



Conference

Programme

SCAN ME

DISP

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ΑΘΗΝΩΝ

ΠΑΝΕΠΙΣΤΗΜΙΟ

Topics of Interest:

• Categorical Data Analysis

ATHENS UNIVERSITY

OF ECONOMICS

AND BUSINESS

ΤΜΗΜΑ ΣΤΑΤΙΣΤΙΚΗΣ

DEPARTMENT

- Graphical Models
- Statistical Models for Financial Analysis
- Latent Variable Models
- Longitudinal Data
- MCMC Methods
- Marginal Models
- Non-Parametric Statistics
- Time Series
- Biostatistics
- Big Data Analytics
- High-Dimensionality Models
- Statistical Process Monitoring



Società Italiana di Statistica

28-30 June 2023

Department of Political Science University of Naples Federico II Aula Spinelli, Via Leopoldo Rodinò, 22 Naples

Statistics@Naples Book of Abstract

The **Statistics@Naples** Conference is the 6th meeting on Statistics held in Naples, organized by the Department of Political Science, University of Naples Federico II and the Department of Statistics, Athens University of Economics and Business.

Edited by: Lucio Palazzo

This is the book of abstract of the sessions admitted to the conference call. For more detailed information about Statistics@Naples please refer to: www.indico.unina.it/e/statistics-at-naples

Napoli, June 27, 2023

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WiFi Connection

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Special Events

- June 28 18:30 Narrated walk in the heart of the historic centre. www.insolitaguida.it
- June 29 19:30 Social Dinner at Palazzo Venezia. www.palazzovenezianapoli.com
- June 30 15:30 The Holy Mile (the Miglio Sacro) is a sacred, one-mile long itinerary, from the tomb of St. Gennaro to his Treasure. www.catacombedinapoli.it/en/percorsi/theholymile

Book your ticket at the registration desk!

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We are pleased to welcome you to **Statistics@Naples**, the 6th meeting on Statistics in Naples.

This series of conferences was initiated as a collaboration between the University of Athens and the University of Pavia. This idea was originally proposed by Petros Dellaportas and Guido Consonni in order to strengthen collaboration between the Greek and the Italian statistical communities and their collaborators.

The inaugural conference was organized by Claudia Tarantola and Ioannis Ntzoufras at Mati of Marathonas, just outside of Athens with the title "*First Athens-Pavia Meeting on Statistics*". On that occasion, the National Technical University was also actively involved in the organization of the conference through Dimitris Fouskakis who became one of the main organizers of these meetings on statistics.

The second meeting was held in 2010 in Sardinia under the title "*Second Greek-Italian Meeting on Statistics*". This meeting was also remembered (at least by the Greek members of the meeting) for the cancellation of the Easy-Jet flight to Sardinia. Some Greek delegates arrived a day late, while others returned to Athens. This meeting owes its success to the remarkable efforts of Clelia Di Serio and her team. Both meetings were a great success, with around 50 participants and high quality speeches.

In 2010-2015, no new meetings were announced, as the crisis hit both organising countries hard. In 2014, the meeting series was revived again under the name "3rd Meeting on Statistics" and the more elegant nickname "Statistics3@Athens". The new name was deliberately more neutral to encourage our non-Greek and non-Italian collaborators to attend. The meeting was held in June 2015 at the Athens University Museum under the Acropolis rock. The entire conference focused on the Acropolis and the old city of Athens. Again, the conference was a success with over 60 participants and a very interesting poster session. At the end of an excellent dinner under the lights of the Acropolis, late in the evening the Greek Prime Minister announced the referendum to approve the EU's economic measures: this was the beginning of capital controls and two difficult years for Greece.

The 4th meeting on Statistics was held at the University of Florence (*Statistics4@Florence*) with a similar number of participants. The conference ended successfully with the Greek members wondering where to organize the next meeting and how they can compete with the choice of the beautiful city of Florence, while the Italian delegates were clamouring for a conference on a Greek island.

The island of Aegina was chosen for the fifth meeting conference, mainly because of its location and its easy proximity to Athens. At this conference there were 53 participants with 37 papers and 10 poster presentations. The meeting ended with an enjoyable lecture by Professor Dimitris Karlis with the title "The funny side of Statistics" (highly recommended - do not miss it) and a dinner in a local restaurant. This year the Conference will be hosted in Naples following the scientific collaboration between loannis Ntzoufras, Maria lannario and Claudia Tarantola. The latter, between a scientific contribution and a discussion on a possible double degree, thought of extending the series to one of the Italian places that for assonance and characteristics most closely reflects the Greek way of life. 65 participants, 28 papers and two invited speakers will enliven the conference. Naples will warmly welcome colleagues not only from Italy and Greece but from as many as several different countries, including non-European countries. One invited talk will be given by Ralitza Gueorguieva, Yale School of Public Health, and one talk (which will present some surprises) will be given by Petros Dellaportas, University College London and Athens University of Economics and Business. The Conference will be closed by Dimitris Karlis discussing "The still funny side of Statistics". There will be no shortage of social initiatives and the discovery of places less known to tourists in the city with two tours dedicated to participants and promoted by cultural associations working to regenerate the area, this year celebrating the victory at the football championship. Furthermore, a social dinner in the historic Palazzo Venezia, a 14th-century building located in the heart of Spaccanapoli, will allow participants to experience the flavours of the city.

The conference is jointly organized by the Department of Political Science, University of Naples Federico II and the Department of Statistics, Athens University of Economics and Business, and is supported by the SIS, the Italian Statistical Society.

Finally, in this electronic book, you can find the complete list of abstracts of the oral and poster presentations of the 6th Statistics@Naples meeting. A detailed index of all authors can be found at the end to facilitate an easy search. We deeply thank all contributors for having submitted their works to the conference and the prestigious keynote speakers: Ralitza Gueorguieva and Petros Dellaportas.

We hope you enjoy Statistics@Naples and your visit to the historical city.

On behalf of the local organizing and the scientific committee of Statistics@Naples. Maria lannario

Contents

Conference Program	8
Wednesday, 28 June	8
Thursday, 29 June	9
Friday, 30 June	10
Sessions	11
Subdata selection for big data regression based on leverage scores	11
Bayesian MANOVA for the combined evaluation of handwriting evidence	12
Objective Shrinkage Priors Via Imaginary Data	13
Variable selection via ranking in generalized linear models	14
Bayesian networks for complementing and building gender equality composite indicators	15
A structural equation model to integrate item responses, response times and item	
positioning in students' ability assessment	16
A model-robust subsampling approach in presence of outliers	17
Modelling ordinal data from repeated surveys	18
$arphi$ -Divergence based Modelling of Categorical and Rank Data $\ldots \ldots \ldots \ldots \ldots$	19
Fast Bayesian Variable Screening Using Correlation Thresholds	20
Assessing replication success via skeptical mixture priors	21
Power Prior's Weight Parameter elicitation via Bayes Factor-Calibrated p-values $\ . \ . \ .$	22
Maximum likelihood estimation of multivariate regime switching Student-t copula models	23
Latent Feeling and Uncertainty of Perception and Expectations of Price levels over time:	
A Change Point Analysis	24
Capturing Correlated Clusters Using Mixtures of Latent Class Models	25
Regularized Joint Mixture Models	26
Semiparametric regression for competing risks data with missing not at random cause of	
failure	27
Bayesian effect measures for a location scale model	28
On the predictability of a class of ordinal data models	29
Sparse Pairwise Likelihood Inference for Multivariate Time Series Models	30
Bayesian learning of network structures from interventional experimental data	31
Integrating model-based clustering and graphical models to explore the	
relationship with the digital self-image in (pre)adolescents	32
Sparse Hierarchical Vector Autoregression for Psychopathological Network Estimation	
from Intensive Longitudinal Data	33
How to peel the network: an algorithm for weighted triad census	34
Network Integration with INet algorithm	35
An approach to structural equation modeling in a multiblock framework	36
Plenary Sessions	37
Mixture Models for Repeatedly Measured Survey Data	37
MCMC or Reservoir computing? A direct sampling approach	38

