Statistical Data Analysis – a course map

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http://cw.felk.cvut.cz/wiki/courses/b4m36san/start

B4M36SAN

Purpose

- This course mainly aims at the statistical methods that help to understand, interpret, visualize and model potentially high-dimensional data. It works with R environment.
- Interactions with other courses



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IDA Highlights



publications



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Abstract

NO. MER

Background sion data in terms of a priori-defined gene sets has recently received significant attention a Analysis of gene expres this approach typically yields more compact and interpretable results than those produced by traditional

Formula display: MathJax

Journal of **Biomedical** Semantics



C Handled Control



18th International Conference on Inductive Logic Programming Prague 2008





organizing conferences

SUPREME*





software projects

The key terms

- Multivariate statistical analysis
 - concerned with data that consists of sets of measurements on a number of individuals,
 - statistical approach based on stochastic data models
 - * a certain model is assumed (a class of models),
 - * its parameters are learned based on data,
 - more than independent testing of the individual variables (i.e., univariate tests known from introductory statistical courses),
 - intertwined variables, **examined simultaneously**,
 - not only the extensions of univariate and bivariate procedures,
 - examples: multivariate analysis of variance, multivariate discriminant analysis.

The key terms

- Applied statistics
 - in general, rather a branch of study than a course,
 - in here, the course could be understood as an opportunity to bring the (previously learned) methods to practice,
 - in labs, stress on applications and their implementation in R.
- Statistical inference/learning
 - close interaction with (statistical) machine learning,
 - sometimes it is difficult to distinguished these two fields
 - * as their goals are interchangeable,
 - the most striking distinctions
 - * different schools statistics is a subfield of mathematics, machine learning is a subfiled of computer science,
 - * different eras for centuries versus modern,
 - * different degree of assumptions larger versus smaller.

B4M36SAN and quotes

- **::** Data do not give up their secrets easily. They must be tortured to confess. Jeff Hopper, Bell Labs
- :: All models are wrong, but some models are useful. George Box, Princeton University
- :: There are two kinds of statistics ...

... the kind you look up and the kind you make up.

Rex Stout, writer

Syllabus

#	Lect	Content
1.	JK	Introduction, course map, review of the basic stat terms/methods.
2.	JK	Dimension reduction (PCA and kernel PCA).
3.	JK	Dimension reduction (other non-linear methods).
4.	JK	Clustering (basic methods, spectral clustering).
5.	JK	Clustering (biclustering, semi-supervised clustering).
6.	JK	Multivariate confirmation analysis (ANOVA and MANOVA).
7.	JK	Discriminant analysis (categorical, LDA, logistic regression).
8.	JK	Multivariate regression (continuous, linear regression, p-vals, overfitting).
9.	JK	Multivariate regression (non-linear, polynomial and local regression).
10.	ΤP	Anomaly detection.
11.	ΤP	Robust statistics.
12.	ZM	Empirical studies, their design and evaluation.
13.	ZM	Power analysis.

R package

R – the platform selected for labs

- the leading tool for statistics,
- one of the main tools in data analysis and machine learning,
- it is free, open-source and platform independent,
- a large community of developers and users \rightarrow a great variety of libraries, tutorials, mailing lists,
- easy to integrate with other languages (C, Java, Python),
- we actually use it,
- bottlenecks in memory management, speed, and efficiency,
- alternatives
 - Python with its data analysis libraries (more general use),
 - Matlab (popular at FEL for its forte in control, Simulink etc.).

- 150 male Egyptian skulls from 5 different periods (30 skulls per group)
 - the early predynastic period (circa 4000 BC),
 - the late predynastic period (circa 3300 BC),
 - the 12th and 13th dynasties (circa 1850 BC),
 - the Ptolemiac period (circa 200 BC),
 - the Roman period (circa AD 150),
- 4 anthropometric measures collected for each skull
 - the maximum breadth (V1, mb),
 - the basibregmatic height (V2, bh),
 - the basialveolar length (V3, bl),
 - the nasal height (V4, nh).



research questions

- are there any differences in the skull sizes between the time periods?
- do they show any (gradual) changes with time?
- how are the four measurements related?
- note: a change in skull size over time could be an evidence of the interbreeding of the Egyptians with immigrant populations over the years.
- statistical tasks (performed in R)
 - data visualization and understanding (exploratory analysis),
 - are there any differences in the skull sizes between the time periods?
 - do they show any changes with time?
 - discriminant analysis, can we tell the period from the skull measurements?

