

# Curriculum Vitae

## Assoc. Prof. Dr. Vilda Purutçuoğlu

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### PERSONAL DETAILS

**Nationality** : Turkish  
**Office** : Middle East Technical University (METU),  
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### EDUCATION

- **Ph.D.**, Statistics, Lancaster University, UK. *Advisor*: Prof. Dr. Ernst Wit. *Thesis Title*: Bayesian methods for gene network analysis. *October 2004-August 2007*.
- **M.Sc.**, Statistics, METU, Turkey. *Advisor*: Prof. Dr. Moti L. Tiku. *Thesis Title*: Unit root problems in time series analysis. *September 2002-January 2004* (By completing a 2 (+1) year programme in 1 and 1/2 years with high honor cumulative).
- **Minor in B.Sc.**, Economics, METU, Turkey. *Degree Title*: Economic theory. *February 2000-June 2002*.
- **B.Sc.**, Statistics, METU, Turkey. *September 1998-June 2002*.

### ACADEMIC POSTS

- *December 2012-current*, **Associate Professor**, Dept. of Statistics, METU.
- *February 2011-current*, **Affiliated Faculty Member**, Dept. of Biomedical Engineering, METU.
- *January 2011-current*, **Affiliated Faculty Member**, Informatics Institute, METU.
- *November 2010-current*, **Affiliated Faculty Member**, Institute of Applied Mathematics, METU.
- *August 2010-December 2012*, **Assistant Professor**, Dept. of Statistics, METU.
- *June 2009-July 2010*, **Instructor**, Dept. of Statistics, METU.
- *August 2007-May 2009*, **Doctor Research Assistant**, Dept. of Statistics, METU.

- *July 2002-July 2007, Research Assistant*, Dept. of Statistics, METU.
- *February 2002-June 2002, Student Assistant*, Dept. of Economics, METU.
- *May 2000 - February 2001, Internships* in the Central Bank (Department of Marketing), State Planning Organization (Department of Economic Modelling and Strategic Researches), Turkish Statistical Institute (Department of National Accounts and Economic Indicators), Ankara.

## RESEARCH INTERESTS

- Bioinformatics, system biology, computational biology, biostatistics, robust inference, high-dimensional inference

## PROFESSIONAL ACTIVITIES

- *September 2017 - current, Faculty Executive Board Member*, Faculty of Arts and Sciences, METU.
- *November 2016 - current, Biomedical Engineering-Biomolecular Track Representative*, Department of Biomedical Engineering, METU.
- *2016 - 2020, Management Committee Member for Turkey* for the COST grant CA15109.
- *2016 - 2020, Awards Fund Committee Member*, International Biometric Society.
- *2017, Programme Committee Member and Referee* of the 3rd International Researchers, Statisticians and Young Statisticians Congress (IRSYC).
- *2015, 2016, Referee* of 2nd and 3rd International Conference on Computational and Experimental Science and Engineering (ICCESEN).
- *February 2015-current, Regular Referee* for the Canadian International Journal of Statistics and Probability.
- *2014-current, Reviewer* for TÜBİTAK (The Scientific and Technological Research Council of Turkey) grant applications.
- *2013-current, Internship Coordinator*, Department of Statistics, METU.
- *2010-current, Advisory Board and Regular Referee* for Journal of Biostatistics - Turkish Clinics.
- *April 2006-current, Referee of Internationale Journals* mostly indexed in SCI: African Journal of Agricultural Research, Annals of Operational Researches, Archives of Current Research International, Asian Journal of Mathematics and Computer Research, Biostatistics, British Journal of Mathematics and Computer Science, British Journal of Economics, Management and Trade, Canadian International Journal of Statistics and Probability, Central European Journal of Operations Research, Cogent Environmental Science, Cogent Food and Agriculture, Cogent Mathematics, Computational Statistics, Communications Faculty of Sciences University of Ankara Series

A1: Mathematics and Statistics, Communications in Nonlinear Science and Numerical Simulation, Journal of Applied Statistics, Journal of Basic and Applied Research International, Journal of Biology and Nature, Journal of Biomedicine and Biotechnology, Journal of Biopharmaceutical Statistics, Journal of Biostatistics-Turkish Clinics, Journal of Clinical Reviews and Opinions, Journal of Computational Science, Journal of Royal Statistical Society-Series C, Journal of Statistical Software, Journal of Statistical Research, Journal of Thermal Engineering, Informatics in Medicine Unlocked, International Journal of Diabetology and Vascular Disease Research, International Journal of Machine Learning and Cybernetics, Mathematical Problems in Engineering, Machine Learning, Microarrays, Optimization, PeerJ, Scientific Research and Essays, Statistical Applications in Genetics and Molecular Biology, Turkish Journal of Biochemistry, Turkish Journal of Biology, Turkish Journal of Mathematics.

- *2013, 2016*, **Stream Organizer** and **Referee** of the 26th and 28th European Conference on Operational Research.
- *2014, 2016, 2017*, **Programme Committee Member** and **Referee** of the 11th, 13th, 14th International Meeting on Computational Intelligence Methods for Bioinformatics and Biostatistics (CIBB).
- *November 2010-2015*, **Editorial Board** and **Regular Referee** for Journal of Clinical Reviews and Opinions.
- *2012, 2013*, **Organization Committee Member** and **Referee** of the 7th and 8th International Symposium on Health Informatics and Bioinformatics (HIBIT).
- *July 2012*, **Invited Researcher**, Department of Mathematics, Brunel University, UK.
- *September 2012-December 2013*, **Minor Programme Coordinator**, Department of Statistics, METU.
- *2009-2012*, **Seminar Coordinator**, Department of Statistics, METU.
- *2009, 2010*, **Invited Researcher**, Institute of Mathematics and Computing Science, Groningen University, The Netherlands.
- *February 2010-July 2010*, **Organization Committee Member** and **Referee** of the 7th National Symposium of Statistical Days.

## SCHOLARSHIPS AND AWARDS

- *2014-2016* : **Academic Performance Award**, Faculty of Art and Science, METU.
- *2015*: **Finalist of Sabri Ülker Science Award**, Sabri Ülker Food Research Foundation.
- *2015*: **TEÇEP Translation Book Honorable Mention Award**, Translated Book Edited by Prof. Dr. Zeki Kaya and co-authored with 22 academicians.
- *2013-2015* : **Academic Performance Award**, Faculty of Art and Science, METU.
- *2012-2014* : **Academic Performance Award**, Faculty of Art and Science, METU.