Conference Program



- **CT** Contributed Talk
- KL Keynote Lecture

Wednesday, 28 June

13:00-14:30	Registration			
14:30-15:00	Welcome Address, speakers: Corrado Crocetta and Francesco Palumbo			
15:00-16:30	First Session, chair: Corrado Crocetta			
15:00-15:20	СТ	Vasilis Chasiotis Athens University of Economics and Business	Subdata selection for big data regression based on leverage scores	
15:20-15:40	СТ	Lampis Tzai University of Lausanne	Bayesian MANOVA for the combined evaluation of handwriting evidence	
15:40-16:00	СТ	Dimitrios Fouskakis National Technical University of Athens	Objective Shrinkage Priors Via Imaginary Data	
16:00-16:20	СТ	Marcella Niglio University of Salerno	Variable selection via ranking in generalized linear models	
16:30-17:00	Coffee Break			
17:00-18:30	Second Session, chair: Guido Consonni			
17:00-17:20	СТ	Paola Vicard Roma Tre University	Bayesian networks for complementing and building gender equality composite indicators	
17:20-17:40	СТ	Rosa Fabbricatore University of Naples Federico II	A structural equation model to integrate item responses, response times and item positioning in students' ability assessment	
17:40-18:00	СТ	Laura Deldossi Università Cattolica del Sacro Cuore	A model-robust subsampling approach in presence of outliers	

Thursday, 29 June

09:30-11:00	Third Session, chair: Dimitrios Fouskakis			
09:30-09:50	СТ	Marcella Corduas University of Naples Federico II	Modelling ordinal data from repeated surveys	
09:50-10:10	СТ	Maria Kateri RWTH Aachen University	arphi-Divergence based Modelling of Categorical and Rank Data	
10:10-10:30	СТ	Ioannis Ntzoufras Athens University of Economics and Business	Fast Bayesian Variable Screening Using Correlation Thresholds	
10:30-10:50	СТ	Leonardo Egidi University of Trieste	Assessing replication success via skeptical mixture priors	
11:00-11:50		Plenary Session, chair: Maria Iannario		
11:00-11:40	KL	Ralitza Gueorguieva Yale University	Mixture Models for Repeatedly Measured Survey Data	
11:50-12:30	Coffee Break			
12:30-14:00		Fourth Session, ch	air: Ioannis Ntzoufras	
12:30-12:50	СТ	Roberto Macrì Demartino University of Padova	Power Prior's Weight Parameter elicitation via Bayes Factor-Calibrated p-values	
12:50-13:10	СТ	Federico Cortese Bicocca, Milan	Maximum likelihood estimation of multivariate regime switching Student-t copula models	
13:10-13:30	СТ	Carmela Cappelli University of Naples Federico II	Latent Feeling and Uncertainty of Perception and Expectations of Price levels over time: A Change Point Analysis	
13:30-13:50	СТ	Gertraud Malsiner-Walli Vienna University of Economics and Business	Capturing Correlated Clusters Using Mixtures of Latent Class Models	
14:00-15:00		Lunch Break		
15:00-16:30		Fifth Session , chai	r: Francesco Palumbo	
15:00-15:20	СТ	Konstantinos Perrakis Durham University	Regularized Joint Mixture Models	
15:20-15:40	СТ	Giorgos Bakoyannis Athens University of Economics and Business	Semiparametric regression for competing risks data with missing not at random cause of failure	
15:40-16:00	СТ	Claudia Tarantola University of Pavia	Bayesian effect measures for a location scale model	
16:00-16:20	СТ	Rosaria Simone University of Naples Federico II	On the predictability of a class of ordinal data models	
16:30-17:00	Coffee Break			
17:00-18:30	Workshop, Discussion, Planning for possible Joint Work			
19:30-22:30		Social Dinner		

Friday, 30 June

09:30-11:00	Sixth Session, chair: Paola Vicard		
09:30-09:50	СТ	Xanthi Pedeli Athens University of Economics and Business	Sparse Pairwise Likelihood Inference for Multivariate Time Series Models
09:50-10:10	СТ	Stefano Peluso Bicocca, Milan	Bayesian learning of network structures from interventional experimental data
10:10-10:30	СТ	Chiara Brombin San Raffaele University	Integrating model-based clustering and graphical models to explore the relationship with the digital self-image in (pre)adolescents
10:30-10:50	СТ	Spyros Balafas UNISR	Sparse Hierarchical Vector Autoregression for Psychopathological Network Estimation from Intensive Longitudinal Data
11:00-11:50	Plenary Session, chair: Claudia Tarantola		
11:00-11:40	KL	Petros Dellaportas Athens University of Economics and Business	MCMC or Reservoir computing? A direct sampling approach
11:50-12:30		Coffee Break	
12:30-13:30		Seventh Session, chair: Clelia Di Serio	
12:30-12:50	СТ	Roberto Rondinelli University of Naples Federico II	How to peel the network: an algorithm for weighted triad census
12:50-13:10	СТ	Valeria Policastro University of Naples Federico II	Network Integration with INet algorithm
13:10-13:30	СТ	Rosaria Romano University of Naples Federico II	An approach to structural equation modeling in a multiblock framework
13:30-14:00	Closing Plenary Session, speaker: Dimitris Karlis		
14:00-15:00	Lunch Break		

Sessions

Subdata selection for big data regression based on leverage scores

Vasilis Chasiotis, Dimitris Karlis

First Session, Wednesday 28 June

15:00 - 15:20

Leverage Scores; Subdata; Large Datasets

Data continues to become more abundant, and so the datasets that contain it. Even though big datasets can present insights and opportunities, they can pose significant challenges when it comes to statistical analysis. One of the biggest challenges, required to process and analyze large datasets, is the computational resources. Regression can be problematic in case of big datasets, due to the huge volumes of data. A standard approach is subsampling that aims at obtaining the most informative portion of the big data.

We consider an approach based on leverages scores, already existing in the current literature for the selection of subdata for linear model discrimination. However, we highlight its importance on the selection of data points that are the most informative for estimating unknown parameters. We conclude that the approach based on leverage scores improves existing approaches, providing simulation experiments as well as a real data application.

