 - use posterior $p(\theta|\mathcal{D})$ as new prior

Cons:

- definition of noninformative priors can be tricky
- often requires numerical integration
- not widely accepted by traditional statistics (frequentism)







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Parametric









Parametric









Parametric









Parametric









Parametric









Parametric









Parametric

non parametric





Parzen window

 \bullet define cell volume as a function of total number of samples n





Parametric

non parametric





Parzen window

- \bullet define cell volume as a function of total number of samples n
- k_n -nearest neighbour
 - \bullet define number of samples in a cell as a function of \boldsymbol{n}





use a linear combination of the components of x to rank a class

$$g_i(\mathbf{x}) = \mathbf{w}_i^t \mathbf{x} + w_{i0}$$

- compare the g_i s to choose the best class
- **\blacksquare** for two categories $g_1(\mathbf{x}) = g_2(\mathbf{x})$ defines a hyperplane





Nonlinear extension







Nonlinear extension





 \blacksquare non-linearly map the features in a higher dimensional space $x \to y$

$$g(\mathbf{x}) = \mathbf{a}^t \mathbf{y}$$





Gradient descent procedures

- \bullet define a criterion $J(\mathbf{a})$ that is maximised if \mathbf{a} is a solution
- \bullet update the current ${\bf a}$ with a fraction of the gradient of J

$$\mathbf{a} \leftarrow \mathbf{a} - \eta \, \, \mathbf{\Delta} J(\mathbf{a})$$



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• Perceptron criterion: $J_p(\mathbf{a}) = \sum_{y \in \mathcal{Y}} (-\mathbf{a}^t \mathbf{y})$ where \mathcal{Y} is the set of misclassified samples



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Support vector machines



Perceptron





Support vector machines



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Perceptron

Support vector machine





Multi layer neural networks







Multi layer neural networks





Multi layer neural networks (cont.)







Multi layer neural networks (cont.)







Multi layer neural networks (cont.)





Backpropagation algorithm







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Stochastic methods



Gradient descent procedure find local minima





- Gradient descent procedure find local minima
- solution: repeat training several times with different initialisations





- Gradient descent procedure find local minima
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- Simulated annealing
 - based on concepts from physics
 - well grounded theoretically
- Boltzmann learning





- Gradient descent procedure find local minima
- solution: repeat training several times with different initialisations
- Simulated annealing
 - based on concepts from physics
 - well grounded theoretically
- Boltzmann learning
- Evolutionary methods (Genetic algorithms)
 - based on concepts from biology
 - no theory behind: heuristic



Genetic algorithms



Generations

Ge	Generation k+1	
chromosomes	after ranking	survival + reproduction
01110110100100101001		10010100101010101010
1001010010101010101010		01010111110100101100
01001010001010010101		10101001000101010010
10101001000101010010		0101001010100001010
01001001010101001000	00101011110100011111	10010100100100101100
00010111101010101010	🗶 🎊 01001010001010010101	01010111110101010010
01010111110100101100	00010111101010101010	10101001001010001010
00101011110100011111	/ \ 01110110100100101001	01010010100101010010
01010010101010001010	∕	\0101001010101010101010



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01001001010101001000	00101011110100011111	10010100100100101100
00010111101010101010	🛛 🔨 01001010001010010101	01010111110101010010
01010111110100101100	00010111101010101010	10101001001010001010
00101011110100011111	∥ 🔨 01110110100100101001	01010010100101010010
01010010101010001010	/ \ 010010010101001000	\01010010101010101010

Genetic operators

¥	replication (survival)	crossover	mutation
gen	01110110100100101001	1001010010 01010111110100101100	10010100101010101010
			¥¥¥
+	01110110100100101001	1001010010 0100101100	100 <mark>0</mark> 01001 1 1010 0 01 1 10
gen		0101011111 <mark>1010101010</mark>	







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Universal principles



No free lunch theorem

• if we make no prior assumptions on the nature of the problem, no learning method can be proved to be superior to any other, not even random guessing