- *June 2012* : **Visitor Fellowship**, Department of Mathematics, Brunel University, UK.
- *2009-2011* : **Academic Performance Award**, Faculty of Art and Science, METU.
- *2009, 2010* : **Visitor Fellowship**, Institute of Mathematics and Computing Science, Groningen University, The Netherlands.
- *2005-2008* : **Fellowships from Research Institutions in UK**: RSS (Royal Statistical Society), ESRC (Economic and Social Research Council), EPSRC (Engineering and Physical Sciences Research Council), BBSRC (Biotechnology and Biological Sciences Research Council), CRISM (Centre for Research in Statistical Methodology, Warwick University) Grants; Faculty Scholarships of Lancaster University (The fellowships cover all expenses for the participation of different conferences and training courses).
- *2005-2007* : **EPSRC Project Grant** (covers all expenses during Ph.D. including home tuition fees and subsistence), UK.
- *2004-2007* : **Overseas Ph.D. Fellowship**, Department of Mathematics and Statistics, Lancaster University, UK (The fellowship covers the difference between home and overseas tuition fees during Ph.D. in UK).
- *2004-2005* : **Ph.D. Fellowship**, Turkish Higher Education Council (The fellowship covers the home tuition fees and subsistence).

## PROJECTS

### A. ONGOING PROJECTS

- **Principal investigator**, *January 2017-August 2018*, Scientific Research Project (BAP1), METU. *Project title*: Inference of the Gaussian graphical model and lasso regression via the modified maximum likelihood method. *Project no*: BAP-01-09-2017-002.
- **Principal investigator**, *January 2017-June 2018*, Scientific Research Project (DAP1), METU. *Project title*: Collection of genetic information, analysis and modeling of the cervical cancer with risk factors. *Project no*: BAP-08-11-2017-035.
- **Co-applicant and management committee member for Turkey**, *January 2016-January 2020*, European Community 7th Framework Programme (COST Programme). *Project title*: European network for statistical network science. *Project no*: CA15109.

### B. COMPLETED PROJECTS

- **Principal investigator**, *January 2016-June 2017*, Scientific Research Project (BAP1), METU. *Project title*: Alternative approaches, inference and copulas in deterministic modellings of complex biological systems. *Project no*: BAP-01-09-2016-002.
- **Principal investigator**, *April 2015-April 2017*, Scientific and Technological Research Council (TÜBİTAK - 1001 Programme). *Project title*: Application of copulas in inference of biological networks via Gaussian graphical models and parameter estimation. *Project no*: 114E636.

- **Principal investigator**, *January 2014-January 2017*, METU Research and Development Project (AGEP Programme). *Project title*: Simulation of biochemical systems via stochastic simulation algorithms and extensions of these methods by including impulses, *Project no*: BAP-08-11-2014-007.
- **Principal investigator**, *April 2013-April 2015*, Scientific and Technological Research Council (TÜBİTAK-3501 Programme). *Project title*: Stochastic inference of the model parameters for the biochemical systems via the particle filtering method. *Project no*: 112T772.
- **Researcher**, *December 2012-July 2013*, Scientific and Technological Research Council (TÜBİTAK-1002 Programme). *Project title*: The classification of histologic and pathologic subtypes of breast cancer with meta-analysis based in miRNA signatures. *Project no*: 112T679.
- **Co-applicant for METU**, *September 2010 - March 2015*, European Community 7th Framework Programme, with Dr. Tolga Can (Dept. of Computer Engineering, METU) and Prof. Gerhard Wilhelm Weber (Institute of Applied Math, METU). *Project title*: PATHOSYS - New Algorithms for Host Pathogen Systems Biology. *Project no*: 260429.
- **Co-applicant**, *1-15 July 2010*, Scientific and Technological Research Council (TÜBİTAK-2229 Programme), with Dr. Barış Bayram and Prof. Mevlüde Gülbin Dural (Dept. of Electric and Electronic Engineering, METU). *Project title*: Teaching promising high-school students for the “Physics Olympiad”.
- **Researcher**, *September 2009-September 2011*, Scientific Research Project (BAP), METU. *Project title*: Identification of Turkish climate zones and development of precipitation forecast models by data mining methods. *Project no*: BAP-2008-01-09-02.
- **Researcher**, *September 2005-August 2007*, Engineering and Physical Sciences Research Council, UK (EPSRC). *Project title*: Stochastic modelling and statistical inference of gene regulatory pathways: integrating multiple sources of data. *Project no*: EP/C010620/1.

## PUBLICATIONS IN JOURNALS

1. Altıntan, D., **Purutçuoğlu, V.** and Uğur, Ö. (2018) “Impulsive expressions in stochastic simulation algorithms”, *International Journal of Computational Methods*, doi: 10.1142/S021987621750075X, 15 (1), 1750075.1-16.
2. Ayyıldız, E., **Purutçuoğlu, V.** and Weber, G.W. (2017) “Loop-based conic multivariate adaptive regression splines is a novel method for advanced construction of complex biological networks”, *European Journal of Operational Research*, doi: 10.1016/j.ejor.2017.12.011, 1-10.
3. Varol, D., **Purutçuoğlu, V.** and Yılmaz, R. (2017) “Whole genome analysis of the heat stress response in commercial baker’s yeast (*Saccharomyces cerevisiae*) with comparative statistical approaches”, *Gene and Genomics*, doi: 10.1007/s13258-017-0616-6, 1-14.
4. Dokuzoğlu, D. and **Purutçuoğlu, V.** (2017) “Comprehensive analyses of Gaussian graphical model under different biological networks”, *Acta Physica Polonica, Series A*, 132, 1106-1111.

5. **Purutçuoğlu, V.** and Farnoudkia, H. (2017) “Gibbs sampling in inference of copula Gaussian graphical model adapted to biological networks”, *Acta Physica Polonica, Series A*, 132, 1112-1117.
6. **Purutçuoğlu, V.** and Ayyıldız, E. (2017) “Mathematical modeling of gene networks (in Turkish)”, *Journal of Biostatistics-Turkish Clinics*, doi: 10.5336/biostatic.2017-55474, 143-155.
7. Akal, T., **Purutçuoğlu, V.** and Weber, G.-W. (2017) “Robust background normalization method for one-channel microarrays”, *Turkish Journal of Biochemistry*, doi: 10.1515/tjb-2016-0231, 42 (2), 111-121.
8. **Purutçuoğlu, V.**, Ayyıldız, E. and Wit, E. (2017) “Comparison of two inference approaches in Gaussian graphical models”, *Turkish Journal of Biochemistry*, 42 (2), 203-211.
9. **Purutçuoğlu, V.**, Ağraz, M. and Wit, E. (2017) “Bernstein approximations in glasso-based estimation of biological networks”, *The Canadian Journal of Statistics*, doi: 10.1002/cjs11309, 45 (1), 62-76.
10. Tuncer, G. and **Purutçuoğlu, V.** (2017) “Application of impulsive deterministic simulation of biochemical networks via simulation tools”, *Proceedings of the Jangjeon Mathematical Society (PJMS)*, 1, 105-119.
11. Ayyıldız, E., Ağraz, M. and **Purutçuoğlu, V.** (2017) “MARS as the alternative approach of Gaussian graphical model for biochemical networks”, *Journal of Applied Statistics*, doi: 10.1080/02664763.2016.1266465, 44 (16), 2858-2876.
12. Ağraz, M. and **Purutçuoğlu, V.** (2016) “Different types of Bernstein operators in inference of Gaussian graphical model”, *Cogent Mathematics*, 3, 1154706.1-11.
13. Tuncer, G. and **Purutçuoğlu, V.** (2015) “Comparative assessment of simulation tools for biochemical networks; *American Review of Mathematics and Statistics*, doi: 10.15640/arms.v3n2a9, 3 (2), 69-82.
14. İyigün, C., Türkeş, M., Batmaz, İ., Yozgatlıgil, C., **Purutçuoğlu, V.**, Koç, E.K., Öztürk, M. (2013) “Clustering current climate regions of Turkey by using a multivariate statistical method”, *Theoretical and Applied Climatology*, doi: 10.1007/s00704-012-0823-7, 1-15.
15. **Purutçuoğlu, V.** and Wit, E. (2012) “Estimating network kinetics of the MAPK/ERK pathway using biochemical data”, *Mathematical Problems in Engineering*, doi:10.1155/2012/752631, 1-34.
16. Ayyıldız, E., **Purutçuoğlu, V.**, and Wit, E. (2012) “A short note on resolving singularity problems in covariance matrices”, *International Journal of Statistics and Probability*, doi:10.5539/ijsp.v1n2p113, 1 (2), 113-118.
17. **Purutçuoğlu, V.** (2012) “Inference of the stochastic MAPK pathway by modified diffusion bridge method”, *Central European Journal of Operations Research*, doi: 10.1007/s10100-012-0237-8, 1-15.
18. **Purutçuoğlu, V.** (2012) “Robust gene expression index”, *Mathematical Problems in Engineering*, doi: 10.1155/2011/182758, 1-12.