Bayesian MANOVA for the combined evaluation of handwriting evidence

Lampis Tzai

First Session, Wednesday 28 June

15:20 - 15:40

Handwriting Evidence; Fourier Analysis; Multivariate Bayesian Modelling; Bayes Factor; Sensitivity

Forensic science is a broad field that uses scientific principles and technical methods to help with the evaluation of evidence in legal proceedings of criminal, civil, or administrative nature. Forensic scientists examine recovered traces that can be given by glass fragments, fingerprints, body fluids, textile fibers, digital device data and handwriting. Consider a case involving a handwritten anonymous document. Handwritten features extracted from this questioned document will be compared to those extracted from a document written by a person that is suspected of being the origin of the anonymous document. The propositions of interest are the following: H_p : the suspect is the author of the manuscript; H_d : the suspect is not the author of the manuscript. Handwriting individualization is still largely dependent on the experience of the document examiner, though studies have been conducted with the aim of supporting handwriting examiners to reduce the degree of subjectivity of their expertise. Marguis et al. (2005) proposed to increase the degree of objectivity of handwriting analyses by implementing elements of Fourier analysis in order to describe the contour shape of loops of characters. Specifically, the characters containing loops can be described by means of Fourier descriptors, which can be used to characterize the shape complexity and other geometric attributes. The analyses conducted showed that these features have a good discriminating power. To implement the use of these handwriting features for handwriting identification, Bozza et al. (2008) proposed a Bayesian probabilistic approach by modeling the data with multivariate Normalinverse-Wishart distribution (NIW). The value of the evidence is subsequently assessed by means of the Bayes factor, which can be interpreted as a measure of the strength of support provided by the evidence in favor of the hypothesis H_p against the hypothesis H_d . This approach was accomplished to take into account the correlation between variables, the variability between-writers and within-writer variability. However, the above model is implemented separately for each different type of handwritten character. This can be problematic because it can lead to a different conclusion depending on the character that is retained.

In this research, it is proposed the implementation of a Bayesian Multivariate Analysis of Variance (MANOVA) via using the loop characters as predictors. The indicator of the loop character is transformed into a dummy variable (corner-point representation), so that it is possible to model variables describing the handwriting characters jointly taking into account the variability between characters, the variability between-writers for every character and withinwriter variability. The Bayesian MANOVA is compared with the two-level random effect model (NIW) proposed by Bozza et al. (2008), that is implemented by modelling all characters jointly or separately. Three different methods for estimating the marginal likelihoods are used; the Generalized Harmonic Mean, the Laplace-Metropolis and the Bridge Sampling. Finally, the performances of the NIW and MANOVA models are compared with those of an alternative one, where a conjugate approach is chosen. This does not allow to model the within and between variation separately, but the marginals can be obtained analytically. Firstly, we estimate the Bayes factor of the two data models for each writer to determine which model is more compatible with the data. Secondly, there have been selected handwriting features originating from the same writer or from different writers to evaluate the rate of false negatives (when the BF is smaller than one for characters originating from the same source) and false positives (if the BF is greater than one for characters originating from different sources). Finally, the sensitivity of models is examined in two critical aspects: the misleading background information and the choice of degrees of freedom for the Wishart-inverse distribution that is used to model the handwriting variability. With reference to the misleading background information, the prior distributions were elicited by selecting writers characterized by either small or marked differences.

References

Bozza, S., Taroni, F., Marquis, R. & Schmittbuhl, M. (2008), 'Probabilistic evaluation of handwriting evidence: likelihood ratio for authorship', Journal of the Royal Statistical Society: Series C (Applied Statistics) 57(3), 329-341. Marquis, R., Schmittbuhl, M., Mazzella, W. D. & Taroni, F. (2005), 'Quantification of the shape of handwritten characters: a step to objective discrimination between writers based on the study of the capital character o', Forensic Science International 150(1), 23-32.

Objective Shrinkage Priors Via Imaginary Data

Dimitrios Fouskakis, George Tzoumerkas

First Session, Wednesday 28 June

Bayesian Variable Selection; Imaginary Data; Objective Priors; Shrinkage Priors

In this work, focus is given in the Bayesian variable selection problem for high-dimensional linear regression problems. The use of shrinkage priors, when the number n of available observations is less than the number p of explanatory variables, is a well-established method, which shares great theoretical and empirical properties.

By using imaginary data and shrinkage priors as baseline priors, under the Power-Expected-Posterior (PEP) prior methodology, objective shrinkage priors are being created. In addition, we explore the idea of augmenting the imaginary design matrix in order to make it with orthogonal columns and thus to produce independent PEP-shrinkage priors, based on default baseline priors.

Under this setup, properly chosen hyperpriors are placed on the power parameters of the PEP methodology, in order to produced mixtures of independent priors suitable for the variable selection problem when $n \ll p$. This second approach provides us with algorithmically flexibility and less time-consuming procedures. We check the theoretical properties of our proposed methods and we explore their behavior via simulated studies.

15:40 - 16:00

Variable selection via ranking in generalized linear models

Francesco Giordano, Marcella Niglio, Marialuisa Restaino

First Session, Wednesday 28 June

16:00 - 16:20

Generalized Linear Models; Ranking Based Variable Selection; BIC; Sure Independence Screening

In many empirical domains, the availability of ultrahigh-dimensional data has led to the development of feature screening and variable selection procedures aiming to detect the informative variables of datasets and consequently remove unimportant features.

In this context, we propose a ranking-based variable selection procedure that extends the Ranking Based Variable Selection technique (Baranowski et al., 2020) to general linear regression models. We explore the performance of our proposal using both simulated and empirical data.

The algorithm is compared to two competitors: i) the Extended BIC (Chen and Chen, 2012); ii) the variable selection procedure based on the combination of the Sure Independence Screening (Fan and Song, 2010) and the Elastic Net (Zou and Hastie, 2005).

References

Baranowski R, Chen Y, Fryzlewicz P (2020), Ranking-based variable selection for high-dimensional data, Statistica Sinica, 30(3), 1485-1516.

Chen J, Chen Z (2012), Extended BIC for small-n-large-P sparse GLM. Statistica Sinica, 22(2), 555-574.

Fan J, Song R (2010), Sure independence screening in generalized linear models with NP-dimensionality. The Annals of Statistics, 38(6), 3567-3604.

Zou H; Hastie T (2005), Regularization and Variable Selection via the Elastic Net. Journal of the Royal Statistical Society, Series B, 67(2), 301-320.

Bayesian networks for complementing and building gender equality composite indicators

Paola Vicard, Lorenzo Giammei, Flaminia Musella, Fulvia Mecatti

Second Session, Wednesday 28 June

17:00 - 17:20

Composite Indicator; Gender Equality; Multivariate Dependencies; Object-oriented Bayesian Networks

Composite indicators are a common choice for synthesizing complex phenomena. Over the years, they have grown in popularity and are now applied in many social and environmental sciences. Among others, a subject of increasing interest is gender equality analysis. Gender composite indicators, even if easy to read, may provide a limited picture of the problem. Here we discuss the potentiality of Bayesian networks (BNs) to complement and build composite indicators. BNs are powerful tools for explaining the complex association structure in the dataset and developing scenarios to orient policy-making.