Universal principles



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Ugly duckling theorem

• if we make no prior assumptions on the nature of the problem, no feature representation should be preferred to any other



Universal principles



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Minimum description length principle

• prefer low complexity solutions. True only asymptotically, but valid in practice

Occam's razor

• avoid overfitting







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Unsupervised learning





- release assumption on class independence
- Iearn a mixture of distributions
- the parametric solution is formally similar, but different in practice





- A maximum likelihood solution is the Expectation Maximisation algorithm
- Problem with missing data (class membership $\forall \mathbf{x}_k \in \mathcal{D}$)
- Solution:
 - assume the missing data is known
 - compute and maximise likelihood
 - estimate the new best guess for the missing data
 - iterate
- guaranteed to find ML solution with marginalised missing data



Heuristic methods



k-means clustering

- use Euclidean distance as similarity measure
- \bullet define k centroids
- assign data points to the nearest centroid
- recompute centroids
- iterate

Properties

• is equivalent to Model Based Clustering with equal and spherical covariances



Heuristic methods (cont.)



hierarchical clustering

- start with one cluster per data point
- iteratively merge most similar clusters
- single linkage, complete linkage, average linkage, ...





Number of clusters



- what if the number of clusters is not known?
- Large number of heuristic methods
 - measure the within and across cluster spread



Number of clusters



- what if the number of clusters is not known?
- Large number of heuristic methods
 - measure the within and across cluster spread
- Bayes Information Criterion
 - model fit to the data: likelihood
 - model complexity in number of parameters (minimum description length principle)
 - number of data points available for parameter estimation







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- Synface: map acoustic to visual information in speech
- Accent analysis with hierarchical agglomerative clustering
- Mille, model first language learning with Model Based Clustering









- idea: use a synthesized talking face derived from speech as a hearing aid for users of voice channels
- **problem:** extract (phonetic) information from the speech signal with very low latencies ($\sim 50ms$)
- **•** it is a regression problem
- Instruction by the second s
 - map acoustic signal to visemes
 - use rules to generate the lip movements



Synface: methods



Recurrent neural network



Hidden Markov models







- aim: analysis of regional pronunciation variation on large data sets (\sim 5000 speakers)
- how? Automate part of the process with data mining techniques





- Analyse differences between groups by comparing distributions
 - metric based on Bhattacharyya distance

$$D_{\text{bhatt}}(\Theta_1, \Theta_2) = \underbrace{\frac{1}{8} (M_2 - M_1)^T \left[\frac{\Sigma_1 + \Sigma_2}{2} \right]^{-1} (M_2 - M_1)}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 \right| \left| \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 + \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\left| \frac{\Sigma_1 + \Sigma_2}{2} \right|}{\sqrt{\left| \Sigma_1 + \Sigma_2 \right|}}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\Sigma_1 + \Sigma_2}{2}}_{\mathbf{I} = \underbrace{\frac{\Sigma_1 + \Sigma_2}{2}}_{\mathbf{I} = \underbrace{\frac{1}{2} \ln \frac{\Sigma_1 + \Sigma_2}{2}}_{\mathbf{I} = \underbrace{\frac{\Sigma_1 + \Sigma_2}{2}}_{\mathbf{I} = \underbrace{\frac{\Sigma_2 + \Sigma_2}{2}}_{\mathbf{I} = \underbrace{\frac{\Sigma$$



Accent clustering (cont.)













- Background: infants have no innate linguistic knowledge
- Aim (long term): mathematical modelling of the learning process
 - acoustic features classification
 - time integration into meaningful sequences
- Aim (so far): spectral features classification
 - unsupervised
 - incremental




start with a MCLUST model get new data adjust old model to new data divide new data into well and poorly[∞]

modelled points

- 5. try a more complex model, if better BIC set as best and go back to 4
- 6. set the current best model and go back to 2







- **1.** start with a MCLUST model
- 2. get new data
- 3. adjust old model to new data
- 4. divide new data into well and poorly^ℵ modelled points
- 5. try a more complex model, if better BIC set as best and go back to 4
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