19. **Purutçuoğlu, V.** and Karagülle, S. (2011) “Stukel’s extended logistic regression analysis with R”, *Journal of Biostatistics-Turkish Clinics*, 3 (1), 49-56.
20. Çuhacı, G. and **Purutçuoğlu, V.** (2011) “Review of the six-sigma methodology and its case studies”, *Selçuk Journal of Applied Mathematics*, 12 (1), 95-108.
21. Kartal, E., Fahmi, F. M., İyigün C., Türkeş, M., Yozgatlıgil, C., **Purutçuoğlu, V.**, Batmaz, İ., and Köksal, G. (2011) “Identifying climate zones of Turkey by hierarchical clustering method (in Turkish)”, *Journal of Statistical Research*, 8 (1), 13-25.
22. **Purutçuoğlu, V.** (2010) “An overview to stochastic simulation algorithms for biochemical systems (in Turkish)”, *Journal of Statistical Research*, 7 (1), 70-82.
23. **Purutçuoğlu, V.** and Wit, E. (2008) “Bayesian inference for the MAPK/ERK pathway by considering the dependency of the kinetic parameters”, *Bayesian Analysis*, doi: 10.1214/08-BA332, 3 (4), 851-886.
24. **Purutçuoğlu, V.** and Wit, E. (2008) “Bayesian inference of the complex MAPK pathway under structural dependency”, *Journal of Statistical Research*, 6 (1), 1-17.
25. **Purutçuoğlu, V.** and Wit, E. (2007) “FGX: a frequentist gene expression index for Affymetrix arrays”, *Biostatistics*, doi: 10.1093/biostatistics/kx1020, 8 (2), 433-437.
26. **Purutçuoğlu, V.** and Wit, E. (2006) “Exact and Approximate Stochastic Simulations of the MAPK Pathway and Comparisons of Simulations’ Results”, *Journal of Integrative Bioinformatics*, doi:10.2390/biecoll-jib-2006-38, 3, 1-13.
27. **Purutçuoğlu, V.** and Wit, E. (2005). Discussion on the paper by Copas and Eguchi, *Journal of the Royal Statistical Society, Series B*, doi: 1369-7412/05/67459, 67, Part 4, 507-508.

## BOOKS

- Wit, E., Vinciotti, V., and **Purutçuoğlu, V.** (January 2018) “Statistics for Biological Networks: How to Infer Networks from Data”, Chapman and Hall/CRC. ISBN: 978-439841470.
- **Purutçuoğlu, V.** and Ayyıldız, E. (December, 2014) “Statistics in the Field of Bioinformatics (Biyoinformatik Alanında İstatistik-in Turkish)”, Nobel Publisher. ISBN: 978-605-320-008-6.

## CHAPTERS IN BOOKS

1. Seçilmiş, D., Ağraz, M. and **Purutçuoğlu, V.** (2018) “Two novel non-parametric models in the construction of biological networks”. Chapter in: *Information Complexity and Statistical Modeling in High Dimensions with Applications*. Editors: O. Arslan. Springer (Accepted).
2. Ayyıldız, E. and **Purutçuoğlu, V.** (2017) “Generating various types of graphical models via MARS”. Chapter in: *Information Complexity and Statistical Modeling in High Dimensions with Applications*. Editors: O. Arslan. Springer (In print).

3. Seçilmiş, D. and **Purutçuoğlu, V.** (2017) “Modeling of biochemical networks via classification and regression tree methods”. Chapter in: *Mathematical Methods in Engineering*. Editors: K. Taş, D. Baleanu and J.A. Mashado. *Springer* (In print).
4. Ağraz, M. and **Purutçuoğlu, V.** (2017) “Empirical copula in the detection of batch effects”. Chapter in: *Mathematical Methods in Engineering*. Editors: K. Taş, D. Baleanu and J.A. Mashado. *Springer* (In print).
5. Tuncer, G. and **Purutçuoğlu, V.** (2017) “Major simulation tools for biochemical networks”. Chapter in: *Modelling, Dynamics, Optimization and Bioeconomics III*. Editors: A. Pinto, Z. Zilberman. *Springer* (In print).
6. **Purutçuoğlu, V.** (2017) “Stochastic modelling of biochemical networks and inference of model parameters”. Chapter in: *Modelling, Dynamics, Optimization and Bioeconomics III*. Editors: A. Pinto, Z. Zilberman. *Springer* (In print).
7. Ağraz, M. and **Purutçuoğlu, V.** (2016) “Transformations of data in deterministic modelling of biological networks”, 343-356. Chapter in: *Intelligent Mathematics II: Applied Mathematics and Approximation Theory*. Editors: O. Duman. *Springer*.
8. **Purutçuoğlu, V.** (2014) “Biological networks uncover evolution, disease, and gene functions (in Turkish)”, 291-314. Chapter in: *Bioinformatics for Biologist* (Translated from the book of P. Pevzner and R. Shamir entitled by the same name). Editor: Z. Kaya, J. *Nobel Publisher*.
9. Defterli, Ö., **Purutçuoğlu, V.**, and Weber, G. W. (2014) “Advanced mathematical and statistical tools in the dynamic modelling and simulation of gene-environment networks”, 235-257. Chapter in: *Modeling, Optimization, Dynamics and Bioeconomy*. Editor: D. Zilberman and A. Pinto. *Springer*.
10. **Purutçuoğlu, V.**, Kayış, E., and Weber, G. W. (2011) “Survey of background normalizations for Affymetrix arrays and a case study”, 199-219. Chapter in: *Advances in Intelligent Modelling and Simulation: Simulation Tools and Applications*. Editor: A. Byrski, Z. Oplatkova, M. Carvalho, and M. Kisiel-Dorohinicki. *Springer*.
11. Fahmi, M. F., Kartal, E., İyigün, C., Türkeş, M., Yozgatlıgil, C., **Purutçuoğlu, V.**, Batmaz, İ., and Köksal, G. (2011) “Determining the climate zones of Turkey by center-based clustering methods”, 171-177. Chapter in: *Nonlinear Dynamics of Complex Systems: Applications in Physical, Biological and Financial Systems*. Editor: J. A. T. Machado, D. Baleanu, and A. Luo. *Springer*.
12. Wit, E., **Purutçuoğlu, V.**, O’Donovan, L., and Zhu, X. (2006) “Gaining weights . . . and feeling good about it”, 31-38. Chapter in: *Microarray Technology and Cancer Gene Profiling*. Editor: S. Mocellin. *Springer*.