Here we propose to use BNs to model the association structure among the gender equality index, its ingredient variables and other context socio-economic variables. In such a way the synergy between composite indicator and BN gives rise to both a monitoring tool for the gender equality gap status and a proactive inferential machine for proposing policies to reduce inequality. BNs can be also used to build the gender equality index, and, in general, any composite indicator.

Specifically, we focus attention on an extension of BNs, namely Object-Oriented Bayesian networks (OOBNs). The modularity of the OOBN ensures a computational logic that is consistent with composite indicators, while also providing additional information about the relational structure of variables. An example is carried out on Italian province-level data.

References:

Cowell, R. G., Dawid, A. P., Lauritzen, S. L., and Spiegelhalter, D. J. (1999). Probabilistic Networks and Expert Systems. Springer Verlag, New York

Musella, F., Vicard, P. (2015). Object-oriented Bayesian networks for complex quality management problems. Quality & Quantity, 49, 115-133

A structural equation model to integrate item responses, response times and item positioning in students' ability assessment

Silvia Bacci, Rosa Fabbricatore, Carla Galluccio, Maria Iannario

Second Session, Wednesday 28 June

17:20 - 17:40

Ability Assessment; Structural Equation Modeling; Moodle platform

In the context of students' ability assessment, considering collateral information in addition to item responses can be helpful in increasing the accuracy of the measurement. In this vein, the evaluation of students' abilities via computer based-devices has made response time data available at the item level (Wang et al., 2019). Besides, the literature (Becker et al., 2022) has highlighted an item position effect when the same items are presented in different positions within multiple test forms.

With the present contribution, we contribute to this research line by proposing a structural equation model (SEM) to jointly consider item responses, response times and item positioning in students' ability assessment. In particular, we assume that the response process is driven by two underlying latent variables: the first latent variable, denoted by Θ_i , represents the ability of individual *i* that is measured by the test items; the second latent variable, denoted by η_i , refers to the speediness of individual *i* to answer the test items.

We formulate the statistical model assuming that the item responses are directly affected by the ability Θ_i , whereas the response times depend both on the ability Θ_i and on the speediness η_i . Accordingly, response accuracy tends to increase with the ability level of individual *i* while response time tends to decrease with the speediness and ability levels. Moreover, we suppose that item positioning affects both item responses and response time. Under this setting, the correlation between Θ_i and η_i is modelled through the cross-relation function that models the relationships between Θ_i and the observed response times.

The empirical application of the proposed model was carried out on first-year Psychology students at the University of Naples Federico II, attending the introductory Statistics course. The test administered was composed of 30 multi-choice questions developed according to three of the five Dublin descriptors: Knowledge (10 items), Application (10 items) and Judgement (10 items). For each question, students' answers were coded as correct (2 credits), partially correct (1 credit) and wrong (0 credits). Data were collected through Moodle platform, which also provided the response time.

References Becker, B., Van Rijn, P., Molenaar, D., and Debeer, D. (2022). Item order and speededness: Implications for test fairness in higher educational high-stakes testing. Assessment & Evaluation in Higher Education, 47(7):1030-1042. Wang, C., Weiss, D. J., and Su, S. (2019). Modeling response time and responses in multidimensional health measurement. Frontiers in psychology, 10:51.

A model-robust subsampling approach in presence of outliers

Laura Deldossi, Chiara Tommasi

Second Session, Wednesday 28 June

17:40 - 18:00

Active learning; D-optimality; Subsampling

In the era of big data, several sampling approaches are proposed to reduce costs (and time) and to help in informed decision making. Some of these proposals (Drovandi et al., 2017; Wang et al., 2019; Deldossi and Tommasi (2022) among others) are inspired to Optimal Experimental Design and require the specification of a model for the big dataset. This model assumption, as well as the possible presence of outliers in the big dataset represent a limitation for the most commonly applied subsampling criterions.

Deldossi et al. (2023) introduced non-informative and informative exchange algorithms to select "nearly" D-optimal subsets without outliers in a linear regression model. In this study, we extend their proposal to account for model uncertainty. More precisely, we propose a model robust approach where a set of candidate models is considered; the optimal subset is obtained by merging the subsamples that would be selected by applying the approach of Deldossi et al. (2023) if each model was considered as the true generating process. The approach is applied in a simulation study and some comparisons with other subsampling procedures are provided.

References

Deldossi, L., Tommasi C. (2022) Optimal design subsampling from Big Datasets. Journal of Quality Technology 54(1): 93-101

Deldossi, L., Pesce, E., Tommasi, C. (2023) Accounting for outliers in optimal subsampling methods, Statistical Papers, https://doi.org/10.1007/s00362-023-01422-3.

Drovandi CC, Holmes CC, McGree JM, Mengersen K, Richardson S, Ryan EG (2017) Principles of experimental design for big data analysis. Statistical Sciences 32(3): 385-404

Wang H, Yang M, Stufken J (2019) Information-based optimal subdata selection for Big Data linear regression. Journal of American Statistical Association 114(525): 393-405

Modelling ordinal data from repeated surveys

Marcella Corduas

Third Session, Thursday 29 June

09:30 - 09:50

Ordinal Data; CUB Model; Consumers' Perceptions; Consumers' Expectations

Business and consumers survey data are the basis for several indicators describing the trend of macro-economic variables that are fundamental for monitoring the overall performance of the economic system. Qualitative surveys typically ask interviewees to express their perceptions or expectations about the current or future tendency of a reference economic variable (such as inflation or industrial output) using a trichotomous or a finer-tuned ordered scale.

Surveys are carried out at regular interval by statistical offices, and collected data are traditionally published in aggregate form, reporting the proportions of positive, neutral or negative assessments. This contribution presents an innovative dynamic model that describes the probability distributions of ordered categorical variables observed over time. For this aim, we extend the definition of the mixture distribution obtained from the Combination of a Uniform and a shifted Binomial distribution (CUB model), introducing time varying parameters.

The model parameters identify the main components ruling the respondent evaluation process: the degree of attraction towards the object under assessment, the uncertainty related to the answer, and the weight of the refuge category that is selected when a respondent is unwilling to elaborate a thoughtful judgment. We suggest to use the model time-varying parameters as indicators of the diversity of respondents' opinions, shifting from an optimistic to a pessimistic state as the surrounding conditions evolve. For illustrative purpose, the dynamic CUB model is applied to the consumers' perception and expectations of inflation in Italy to investigate: a) the effect of the COVID pandemic on the respondents' perceptions; b) the impact of the respondents' income level on expectations.

References

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Piccolo, D., Simone, R. The class of cub models: statistical foundations, inferential issues and empirical evidence, (with discussion and rejoinder). "Stat. Meth.& Appl." 2019, 28, 389-435.

arphi-Divergence based Modelling of Categorical and Rank Data

Maria Kateri

Third Session, Thursday 29 June

09:50 - 10:10

Cressie-Read Power Divergence; Distance-based Probability Models; Maximum Likelihood Estimation

Standard models for categorical and ordinal data, such as log-linear, association models and logistic regression models for binary or ordinal responses, as well as the Mallows model for rank data are revisited and defined through statistical information theoretic properties in terms of the Kullback-Leibler (KL) divergence.