- upload into R, analysis there too,
- see the skript *skulls*.*R* at the course page,
 - > library(HSAUR) # load the library, must be installed before
 > data("skulls", package = "HSAUR") # load into the workspace
 > skulls # see the content of the data.frame
 - epochmbbhblnh1c4000BC13113889492c4000BC12513192483c4000BC13113299504c4000BC1191329644
 - > help(skulls) # see the info about the dataset
 > summary(skulls) # see the basic univariate stats

use boxplots to visualize the data (univariate analysis)



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• use scatterplots to see interactions between variable group means



• employ **dimension reduction** to understand the epoch distribution



Dimension reduction, PCA

• only 64% of variance captured in this plot.

- cluster analysis may help to understand object distribution in skulls data,
- run hierarchical clustering and inspect the resulting **dendrogram**,



 conclusion: no evidence for the existence of 5 clusters, however epochs not randomly distributed along x axis.

- cluster analysis may propose the optimal number of clusters in skulls data,
- the expected/hoped-for value is 5, i.e., the number of epochs,



• run gaussian mixture model for different k values, compare in terms of **BIC**.

- multiple dependent variables \rightarrow multivariate analysis,
- assume that time is a discrete variable, employ MANOVA (multivariate analysis of variance)

$$-H_0: \mu_{4000BC} = \mu_{3300BC} = \cdots = \mu_{AD150}$$

- $-H_a: \mu_{ik} \neq \mu_{jk}$ for at least one $i \neq j$ and at least one variable k
- assumptions: normal within-group distributions, and equal covariance matrices across groups,

MANOVA outcome

 Df
 Wilks approx F num Df den Df
 Pr(>F)

 epoch
 4 0.66359
 3.9009
 16 434.45 7.01e-07 ***

 Residuals 145

 conclusion: it is very likely that there are differences in the skull sizes between the time periods.

test MANOVA assumptions

- start with the homogeneity of covariance matrices, Box's M-test,

- the clusters for the individual epochs must have a similar shape,

> boxM(skulls[c(2:5)], skulls\$epoch)
Chi-Sq (approx.) = 45.667, df = 40, p-value = 0.2483

• conclusion: H_0 about homogeneous covariance matrices cannot be rejected.



- test MANOVA assumptions
 - continue with the normal within-group distributions,
 - MANOVA not sensitive to shape deviations, rather to outliers,
 - see PCA plot to get an overall image,
 - perform an outlier test based on Mahalanobis distance, remove outliers.



- why MANOVA?
 - it has a higher statistical power than (a series of) simple tests,
 - avoids multiple comparisons and problems with their corrections,
- in our case, e.g., **t-test** provides a simple alternative
 - the null hypothesis is that the means of two populations are equal,
 - assumes normal distribution, in here Welch's test for unequal variances,
 - experiment: test two most distant epochs and the most promising variable,



t-test call and outcome

t = 4.0004, df = 56.716, p-value = 0.0001852 alt hypothesis: true difference in means is not equal to 0 95 percent confidence interval: 2.83 8.50 sample estimates: mean of x 99.2 mean of y 93.5

conclusion is same as in MANOVA

- there are differences in mean skull sizes between epochs, namely in bl,

however, p-value is higher and should further be corrected.