## SUBMITTED AND ONGOING MANUSCRIPTS

1. Ayyıldız, E. and **Purutçuoğlu, V.** “Modeling of various biological networks via LCMARS”.
2. Farnoudkia, H. and **Purutçuoğlu, V.** “Modelling of biological networks via copula Gaussian graphical model and bayesian inference of model parameters”.



3. **Purutçuoğlu, V.** and Altıntan, D. “Exact stochastic simulation algorithms under different scenarios of biological systems”.
4. **Purutçuoğlu, V.** and Wit, E. “Detection of the degree distribution in biological networks”.
5. Ağraz, M. and **Purutçuoğlu, V.** “Extended lasso-type MARS (LMARS) model in the description of biological network”.
6. Ağraz, M. and **Purutçuoğlu, V.** “Inference of the steady-state activation of biological networks via new frequentist approach”.
7. Akkuş, E.C. and **Purutçuoğlu, V.** “A new non-parametric outlier detection method”.
8. Bahçivancı, B. and **Purutçuoğlu, V.** “Mathematical modeling of gynecological caners via quasi target proteins”.

## SELECTED PROCEEDINGS

### A. FULL PAPERS

1. Varol, D., Akal, T. and **Purutçuoğlu, V.** (2017) “Is the log-normal density appropriate for one-channel microarray data?”, *Proceeding of the International Conference on Advances in Science (ICAS 2017)*, İstanbul, Turkey.
2. Seçilmiş, D. and **Purutçuoğlu, V.** (2017) “Nonparametric versus parametric models in inference of protein-protein interaction networks”, *Proceeding of the International Conference on Advances in Science and Arts İstanbul (ICASA 2017)*, İstanbul, Turkey.
3. Ağraz, M. and **Purutçuoğlu, V.** (2017) “Inference of the Gaussian graphical model via the modified maximum likelihood approach”, *Proceeding of İstanbul International Conference on Progress in Applied Science İstanbul (ICPAS 2017)*, İstanbul, Turkey.
4. **Purutçuoğlu, V.** and Farnoudkia, H. (2017) “Bayesian inference of biological networks whose components are linearly dependent”, *Proceeding of İstanbul International Conference on Progress in Applied Science (ICPAS 2017)*, İstanbul, Turkey.
5. Yazıcı, M., **Purutçuoğlu, V.** and Altıntan, D. (2016) “Application of stochastic simulations of biological networks under different impulsive scenarios”, *Proceeding of the 3rd International Conference on Data Mining, Internet Computing, and Big Data (BigData2016)*, Konya, Turkey.
6. **Purutçuoğlu, V.** and Odunsi, O. (2016) “Degree distribution of real biological networks”, *Proceeding of the 3rd International Conference on Data Mining, Internet Computing, and Big Data (BigData2016)*, Konya, Turkey.
7. Yazıcı, C., **Purutçuoğlu, V.**, Yozgatlıgil, C., and Batmaz, İ. (2013) “Detection of unreliable measurements in long-term time-series via data mining techniques: case in Turkish climate data”, *Proceeding of the Global Reach of Industrial Engineering*, İstanbul, Turkey.
8. **Purutçuoğlu, V.** and Kayış, E. (2012) “Comparing clustering techniques for real microarray data”, *Proceeding of IEEE/ACM International Conference on Social Networks Analysis and Mining*, İstanbul, Turkey.

9. Fahmi, F., Kartal, E., İyigün, C., Yozgatlıgil, C., **Purutçuoğlu, V.**, Çınar, İ., Aslan, S., Öztürk, M. Z., Türkeş, M., and Batmaz, İ. (2011) “Studying the effect of climate change on Turkey by clustering techniques (in Turkish)”, *Proceeding of the National Geographical Congress*, İstanbul, Turkey.
10. **Purutçuoğlu, V.**, Erdem, T., and Weber, G. W. (2011) “Inference of the JAK-STAT gene network via graphical models”, *Proceeding of the 23rd International Conference on Systems Research, Informatics and Cybernetics*, Baden, Germany.
11. Yozgatlıgil, C., Yazıcı, C., **Purutçuoğlu, V.**, and Batmaz, İ. (2011) “Validity of homogeneity tests for meteorological time series data: a simulation study”, *Proceeding of the 58th ISI World Statistics Congress*, Dublin, Ireland.
12. Quek, E. M. P., **Purutçuoğlu, V.**, Sambanthamurthi, R., and Weber, G. W. (2011) “Modelling lipid biosynthesis pathways of oil palms by boolean and graphical approaches”, *Proceeding of the 6th International Symposium on Health, Informatics and Bioinformatics (HIBIT 2011)*, İzmir, Turkey.
13. Konak, D. and **Purutçuoğlu, V.** (2010) “Detection of calendar effects on Esenboğa, Atatürk, and London City International Airports by dummy variable method (in Turkish)”, *Proceeding of the 7th National Symposium of Statistical Days*.
14. **Purutçuoğlu, V.** (2010) “Stochastic simulation of large biochemical systems by approximate Gillespie algorithm”, *Proceeding of the 5th International Symposium on Health, Informatics and Bioinformatics*, Antalya, Turkey.
15. **Purutçuoğlu, V.** and Wit, E. “Inclusion of convoluted measurements in Bayesian inference of the MAPK/ERK pathway via multivariate diffusion model”, *Proceeding of the 3rd International Symposium on Health, Informatics and Bioinformatics*, İstanbul, Turkey, 2008.
16. **Purutçuoğlu, V.** and Wit, E. (2006) “Stochastic modelling of the MAPK signalling pathway”, *Proceeding of the 21st International Workshop on Statistical Modelling*, Galway, Ireland.

## B. PUBLISHED ABSTRACTS

1. Akkuş, E.C., **Purutçuoğlu, V.** and Ağraz, M. (2017) “Detection of outliers using Fourier transform”, *Proceeding of the 10th International Statistics Congress (ISC2017)*, Ankara, Turkey.
2. Bahçivancı, B. and **Purutçuoğlu, V.** (2017) “Modeling of gynecological cancers by using all risk factors”, *Proceeding of the International Conference on Engineering Technologies (ICENTE 2017)*, Konya, Turkey.
3. **Purutçuoğlu, V.** (2017) “Comparison of various biological network constructions”, *Proceeding of the 1st International Conference on Computational and Statistical Methods in Applied Sciences (COSTAS 2017)*, Samsun, Turkey.
4. **Purutçuoğlu, V.** and Bahçivancı (2017) “Mathematical modeling of gynecological cancers via quasi target proteins”, *Proceeding of the 4rd International Conference on Computational and Experimental Science and Engineering (ICCESEN-2017)*, Antalya, Turkey.