In the sequel, replacing the KL by the φ -divergence, which is a family of divergences including the KL as special case, these models are generalized to flexible families of models. The suggested models are discussed in terms of their properties, estimation and fit. Finally, their potential is illustrated by characteristic examples.

Fast Bayesian Variable Screening Using Correlation Thresholds

Ioannis Ntzoufras, Roberta Paroli

Third Session, Thursday 29 June

10:10 - 10:30

Zellner's g-prior; Bayesian Variable Screening

We propose a fast Bayesian variable selection method for Normal regression models, using Zellner's *g*-prior specification. The approach is based on using thresholds on Pearson and partial correlation coefficients. Nevertheless, the proposed methodology is derived using purely Bayesian arguments derived from thresholds on Bayes factors and posterior model odds.

The proposed method can be used to screen out the non-important covariates and reduce the model space size. Then, traditional, computer-intensive, Bayesian variable selection methods can be implemented without any problem with the derived reduced model space. We focus on the *g*-prior where the Bayes factor computations and the corresponding correlation thresholds are exact. Nevertheless, the approach is general and can be easily extended to any prior setup. The proposed method is illustrated using simulated examples.

Assessing replication success via skeptical mixture priors

Guido Consonni, Leonardo Egidi

Third Session, Thursday 29 June

10:30 - 10:50

Bayes Factor; Consistency; Prior-data Conflict; Replication Studies

There is a growing interest in the analysis of replication studies of original findings across many disciplines. When testing a hypothesis for an effect size, two Bayesian approaches stand out for their principled use of the Bayes factor (BF), namely the replication BF (Verhagen and Wakenmalers, 2014) and the skeptical BF (Pawel and Held, 2022). In both cases replication data are used to compare an "advocacy" prior against a benchmark. For the replication BF, the latter is the standard point null hypothesis of no effect while for the skeptical BF it represents the prior of somebody who is unconvinced by the original findings.

We propose a novel skeptical mixture prior which incorporates skepticism and limits prior-data conflict. We support our proposal with theoretical results on consistency of the resulting BF, we illustrate its features on an extended example, and we apply it to case studies from the Social Sciences Replication Project.

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Verhagen, J., & Wagenmakers, E. J. (2014). Bayesian tests to quantify the result of a replication attempt. "Journal of Experimental Psychology: General", 143(4), 1457.

Power Prior's Weight Parameter elicitation via Bayes Factor-Calibrated p-values

Roberto Macrì Demartino

Fourth Session, Thursday 29 June

12:30 - 12:50

Beta Distribution; Clinical Trial; Historical Information; Robust Selection

In recent times, the integration of historical data in the design and analysis of new clinical trials has gained considerable attention, owing to ethical reasons and difficulties encountered in recruiting patients. In the Bayesian framework, the process of informative prior elicitation is widely recognized as a complex and multifaceted undertaking, requiring the careful quantification and synthesis of prior information into an appropriate prior distribution. Hence, there is a pressing need for developing techniques and methods that can facilitate synthesizing and quantifying prior information more effectively and efficiently. Within this context, the concept of "power priors" (Chen and Ibrahim, 2000) has emerged as a popular approach for incorporating historical data into the prior distribution of a treatment effect, in a flexible and controlled manner.

The power prior methodology heavily relies on the "weight parameter" δ , ranging between 0 and 1, that is a crucial factor in determining the degree to which the historical data influences the prior distribution, and for which multiple elicitation strategies are available. A modification of the power prior allows a hierarchical prior specification by taking δ as a random quantity $\pi(\theta, \delta \mid D_0) \propto L(\theta \mid D_0)^{\delta} \pi_0(\theta)\pi_0(\delta)$, where D_0 is an historical dataset with corresponding likelihood $L(\theta \mid D_0), \pi_0(\theta)$ and $\pi_0(\delta)$ are the initial priors for θ and δ , respectively. Furthermore, a significant benefit of incorporating a normalizing factor in the power prior methodology is its adherence to the likelihood principle, as demonstrated by the joint normalized power prior

$$\pi\left(\theta,\delta\mid D_{0}\right) = \frac{L\left(\theta\mid D_{0}\right)^{\delta}\pi_{0}(\theta)\pi_{0}(\delta)}{\int_{\Omega}L\left(\theta\mid D_{0}\right)^{\delta}\pi_{0}(\theta)d\theta}.$$

Consequently, in a fully Bayesian approach, the ability to effectively elicit an appropriate initial prior distribution for the weight parameter δ is a crucial step. As far as we know from reviewing the existing literature, a comprehensive justification underlying the choice of a Beta distribution with fixed hyper-parameters, that is an usual choice for this framework, is pretty vague. The Bayes factor (BF) constitutes a valuable statistical tool for model comparison; however, we explore the use of the Bayes Factor to discriminate between competing models that incorporate distinct initial Beta prior distributions for the weight parameter by exploiting some BF *p*-value calibration techniques (Garcia-Donato and Chen, 2005). This would enable the selection of candidate models based on a more accurate and reliable assessment of the available evidence, thereby enhancing the validity and robustness of statistical inference.

References

Chen, M.-H. and Ibrahim, J. G. (2000). Power prior distributions for regression models. Statistical Science, 15(1):46 - 60. Garcia-Donato, G. and Chen, M.-H. (2005). Calibrating Bayes factor under prior predictive distributions. Statistica Sinica, 15(2):359-380.

Maximum likelihood estimation of multivariate regime switching Student-t copula models

Federico Cortese, Fulvia Pennoni, Francesco Bartolucci

Fourth Session, Thursday 29 June

12:50 - 13:10

Statistical Models for Financial Analysis; Cryptocurrencies; Time Series; Expectation-Maximization Algorithm; Latent Variable Models

We propose a novel estimation method for multivariate regime switching models based on a Student-t copula function. These models account for the interdependencies between multiple variables by considering the correlation strength controlled by specific parameters. Moreover, they address fat-tailed distributions through the number of degrees of freedom. These parameters, in turn, are governed by a latent Markov process. We consider a two-steps procedure carried out through the Expectation-Maximization algorithm to estimate model parameters by maximum likelihood.

The primary computational challenge lies in estimating both the matrix of dependence parameters and determining the number of degrees of freedom for the Student t-copula. To address this, we introduce a new approach that leverages Lagrange multipliers, simplifying the estimation process. Through a comprehensive simulation study, we demonstrate that our estimators possess desirable properties in finite samples. Additionally, the estimation procedure shows good computational efficiency.

We apply our model to analyze the log-returns of five different cryptocurrencies. The results enable us to identify distinct bull and bear market periods based on the intensity of correlations observed between the crypto assets. This finding highlights the model's efficacy in capturing and characterizing market dynamics within the cryptocurrency domain.