- assume that time is an ordinal variable, perform a trend test,
- in particular, **Jonckheere-Terpstra** test for ordered differences among classes

$$-H_0: \theta_{4000BC,k} = \theta_{3300BC,k} = \cdots = \theta_{AD150,k}$$

- $-H_a: heta_{4000BC,k} \ge heta_{3300BC,k} \ge \cdots \ge heta_{AD150,k}$ (at least 1 strict inequality),
- non-parametric rank-based test (θ stands for group medians),
- not multivariate, perform independently for the individual variables,
- > jonckheere.test(skulls\$bl,skulls\$epoch)

Jonckheere-Terpstra test

JT = 2989, p-value = 5.281e-07
alternative hypothesis: two.sided

• conclusion: *bl* manifests a monotonic trend.

assume that time is a real variable, perform multiple linear regression,

$$-year = \beta_0 + \beta_1 mb + \beta_2 bh + \beta_3 bl + \beta_4 nh$$

- in this task, a non-realistic linearity and additivity assumption,
- multivariate, partial effects of the individual variables on the model,
- > skulls_lm <- lm(year \sim mb+bh+bl+nh,data=skulls)

Coeffs	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	-3687.40	4946.86	-0.745	0.457235	
mb	96.40	24.19	3.986	0.000106	***
bh	-29.38	24.40	-1.204	0.230384	
bl	-109.03	22.35	-4.877	2.8e-06	***
nh	65.64	36.85	1.782	0.076918	•

Multiple R-squared: 0.2957, Adjusted R-squared: 0.2763 F-statistic: 15.22 on 4 and 145 DF, p-value: 2.06e-10

- perform multiple linear regression with two independent variables only
 - $-year = \beta_0 + \beta_1 mb + \beta_2 bl$
 - take only those with partial effect on the model,
 - > skulls_lm_small <- lm(year \sim mb+bl,data=skulls)

Coeffs	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	-4802.12	4096.13	-1.172	0.243	
bl	-115.18	21.74	-5.298	4.20e-07	***
mb	105.04	23.91	4.394	2.12e-05	***

- Multiple R-squared: 0.2761, Adjusted R-squared: 0.2662 F-statistic: 28.03 on 2 and 147 DF, p-value: 4.873e-11
- about the same performance in terms of Adjusted R-squared as before.

multiple linear regression, assumption violations through residual analysis



- the model is unbiased, i.e., the error is not dependent on skull measures,
- the relationship is probably not linear as the error is not normally distributed.

- multiple linear regression plane,
- for small models brings similar information as the previous residual plot.



- can we tell the period from the skull measurements?
- perform linear discriminant analysis,
- find linear combinations of features that separate two or more classes of objects
 - maximize the difference between class means and minimize class variances,



linear discriminant analysis, projections into new bases.



linear discriminant analysis, classification accuracy

- > dis1 <- lda(epochs \sim mb+bh+bl+nh, skulls)
- > plot(dis1, col = rainbow(5)[skulls\$epoch])
- > table(skulls\$epochs,predict(dis1, skulls)\$epochs)

	01A	02B	18B	33B	40B
01A	11	9	4	4	2
02B	12	5	7	3	3
18B	5	2	15	4	4
33B	3	4	5	8	10
40B	2	4	4	8	12

- around 34% accuracy on train data,
- namely intermediate epochs difficult to discriminate.

- Conclusions, answers to previous questions
 - the skull measures developed during time,
 - maximum breadth and basialveolar length manifest monotonic change with epochs,
 - the epoch cannot be reliably reconstructed solely from the skull measures,
 - there are no obvious additional dependencies between skull measures.

The key prerequisities – a brief review

- probability, independence, conditional probability, Bayes theorem,
- random variables, random vector,
- their description, distribution function, quantile function,
- categorical and continuous random variables,
- characteristics of random variables,
- the most common probability distributions,
- random vector characteristics, covariance, correlation, central limit theorem,
- measures of central tendency and dispersion, sample mean and variance,
- point and interval estimates of population mean and variance,
- maximum likelihood estimation, EM algorithm,
- statistical hypotheses testing,
- parametric and non-parametric tests,
- multiple comparisons problem, family wise error rate and false discovery rate.

The main references

- :: Resources (slides, scripts, tasks) and reading
 - G. James, D. Witten, T. Hastie and R. Tibshirani: An Introduction to Statistical Learning with Applications in R. Springer, 2014.
 - A. C. Rencher, W. F. Christensen: Methods of Multivariate Analysis.
 3rd Edition, Wiley, 2012.
 - T. Hastie, R. Tibshirani and J. Friedman: The Elements of Statistical Learning: Data Mining, Inference, and Prediction. Springer, 2009.