5. **Purutçuoğlu, V.** (2017) “Detection of the best model selection criterion in Gaussian graphical model”, *Proceeding of the 4rd International Conference on Computational and Experimental Science and Engineering (ICCESEN-2017)*, Antalya, Turkey.
6. Karakelle, S., **Purutçuoğlu, V.** and Ürün (2017) “Cancer modeling via biologically validated genes”, *Proceeding of the International Conference on Mathematics and Engineering (ICOME-2017)*, İstanbul, Turkey.
7. **Purutçuoğlu, V.** and Akal, T. (2017) “Accuracy assessment of recent gene expression indices via multi-RGX for one-channel microarrays”, *Proceeding of the International Conference on Mathematics and Engineering (ICOME-2017)*, İstanbul, Turkey.
8. Ayyıldız, E. and **Purutçuoğlu, V.** (2017) “Modeling of various biological networks via LCMARS”, *Proceeding of the International Workshop on Mathematical Methods in Engineering (MME 2017)*, Ankara, Turkey.
9. **Purutçuoğlu, V.** and Ağraz M. (2017) “Steady-state modeling of the biological network via long-tailed symmetric distribution”, *Proceeding of International Conference on Recent Advances in Pure and Applied Mathematics (ICRAPAM 2017)*, Kuşadası, Turkey.
10. **Purutçuoğlu, V.** and Farnoudkia, H. (2016) “Bayesian inference of deterministic MAPK-ERK pathway via reversible jumps Monte Carlo method”, *Proceeding of the 8th International Conference Inverse Problems: Modeling and Simulation (IPMS-2016)*, Fethiye, Turkey.
11. Ağraz, M. and **Purutçuoğlu, V.** (2016) “Deterministic modelling of linear and nonlinear interactions in biological systems”, *Proceeding of the International Conference on Information Complexity and Statistical Modeling in High Dimensions with Applications (IC-SMHD)*, Kapadokya, Turkey.
12. Ağraz, M. and **Purutçuoğlu, V.** (2016) “A non-parametric model in the construction of biological networks”, *Proceeding of the 2nd Researchers-Statisticians and Young Statisticians Congress (IRSYSC)*, Ankara, Turkey.
13. **Purutçuoğlu, V.** and Farnoudkia, H. (2016) “Modelling of biological networks via copula Gaussian graphical model and bayesian inference of model parameters”, *Proceeding of the 2nd Researchers-Statisticians and Young Statisticians Congress (IRSYSC)*, Ankara, Turkey.
14. **Purutçuoğlu, V.** and Altıntan, D. (2016) “Exact stochastic simulation algorithms under different scenarios of biological systems”, *Proceeding of the 2nd Researchers-Statisticians and Young Statisticians Congress (IRSYSC)*, Ankara, Turkey.
15. Tuncer, G. and **Purutçuoğlu, V.** (2015) “Bifurcation Analyses in Biochemical Networks via Simulation Tools”, *Proceeding of the 2nd International Conference on Computational and Experimental Science and Engineering*, Antalya, Turkey.
16. Seçilmiş, D. and **Purutçuoğlu, V.** (2015) “Inference of Biological Networks via Random Forest Algorithm”, *Proceeding of the 7th Bioengineering Congress*, İzmir, Turkey.
17. Altıntan, D., **Purutçuoğlu, V.** and Uğur, Ö. (2015) “Functional impulses in exact stochastic simulation”, *Proceeding of the International Conference on Pure and Applied Mathematics*, Van, Turkey.

18. **Purutçuoğlu, V.** (2015) “ Applications of copulas in deterministic models of biochemical networks”, *Proceeding of the International Conference on Recent Advances in Pure and Applied Mathematics*, İstanbul, Turkey.
19. Dokuzoğlu, D. and **Purutçuoğlu, V.** (2015) “ Simulation study for the application of copulas in the Gaussian graphical model”, *Proceeding of the 8th Eastern Mediterranean Region International Biometric Society*, Kapadokya, Turkey.
20. **Purutçuoğlu, V.** and Şeker, T. (2015) “ Estimation of reaction rates for the Mapk-Erk pathway via western-blotting data”, *Proceeding of the 41st Annual Meeting of the European Society for Blood and Marrow Transplantation*, İstanbul, Turkey.
21. **Purutçuoğlu, V.** (2014) “ Stochastic inference of the complex MAPK/ERK pathway when the system has impulsive changes”, *Proceeding of the 17th International Symposium on the Biology of Actinomycetes*, Kuşadası, Turkey.
22. Ayyıldız, E. and **Purutçuoğlu, V.** (2013) “ Inference of the biological systems via  $L_1$ -penalized lasso regression”, *Proceeding of the 29th Meeting of Statisticians*, Budapest, Hungary.
23. Varol, D. and **Purutçuoğlu, V.** (2013) “ Comparative analysis of a one-channel microarray dataset by different methods”, *Proceeding of the 29th Meeting of Statisticians*, Budapest, Hungary.
24. Ağraz, M., Kılıç, B., and **Purutçuoğlu, V.** (2013) “ Deterministic modelling of gene network via parametric and non-parametric approaches”, *Proceeding of the 29th Meeting of Statisticians*, Budapest, Hungary.
25. Altıntan, D., **Purutçuoğlu, V.**, Uğur, Ö. (2013) “ Impulsive expression in chemical master equation and stochastic simulation algorithms”, *Proceeding of the 26th European Conference on Operational Research*, Rome, Italy.
26. Aksoy, B., **Purutçuoğlu, V.**, Batmaz, İ, and Yozgatlıgil, C. (2013) “ Modelling the extreme precipitation data: case study from Turkey”, *Proceeding of the 26th European Conference on Operational Research*, Rome, Italy.
27. **Purutçuoğlu, V.** and Kayış, E. (2012) “ Comparing clustering techniques for real microarray data”, *Proceeding of IEEE/ACM International Conference on Social Networks Analysis and Mining*, İstanbul, Turkey.
28. Akal, T., Fahmi, F., Kartal Koç, E., **Purutçuoğlu, V.**, İyigün, C., Yozgatlıgil, C., and Batmaz., İ. (2012) “ Detection of seasonal changes in climate data by comparative analysis: a case study for Turkey”, *Proceeding of the 25th European Conference on Operational Research*, Vilnius, Lithuania.
29. Aykan, F., Kartal Koç, E., Yozgatlıgil, C., İyigün, C., **Purutçuoğlu, V.**, and Batmaz., İ. (2012) “ Developing precipitation models for continental Central Anatolia, Turkey ”, *Proceeding of the 25th European Conference on Operational Research*, Vilnius, Lithuania.
30. **Purutçuoğlu, V.** and Akal,T. (2012) “ multi-RGX: a novel background normalization method for oligonucleotides”, *Proceeding of the 8th World Congress in Probability and Statistics*, İstanbul, Turkey.