Latent Feeling and Uncertainty of Perception and Expectations of Price levels over time: A Change Point Analysis

Carmela Cappelli, Rosaria Simone

Fourth Session, Thursday 29 June

13:10 - 13:30

Price Expectation; Price Judgment; Atheoretical Regression Trees; CUB model; Change-point Detection

For the analysis of ordered categorical data, CUB modelling approach entails the estimation of two main structural latent components of the rating process: feeling and uncertainty, parameterized within a two-component mixture of Binomial and uniform distributions: see Piccolo and Simone 2019 for an overview. Featuring parameters can be possibly linked to subject covariates to determine twofold response patterns and they can be promptly estimated using the EM algorithm (as implemented in the R package 'CUB' available on CRAN).

The contribution aims at presenting how change point detection of temporal series of estimated feeling and uncertainty can be pursued to identify if and to what extent Italian people modified their perception and judgments of price levels from 1994 to 2019. To this goal, we resort to the framework of Atheoretical Regression Trees (ART, Cappelli et al. 2008) considering the series of monthly response distributions to questions: 1-(Judgments): How do you think the price level changed over the previous 12 months? 2-(Expectations): How do you think the price level will change over the next 12 months? issued by the Italian National Statistical Institute (ISTAT) within the consumers' confidence survey.

Responses are collected over a scale with m=5 categories (1 = "fall", 2 = "stay about the same", 3 = "rise slightly", 4 = "rise moderately", 5 = "rise a lot"). Preliminary results indicate that ART is effective in partitioning the series into sub-intervals characterized by different levels of the estimated model parameters, allowing to study and compare over time, the change points of both feeling and uncertainty.

It is worth noticing that the model parameters refer to two different aspects of the respondents' perception and judgment of price level, thus the study of their change points may reveal that they show different number and location of break dates providing a further and valuable insight into the two components of respondents' answers. Performances of ART are also discussed comparatively with those of other techniques for structural change point detection, in particular with respect to Bai and Perron's procedure as ART mimics this procedure.

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Capturing Correlated Clusters Using Mixtures of Latent Class Models

Gertraud Malsiner-Walli, Bettina Gruen, Sylvia Fruehwirth-Schnatter

Fourth Session, Thursday 29 June

13:30 - 13:50

Bayesian Inference; Model-based Clustering; Prior on the Number of Components, Telescoping Sampler

Latent class models rely on the conditional independence assumption, i.e., it is assumed that the categorical variables are independent given the cluster memberships. Within the Bayesian framework, we propose a suitable specification of priors for the latent class model to identify the clusters in multivariate categorical data where the independence assumption is not fulfilled. Each cluster distribution is approximated by a latent class model, leading overall to a mixture of latent class models.

The Bayesian approach allows to identify the clusters and fit their cluster distributions using a one-step procedure. We provide suitable estimation and inference methods for the mixture of latent class models and illustrate the performance of this approach on artificial and real data.

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Fop, M., K. M. Smart, and T. B. Murphy (2017). Variable selection for latent class analysis with application to low back pain diagnosis. The Annals of Applied Statistics 11 (4), 2080-2110.

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Regularized Joint Mixture Models

Konstantinos Perrakis, Thomas Lartigue, Frank Dondelinger, Sach Mukherjee

Fifth Session, Thursday 29 June

15:00 - 15:20

Distribution Shifts; Heterogeneous Data; Joint Learning; Latent Groups; Mixture Models; Sparse Regression

Regularized regression models are well studied and, under appropriate conditions, offer fast and statistically interpretable results. However, large data in many applications are heterogeneous in the sense of harboring distributional differences between latent groups. Then, the assumption that the conditional distribution of response Y given features X is the same for all samples may not hold. Furthermore, in scientific applications, the covariance structure of the features may contain important signals and its learning is also affected by latent group structure.

We propose a class of mixture models for paired data (X, Y) that couples together the distribution of X (using sparse graphical models) and the conditional Y|X (using sparse regression models). The regression and graphical models are specific to the latent groups and model parameters are estimated jointly (hence the name "regularized joint mixtures"). This allows signals in either or both of the feature distribution and regression model to inform learning of latent structure and provides automatic control of confounding by such structure. Estimation is handled via an expectation-maximization algorithm, whose convergence is established theoretically. We illustrate the key ideas via empirical examples. An R package is available at https://github.com/k-perrakis/regjmix.

Semiparametric regression for competing risks data with missing not at random cause of failure

Giorgos Bakoyannis, Constantin T. Yiannoutsos

Fifth Session, Thursday 29 June

15:20 - 15:40

Missing Not At Random; Missing At Random; Semiparametric Regression; Multicenter HIV

The cause of failure in cohort studies that involve competing risks is frequently incompletely observed. Failure to deal with this issue can lead to substantially biased estimates. To the best of our knowledge, all the methods that have addressed the issue in the context of semiparametric competing risks models rely on a missing at random (MAR) assumption. Nevertheless, the MAR assumption is not realistic in many real-world settings.

In this work we relax the latter assumption by allowing for a class of missing not at random (MNAR) mechanisms, which contain the MAR mechanism as a special case. Due to the inherent non-identifiability issues under MNAR, we propose an approach for hypothesis testing that does not require the estimation of the non-estimable parameters. Using modern empirical process theory, we show that the proposed estimators are uniformly consistent under the assumed class of MNAR mechanisms. We also show that our estimators converge weakly to tight zero mean Gaussian processes and propose rigorous methodology for the computation of confidence intervals which achieve a coverage rate of at least $100 \times (1 - \alpha)\%$, asymptotically, for the true unknown parameters of interest.

The proposed methodology is applied to competing risks data from a large multicenter HIV study in sub-Saharan Africa where a substantial portion of causes of failure is missing not at random.

Bayesian effect measures for a location scale model

Maria Iannario, Maria Kateri, Claudia Tarantola

Fifth Session, Thursday 29 June

15:40 - 16:00

Rating Data; Markov Chain Monte Carlo; Heteroskedasticity

We consider a Bayesian approach for the analysis of rating data when a scaling component is taken into account, thus incorporating a specific form of heteroskedasticity. Our approach includes model-based probability effect measures that enable comparisons of distributions among multiple groups. These effect measures are adjusted for explanatory variables that have an impact on both the location and scale components.

To estimate the parameters of our fitted model and derive the associated effect measures, we employ Markov Chain Monte Carlo techniques. Through an analysis of students' evaluations of a university curriculum counselor service, we assess the performance of our method and highlight its valuable support in the decision-making process. Our findings demonstrate the effectiveness of our approach and emphasize its ability to enhance decision-making processes by providing valuable insights and support to stakeholders involved.

On the predictability of a class of ordinal data models

Rosaria Simone, Piccolo Domenico

Fifth Session, Thursday 29 June

16:00 - 16:20

CUB models; Predictability; Ranked Probability Score; Ordinal Data

The contribution aims at discussing some preliminary results on the evaluation of prediction performance for the class of mixture models with uncertainty (Piccolo and Simone, 2019). The ultimate goal of the analysis is the evaluation of the extent by which the uncertainty specification constitutes an added value for prediction of ordinal scores. A small simulation study is presented to assess prediction performance of competing models under miss-specification. The Ranked Probability Score is chosen as scoring rule since it is the most suited to deal with ordinal data, without the assignment of numerical scores to category. Finally, a variable selection procedure based on prediction performance can be outlined on a case study for the prediction of subjective probability to survive.