31. **Purutçuoğlu, V.**, Defterli, Ö., Fügenschu, A., and Wilhelm-Weber, G. (2012) “Reconstruction of a complex biochemical system and forecasting its behaviour”, *Proceeding of the 25th Conference of European Chapter on Combinatorial Optimization*, Antalya, Turkey.
32. Akay, T., Kartal Koç, E., **Purutçuoğlu, V.**, İyigün, C., Yozgatlıgil, C., and Batmaz, İ. (2012) “Investigating the seasonal patterns of Continental Central Anatolia by clustering”, *Proceeding of the 25th Conference of European Chapter on Combinatorial Optimization*, Antalya, Turkey.
33. Yazıcı, C., **Purutçuoğlu, V.**, Yozgatlıgil, C., Batmaz, İ, and Bayramoğlu, K. (2011) “Homogeneity analysis for dependent climate data”, *Proceeding of the 7th International Statistical Congress*, Antalya, Turkey.
34. Yozgatlıgil, C., **Purutçuoğlu, V.**, Yazıcı, C., and Batmaz, İ. (2010) “Plausibility of the standard normal homogeneity test on Turkish precipitation data (in Turkish)”, *Proceeding of the 7th National Symposium of Statistical Days*.
35. **Purutçuoğlu, V.** and Tiku, M. L. (2009) “Testing unit root and comparison of estimates”, *Proceeding of the 14th International Congress on Computational and Applied Mathematics*, Antalya, Turkey.
36. **Purutçuoğlu, V.** and Wit, E. (2009) “Rao-blackwellized estimates for the multivariate Bayesian inference”, *Proceeding of the International Conference of Mathematical Sciences*, İstanbul, Turkey.
37. **Purutçuoğlu, V.** and Wit, E. (2008) “Variational approximation in inference of the kinetic parameters of the MAPK/ERK pathway”, *Proceeding of CRISM Workshop: Bayesian Inference for high-dimensional data*, Warwick, UK.
38. **Purutçuoğlu, V.** and Wit, E. (2008) “An approximation algorithm based on leap condition for stochastic simulation of biomedical systems”, *Proceeding of the 4th International Conference “Inverse problems: Modelling and Simulation”*, Fethiye, Turkey.
39. **Purutçuoğlu, V.** and Wit, E. (2007) “A new method in approximate stochastic simulations and application in real biochemical systems”, *Proceeding of the 17th International Workshop on Mathematical and Statistical Aspects of Molecular Biology*, Manchester, UK.

## TECHNICAL REPORTS AND OTHERS

- Yazıcı, C., **Purutçuoğlu, V.**, Yozgatlıgil, C., Bayramoğlu, K., İyigün, C., and Batmaz, İ. “Homogeneity analysis of Turkish climate data”, 2012, No: METU-STAT-Technical Report-2012-001.
- Asar, Ö., Kartal, E., Aslan, S., Öztürk, M. Z., Yozgatlıgil, C., Çınar, İ., Batmaz, İ., **Purutçuoğlu, V.**, İyigün, C., Fahmi, F., Köksal, G., Türkeş, M., and Tath, H. “Handling and analysis of Turkish precipitation data for the period 1950-2006 using descriptive data mining techniques (in Turkish)”, 2011, No: METU-STAT-Technical Report-2011-1.
- Wit, E., Vinciotti, V., **Purutçuoğlu, V.** “Statistics for biological networks”, 2010, 25th International Biometric Conference, Short Course’s Notes, 1-97.

## TALKS

### Conferences Talks

1. “Modeling of gynecological cancers by using all risk factors, *International Conference on Engineering Technologies (ICENTE 2017)*, Konya, Turkey, 7-9 December, 2017.
2. “Comparison of various biological network constructions, *1st International Conference on Computational and Statistical Methods in Applied Sciences (COSTAS 2017)*, Samsun, Turkey, 9-11 November, 2017.
3. “Mathematical modeling of gynecological cancers via quasi target proteins”, *4rd International Conference on Computational and Experimental Science and Engineering (ICCESEN-2017)*, Antalya, Turkey, 4-8 October, 2017.
4. “Nonparametric versus parametric models in inference of protein-protein interaction networks, *International Conference on Advances in Science and Arts İstanbul (ICASA 2017)*, İstanbul, Turkey, 29-31 March, 2017.
5. “Accuracy assessment of recent gene expression indices via multi-RGX for one-channel microarrays”, *International Conference on Mathematics and Engineering (ICOME-2017)*, İstanbul, Turkey, 10-12 May, 2017.
6. “Inference of the Gaussian graphical model via the modified maximum likelihood approach, *İstanbul International Conference on Progress in Applied Science İstanbul (ICPAS 2017)*, İstanbul, Turkey, 4-6 January, 2017.
7. “Bayesian inference of biological networks whose components are linearly dependent”, *İstanbul International Conference on Progress in Applied Science (ICPAS 2017)*, İstanbul, Turkey, 4-6 January, 2017.
8. “Construction of ovarian cancer pathway via different mathematical models, *3rd International Researchers, Statisticians and Young Statisticians Congress (IRSYC 2017)*, Konya, Turkey, 24-26 April, 2017.
9. “Modelling of biochemical networks via classification and regression tree method, *International Workshop on Mathematical Methods in Engineering (MME 2017)*, Ankara, Turkey, 27-29 April, 2017.
10. “Steady-state modeling of the biological network via long-tailed symmetric distributio, *International Conference on Recent Advances in Pure and Applied Mathematics (ICRAPAM 2017)*, Kuşadası, Turkey, 11-15 May, 2017.
11. “Bayesian inference of biological networks whose components are linearly dependent”, *stanbul International Conference on Progress in Applied Science (ICPAS 2017)*, İstanbul, Turkey, 4-6 January, 2017.
12. “Acceleration of reversible jump Markov chain Monte Carlo method within Gaussian copula graphical model”, *Proceeding of the 3rd International Conference on Computational and Experimental Science and Engineering (ICCESEN-2016)*, Antalya, Turkey, 19-24 October, 2016.