Comparisons with cumulative link models are illustrated for the sake of completeness. Preliminary findings discussed in Simone and Piccolo (2022) indicate that uncertainty modelling improves prediction performance substantially. Hence, it is important to assess the information quality of the baseline preference model (the Binomial, for instance).

To this aim, we introduce a new utility measure for preference models when contaminated with alternative uncertainty specifications in the sense proper to the framework of Information Quality. As a result, the mixing weight of the chosen feeling component within the mixture can be explicitly interpreted in terms of model predictive ability.

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D. Piccolo, R. Simone (2019). The class of CUB models: statistical foundations, inferential issues and empirical evidence. Satistical Method and Applications, Volume 28, pages 389-435.

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Sparse Pairwise Likelihood Inference for Multivariate Time Series Models

Xanthi Pedeli

Sixth Session, Friday 30 June

09:30 - 09:50

Full Likelihood; Pairwise Likelihood; Bivariate Marginals

Multivariate time series data is becoming an increasingly common research topic. Unlike univariate time series, the temporal dependence of a multivariate series includes both serial dependences and interdependences across different marginal series. Consequently, as the number of component series increases, multivariate time series models become overparameterized. In addition, there are many cases where the conditional distribution of the multivariate series given its past might have a complicated form.

Given these challenges we develop methodology by replacing the full likelihood function by a pairwise likelihood that only requires the specification of bivariate marginals instead of the multivariate distribution. Clearly, the computational task of maximization of the pairwise likelihood is much simpler than maximization of the full likelihood function but still it poses the problem of combining all estimators.

For this purpose, we rely on maximization of an approximate weighted least squares estimation criterion subject to a shrinkage penalty that allows for model selection. The suggested approach provides a general framework for multidimensional time series since it can be applied to both continuous and discrete time series but also to mixed mode time series data.

Bayesian learning of network structures from interventional experimental data

Stefano Peluso

Sixth Session, Friday 30 June

09:50 - 10:10

Bayesian Network; Multivariate Data; Directed Acyclic Graphs

Directed Acyclic Graphs (DAGs) provide an effective framework for learning causal relationships among variables given multivariate observations. Under pure observational data, DAGs encoding the same conditional independencies cannot be distinguished and are collected into Markov equivalence classes. In many contexts however, observational measurements are supplemented by interventional data that improve DAG identifiability and enhance causal effect estimation.

We propose a Bayesian framework for multivariate data partially generated after stochastic interventions. To this end, we introduce an effective prior elicitation procedure leading to a closed-form expression for the DAG marginal likelihood and guaranteeing score equivalence among DAGs that are Markov equivalent post intervention. Under the Gaussian setting we show, in terms of posterior ratio consistency, that the true network will be asymptotically recovered, regardless of the specific distribution of the intervened variables and of the relative asymptotic dominance between observational and interventional measurements.

We validate our theoretical results in simulation and we implement on both synthetic and biological protein expression data a Markov chain Monte Carlo sampler for posterior inference on the space of DAGs.

Integrating model-based clustering and graphical models to explore the relationship with the digital self-image in (pre)adolescents

Federica Cugnata, Clelia Di Serio, Chiara Brombin

Sixth Session, Friday 30 June

10:10 - 10:30

Graphical Models; Two-clusters Solution; Social Network

Digital revolution has dramatically changed not only the way people interact but also the relationship with the self-image. Increased data availability and computational power have significantly improved algorithms for facial feature detection which have been also successfully applied to develop face filter apps enhancing and "beautifying" self-portraits. Potential of these filters in altering facial appearance has raised concerns in parents, educators and health professionals as they promote unrealistic beauty standards increasing discrepancy between real and digital self. Actually taking, sharing and viewing edited selfies may have detrimental effects especially on younger users in a developmental phase where they are already facing significant identity construction processes, possibly giving rise to appearance-related cyberbullying.

To investigate selfie-sharing/editing behavior in (pre)adolescents, their relationship with digital self-image, problematic use of social network and possible internalizing symptoms an online questionnaire, including both validated and ad-hoc realized scales, has been developed. When examining the digital-self image, here the attention is narrowed to the face only, the protagonist of real and virtual interactions, and not to the whole body.

In this setting, graphical models represent an appealing tool to model dependence structure between collected variables. To properly analyze collected data, the procedure should account for the fact that (i) data from psychological questionnaires are measured on discrete/ordinal levels thus violating the normality assumption and (ii) measured behaviors are rarely homogeneous and this heterogeneity should be properly modeled to obtain unbiased results.

To tackle these issues, an approach integrating model-based clustering and graphical models (Fop et al., 2019), has been applied to copula transformed data collected on a sample of 229 middle school (pre)adolescents which took part to the online survey. A two-clusters solution was selected as best based on BIC criterion: the two clusters actually showed different covariance network and different management of online self-image and psychological status. Participants in the cluster displaying a worse management of online self-image and psychological status were mainly female reporting higher use of social networks. To better examine the relationships among variables within each cluster, partial correlation networks were estimated separately for the two clusters and compared using both global and local network statistics and inferential procedure for network comparison.

Although graphical models have been widely used to model psychological phenomena as complex networks, the application to selfie behavior is original. Moreover, identifying clusters within a graphical model framework has important practical implications such as (i) aiding in the development of tailored training programs suited for improving digital wellbeing in younger users and (ii) uncovering new data-driven relationships among constructs thus generating new hypothesis to test in successive studies.

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Sparse Hierarchical Vector Autoregression for Psychopathological Network Estimation from Intensive Longitudinal Data

Spyros Balafas, Grzegorczyk Marco, Federica Cugnata, Sanne Booij, Johanna T. W. Wigman, Ernst C. Wit

Sixth Session, Friday 30 June

10:30 - 10:50

Graphical Vector Autoregressive Model; Network Density; Graphical LASSO

The use of networks as a tool for studying complex systems gained popularity in various scientific disciplines. In the past decade, the "network takeover" reached psychology, and networks were utilized to abstract complex psychological phenomena.

In psychopathology, a network-based framework known as the *network theory of mental illness*, posits that mental disorders emerge as systems of causally interacting psychopathological symptoms. According to this framework, symptoms and other psychological or sociological factors are nodes in a psychopathological network, and the absence of an edge between two nodes corresponds to a conditional independence relationship.

In contrast to other types of networks (e.g., social networks) where the structure is observed, here the dependence structure between the nodes is not known a priori and needs to be estimated from data. Typically, after estimating a psychological network, summary statistics are used to describe its structural properties both at the global and local levels. In the psychological literature, clinical outcomes such as illness severity have been associated with network summary statistics such as network density.

The aim of this study is to test whether i) the network density differs across populations of increasing illness severity, and ii) the local network statistics can be used to identify symptoms that are associated with illness severity. For this purpose, we use intensive longitudinal data from a 90-day diary study called Mapping Individual Routes of Risk and Resilience (MIRORR).