13. “Comprehensive analyses of Gaussian graphical model under different biological networks”, *Proceeding of the 3rd International Conference on Computational and Experimental Science and Engineering (ICCESEN-2016)*, Antalya, Turkey, 19-24 October, 2016.
14. “Bayesian inference of deterministic MAPK-ERK pathway via reversible jumps Monte Carlo method”, *8th International Conference Inverse Problems: Modeling and Simulation (IPMS-2016)*, Fethiye, Turkey, 23-28 May, 2016.
15. “Copulas in random forest algorithm for modelling biochemical networks”, *International Conference on Information Complexity and Statistical Modeling in High Dimensions with Applications (IC-SMHD)*, Kapadokya, Turkey, 18-21 May, 2016.
16. “Modelling of biological networks via copula Gaussian graphical model and bayesian inference of model parameters”, *2nd Researchers-Statisticians and Young Statisticians Congress (IRSYSC)*, Ankara, Turkey, 4-8 May, 2016.
17. “Reconstruction of biological networks via MARS model based on various precision matrices”, *7th Bioengineering Congress (BEC 2015)*, İzmir, Turkey, 19-21 November, 2015.
18. “Comparison of performances for stochastic simulation algorithms of biochemical networks with/without impulses”, *2nd International Conference on Computational and Experimental Science and Engineering (ICCESEN 2015)*, Antalya, Turkey, 14-19 October, 2015.
19. “Applications of copulas in deterministic models of biochemical networks”, *International Conference on Recent Advances in Pure and Applied Mathematics (ICRAPAM 2015)*, İstanbul, Turkey, 3-6 June, 2015.
20. “Application of impulsive deterministic simulation of biochemical networks via simulation tools”, *28th International Conference of The Jangjeon Mathematical Society (ICJMS 2015)*, Antalya, Turkey, 15-19 May, 2015.
21. “Comparative analysis of simulation tools in biochemical networks”, *28th International Conference of The Jangjeon Mathematical Society (ICJMS 2015)*, Antalya, Turkey, 15-19 May, 2015.
22. “Transformations of data in deterministic modellings of biological networks”, *3rd International Conference on Applied Mathematics and Approximation Theory (AMAT)*, Ankara, Turkey, 28-31 May, 2015.
23. “Stochastic modelling and inference of model parameters in complex biochemical systems”, *International Conference on Recent Advances in Pure and Applied Mathematics (ICRAPAM)*, Antalya, Turkey, 6-9 November, 2014.
24. “Stochastic inference of the complex MAPK/ERK pathway when the system has impulsive changes”, *17th International Symposium on the Biology of Actinomycetes (ISBA17)*, Kuşadası, Turkey, 8-12 October, 2014.
25. “Particle filtering approach in inference of biochemical pathways”, *11th International Conference on Monte Carlo and Quasi-Monte Carlo Methods in Scientific Computing, (MCQMC2014)* Leuven, Belgium, 6-11 April, 2014.

26. “multi-RGX: a novel background normalization method for oligonucleotides”, *8th World Congress in Probability and Statistics*, İstanbul, Turkey, 9-14 July, 2012.
27. “Reconstruction of a complex biochemical system and forecasting its behaviour”, *25th Conference of European Chapter on Combinatorial Optimization (ECCO)*, Antalya, Turkey, 26-28 April, 2012.
28. “Stochastic modelling and parameter estimation of the HCV network”, *16th INFORMS Applied Probability Conference*, Stockholm, Sweden, 6-8 July, 2011.
29. “Diffusion bridge method in inference of complex biochemical systems”, *2nd International Symposium on Computing in Science and Engineering (ISCSE’11)*, Kuşadası, Turkey, 1-4 June, 2011.
30. “Modelling lipid biosynthesis pathways of oil palms by boolean and graphical approaches” *6th International Symposium on Health, Informatics and Bioinformatics (HIBIT’11)*, İzmir, Turkey, 3-5 May, 2011.
31. “FGX implementation in two case studies”, *International Symposium on Computing in Science and Engineering (ISCSE’10)*, Kuşadası, Turkey, 3-5 June 2010.
32. “Stochastic simulation of large biochemical systems by approximate Gillespie algorithm”, *5th International Symposium on Health, Informatics and Bioinformatics (HIBIT’10)*, Antalya, Turkey, 22-25 April 2010.
33. “Testing unit root and comparison of estimates”, *14th International Congress on Computational and Applied Mathematics*, Antalya, Turkey, 29 September - 2 October 2009.
34. “Rao-blackwellized estimates for the multivariate Bayesian inference”, *International Conference of Mathematical Sciences*, İstanbul, Turkey, 4-10 August 2009.
35. “Estimation of gene expression levels for Affymetrix arrays”, *5th Conference of the Eastern Mediterranean Region of the International Biometric Society (EMR-IBS’09)*, İstanbul, Turkey, 10-14 May 2009.
36. “FGX (frequentist gene expression index) for Affymetrix arrays under non-normality assumption”, *8th International Conference on Robust Statistics (ICORS08)*, Kemer, Turkey, 8-12 September 2008.
37. “An approximation algorithm based on leap condition for stochastic simulation of biomedical systems”, *4th International Conference “Inverse problems: Modelling and Simulation” (IPM&S)*, Fethiye, Turkey, 26-30 May 2008.
38. “Inclusion of convoluted measurements in Bayesian inference of the MAPK/ERK pathway via multivariate diffusion model”, *International Symposium on Health, Informatics and Bioinformatics (HIBIT’08)*, İstanbul, Turkey, 18-20 May 2008.
39. “Bayesian inference for the MAPK/ERK pathway by considering the dependency of the kinetic parameters”, *5th Workshop on Bayesian Inference in Stochastic Processes (BISP5)*, Valencia, Spain, 14-16 June 2007.



40. “Identification and Bayesian inference of the MAPK pathway using western blot data”, *Young Statistician Meeting* (YSM), Bristol, UK, 11-12 April 2007.
41. “Exact and approximate stochastic simulations of the MAPK pathway and comparisons of simulations’ results”, *3rd International Workshop of Integrative Bioinformatics*, Harpenden, UK, 4-6 September 2006.
42. “Stochastic modelling of the MAPK signalling pathway”, *21st International Workshop on Statistical Modelling* (IWSM), Galway, Ireland, 3-7 July 2006.
43. “Stochastic modelling of the MAPK signalling pathway”, *29th Research Students’ Conference in Probability and Statistics* (RSC), Glasgow, UK, 20-23 March 2006.

### Invited Talks

1. “European cooperation for statistics of network data,”, Turkey representative in the second kick-off meeting of COST project COSTNET, Palma De Mallorca, Spain, 27 October, 2017.
2. “Protein-protein interaction networks’ data”, Institute of Applied Mathematics, METU, Ankara, Turkey, 13 December 2016.
3. “European cooperation for statistics of network data,”, Turkey representative in the first kick-off meeting of COST project COSTNET, Ribno, Slovenia, 20 September, 2016.
4. “European cooperation for statistics of network data,”, Turkey representative in the first management committee meeting of COST project COSTNET, Brussels, Belgium, 11 May 2016.
5. “Inference of the network via Gaussian graphical models”, METU-group co-representative in the fifth kick-off meeting of the joint EU-7 project PathoSys, SysPatho Workshop: System Biology ad Medicine, Ecole Sup er de Lyon, Lyon, France, 20 September 2013.
6. “Estimating the network by different inference methods”, METU-group co-representative in the third kick-off meeting of the joint EU-7 project PathoSys, SysPatho Workshop: System Biology ad Medicine, St. Petersburg, Russia, 11 September 2012.
7. “Dynamic modelling of biochemical system via graphical models”, METU-group representative in the second kick-off meeting of the joint EU-7 project PathoSys, Bah eşehir University, İstanbul, 16 February 2012.
8. “Inference of a complex network via stochastic and deterministic approaches”, Institute of Applied Mathematics, METU, Ankara, Turkey, 25 December 2012.
9. “Diffusion bridge modelling of stochastic networks”, *Workshop: Statistics for Biological Networks*, Groningen, The Netherlands, 9 September 2011.
10. “Inference of the JAK-STAT gene network via graphical models”, *the 23rd International Conference on Systems Research, Informatics and Cybernetics* (IIAS 2011), Baden, Germany, 3 August 2011.