Data consists of 8640 observations within N = 96 individuals, divided over four subgroups representing different early clinical stages ($n_1 = 25$, $n_2 = 27$, $n_3 = 24$, $n_4 = 20$). Participants in the lowest risk group were randomly selected from the general population in the north of the Netherlands based on their score on the Community Assessment of Psychic Experiences (CAPE) test.

Inclusion criteria for the study were: aged between 18 and 35 years, reading and speaking Dutch fluently, being capable of following the research procedures, provide informed consent. Exclusion criteria for participating in the study were: psychotic episodes (current or in the past) according to the Diagnostic and Statistical Manual of Mental Disorders 4 (DSM-4), hearing or visual problems and pregnancy. Participants were excluded from the study when they missed more than 22 measurements in total or missed five or more measurements in a row. Items in the diary assessment covered a wide range of feelings and (subclinical) psychotic experiences, depression, anxiety, mania, obsessive-compulsive behavior, and anger. Participants were required to complete a daily digital questionnaire on psychopathological symptoms, emotions, functioning and stress for 90 consecutive days.

For estimating the network structure for each group of participants, we propose a hierarchical extension of the graphical vector autoregressive (GVAR) model that aims to model the heterogeneity in intensive longitudinal data. The parameters of the proposed hierarchical GVAR model are estimated within a two-step procedure that combines penalized linear mixed models with graphical LASSO (gLASSO).

The estimated networks are then used to calculate global and local network statistics, which are compared across groups using statistical tests. Our results showed that global network statistics such as network density and connectivity do not significantly differ as mental illness becomes more severe. However, we propose the use of local network characteristics such as centrality indices to identify emotions that correlate significantly with increasing illness severity.

How to peel the network: an algorithm for weighted triad census

Roberto Rondinelli, Riccardo Ievoli, Lucio Palazzo

Seventh Session, Friday 30 June

12:30 - 12:50

Network Peeling; Weighted Triadic Census; Network Topology

In Network Analysis, the interaction between three nodes is called a "triad" and represents the minimal group structure that can be observed. According to the presence and the type of the relations between three nodes, sixteen triadic configurations (called the isomorphism classes) are defined and their distribution is denoted as "triad census". This kind of analysis is used in different situations concerning relational data and the conventional approach is well-defined for unweighted networks. As a consequence, the information regarding the weights is not taken into account.

To exploit this information in the triad analysis, we propose a new algorithm denoted as "network peeling" to count the different configurations of triads in weighted networks. The algorithm computes the triad census over the network layers generated at each step. The resulting matrix (with dimensions layers \times isomorphism classes) can be summarized through a set of descriptive measures representing the weighted triad census.

With the aim to highlight the appropriateness of our approach, we consider some real scenarios and a simulation study, comparing weighted and conventional triad censuses.

Network Integration with INet algorithm

Valeria Policastro, Matteo Magnani, Claudia Angelini, Annamaria Carissimo

Seventh Session, Friday 30 June

12:50 - 13:10

Consensus; Weighted Multilayer Networks; INet

Nowadays, network data integration is a demanding problem and still an open challenge, especially when dealing with large datasets. When collecting several data sets and heterogeneous data types on a given phenomenon of interest, the individual analysis of each data set will give only a particular view of such phenomenon. In contrast, integrating all the data will widen and deepen the results giving a more global view of the entire system.

We developed a novel statistical method named INet algorithm, for data integration based on weighted multilayer networks. Under the assumption that the structure underneath the different layers has some similarity that we want to emerge in the integrated network, we generate a "consensus network" through an iterative procedure based on structure comparison, capable of pulling out important information about the phenomenon under study. The procedure tries to preserve common higher-order structures of the original networks in the integrated one, i.e. neighborhood. Once obtained the consensus network, we compared it with the starting networks extracting "specific networks", one for each layer, containing peculiar information of the single data type not present in all the others.

We tested our method on simulated networks to analyze the performance of our algorithm and we analyzed virus and vaccine gene co-expression networks to better understand infectious diseases.

An approach to structural equation modeling in a multiblock framework

Rosaria Romano

Seventh Session, Friday 30 June

12:50 - 13:10

Path Analysis; Graphical Modelling; Multiblock Regression

In many application fields, the variables used to measure a phenomenon are gathered into homogeneous blocks that measure partial aspects of the phenomenon. For example, in sensory analysis, the overall quality of products may depend on the taste and odor variables, etc. In consumer analysis, consumer preferences may depend on physical-chemical and sensory variables. In some contexts, a structure of relations between the different blocks may exist that gives rise to a chain of influences. Within each link, the blocks of predictor variables are called input blocks, while the block of dependent variables is called the output block. If the input blocks do not depend on any other block, then they are defined as exogenous blocks, while those that rely on other input blocks in the same relation are called intermediate blocks. If there is a chain of the relationship between the blocks, we are then dealing with what is often called a mediation model and must interpret both indirect and direct effects among blocks.

Within the scope of multiblock data analysis with a directional path among the blocks, we will present a new approach named SO-PLS path modelling (SO-PLS-PM). The approach splits the estimation into separate sequential orthogonalized PLS regressions (SO-PLS) for each output block. The new method is flexible and graphically oriented and allows for handling multidimensional blocks and diagnosing missing paths. New definitions of total, direct, indirect, and additional effects in terms of explained variances will be proposed, along with new methods for graphical representation.

In this research, some interesting properties of the method will be shown both on simulated and real data. The actual data concerns consumer, sensory and process modelling data. Results will also be compared to those of alternative path modelling methods. Keywords:

References

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Plenary Sessions

Mixture Models for Repeatedly Measured Survey Data

Ralitza Gueorguieva

First Plenary Session, Thursday 29 June

11:00 - 11:40

Survey data items are commonly collected on a Likert scale and may have an additional "don't know" category. It is also typical to have questions that are not applicable to some individuals or to observe floor or ceiling effects on ordinal or interval responses. These situations necessitate the use of mixture models to properly account for the structure of the data. The model formulation also needs to account for correlations among repeated measures within individual. We present a couple of mixture models with random effects for such situations.

In particular, we use logistic sub-models to handle "don't know", inapplicable or floor effects and appropriate generalized linear sub-models for the remaining data. Correlated random effects link the sub-models together. For illustration we use data from tobacco surveys. Maximum likelihood estimation methods are used for model fitting and inference. The software implementation is using PROC NLMIXED in SAS. Simulation studies evaluate bias and efficiency of the parameter estimates.

MCMC or Reservoir computing? A direct sampling approach

Petros Dellaportas

Second Plenary Session, Friday 30 June

11:00 - 11:40

Assume that we would like to estimate the expected value of a function f with respect to a density π by using an importance density function q. We prove that if π and q are close enough under KL divergence, an independent Metropolis sampler estimator that obtains samplers from π with proposal density q, enriched with a variance reduction computational strategy based on control variates, achieves smaller asymptotic variance than the one from importance sampling.

We illustrate our results in challenging option pricing problems that require Monte Carlo estimation. Furthermore, we propose an automatic sampling methodology based on adaptive independent Metropolis that can successfully reduce the asymptotic variance of an importance sampling estimator and we demonstrate its applicability in a Bayesian inference problems.