11. “Modelling of the biological networks by probabilistic approaches”, *invited short course, International Symposium on Innovations in Intelligent Systems and Applications (INISTA 2011)*, İstanbul, Turkey, 17 June 2011.
12. “Network structure”, *invited short course, 6th International Symposium on Health, Informatics and Bioinformatics (HIBIT’11)*, İzmir, Turkey, 2 May 2011.
13. “Topics in statistical bioinformatics”, Informatics Institute, METU, Ankara, Turkey, 15 April 2011.
14. “Statistical approaches for biological networks: Simulation, modeling, and inference”, Department of Biomedical Engineering, METU, Ankara, Turkey, 8 April 2011.
15. “Network inference”, METU-group representative in the kick-off meeting of the joint EU-7 project PathoSys, BioQuant Center, University of Heidelberg, Germany, 29 October 2010.
16. “Inference in single cell dynamics”, Institute of Applied Mathematics, METU, Ankara, Turkey, 24 December 2010.
17. “Statistics for biological networks”, *invited short course* with Prof. Dr. Ernst Wit and Dr. Veronica Vinciotti, *25th International Biometric Conference (IBC2010)*, Florianópolis, Brazil, 5 December 2010.
18. “Statistical methods in system biology”, Department of Biology, METU, Ankara, Turkey, 1 May 2008.
19. “Variational approximation in inference of the kinetic parameters of the MAPK/ERK pathway”, *CRISM Workshop: Bayesian Inference for High-dimensional Data*, Warwick, UK, 14 April 2008.
20. “Analysis of the MAPK pathway: from simulation to inference”, Bioinformatic Research Center (BRC), University of Glasgow, Glasgow, UK, 18 May 2007.

## Seminar Talks

1. “A Few Project Topics in Bioinformatics”, Department of Statistics, METU, Ankara, 5 December 2013.
2. “Academic Development Programme (AGEP)-METU”, METU Northern Cyprus Campus, Güzelyurt, Cyprus, 31 March 2012.
3. “Stochastic simulation of biochemical networks”, Institute of Applied Mathematics, METU, Ankara, Turkey, 26 May 2011.
4. “Brief overview of statistical bioinformatics”, Department of Statistics, METU, Ankara, Turkey, 29 April 2010.
5. “Statistical methods in the analysis of complex gene networks”, Department of Statistics, METU, Ankara, Turkey, 3 January 2008.
6. “Two case studies in bioinformatics”, Department of Mathematics and Statistics, Lancaster University, Lancaster, UK, 26 October 2006.

7. “FGX: a Frequentist gene expression index for oligonucleotides”, Department of Statistics, University of Glasgow, Glasgow, UK, 12 April 2005.
8. “Unit root problems in time series”, Department of Statistics, METU, Ankara, Turkey, 23 December 2003.

## TEACHING EXPERIENCE

- *2017-2018* : Principle of Statistics I - II (Stat 155 and Stat 156), Decision Theory and Bayesian Analysis (Stat 565 - MSc Course), Nonparametric Statistics (Stat 460).
- *2016-2017* : Statistics for Bioinformatics (Stat 730-MSc/PhD Course) Multivariate Analysis I - II (Stat 465 and Stat 466), Principle of Statistics I - II (Stat 155 and Stat 156).
- *2015-2016* : Statistics for Bioinformatics (Stat 730-MSc/PhD Course) Multivariate Analysis I - II (Stat 465 and Stat 466), Computational Statistics (Stat 361).
- *2014-2015* : Statistics for Bioinformatics (Stat 730-MSc/PhD Course) Nonparametric Statistics (Stat 460), Applications in Statistics (Stat 495).
- *2013-2014* : Statistics for Engineers I (Stat 221), Elements of Probability and Statistics (IAM 530 - MSc Course)
- *2012-2013* : Multivariate Analysis I - II (Stat 465 and Stat 466), Decision Theory and Bayesian Analysis (Stat 565 - MSc Course)
- *2011-2012* : Multivariate Analysis I - II (Stat 465 and Stat 466), Probability Theory (Stat 391), Nonparametric Statistics (Stat 460), Statistical Methods for Informatics (Bin 502)
- *2010-2011* : Multivariate Analysis I - II (Stat 465 and Stat 466), Nonparametric Statistics (Stat 460), Decision Theory and Bayesian Analysis (Stat 565 - MSc Course)
- *2009-2010* : Multivariate Analysis I - II (Stat 465 and Stat 466), Undergraduate Research (Stat 499), Introduction to Probability and Statistics I (Stat 201), Statistics for Engineers I (Stat 221)
- *2008-2009* : Probability I - II (Stat 153 and Stat 154), Introduction to Probability and Statistics I - II (Stat 201 and Stat 202)
- *2007-2008* : Probability I - II (Stat 153 and Stat 154), Introduction to Probability and Statistics I - II (Stat 201 and Stat 202)
- *2003-2004* : Operational Research (Stat 496), Probability (Stat 253) (as a teaching assist.)
- *2002-2003* : Operational Research (Stat 496), Probability (Stat 253) (as a teaching assist.)
- *2001-2002 (Spring Semester)*: Statistics for Economists II (Econ 206) (as a student assist.)

## POSTGRADUATE SUPERVISION AND THESIS TITLES

- **PhD thesis, ongoing.** “Modeling high dimensional biological data”.

- **PhD thesis, ongoing.** “Description of complex biological networks via model-based approaches”.
- **PhD thesis, ongoing.** “Modelling breast cancer for women located in Ankara and developing a risk assessment tool with epigenetic and genetic networks via bayesian networks”.
- **PhD thesis, ongoing.** “Bayesian inference of the Gaussian graphical model for biochemical networks”.
- **PhD thesis, ongoing.** “Statistical analyses for large scale and multiple sources of biological data”.
- **PhD thesis, ongoing.** “Detection of functionally related subgraphs in biological networks”.
- **PhD thesis, ongoing.** “Inference of large-scale networks via statistical approaches”.
- **MSc thesis, ongoing.** “Analyses and modelling of ovarian cancer microarray data”.
- **MSc thesis, ongoing.** “Biological pathway construction via continuous and categorical data”.
- **PhD thesis, May 2017.** Melih Ağraz, “Different types of modellings and the inference of model parameters for complex biological systems”.
- **MSc thesis, April 2017.** Deniz Seçilmiş, “Deterministic modelling and inference of biochemical networks”.
- **MSc thesis, June 2016.** Damla Dokuzoğlu, “Application of copulas in graphical models for inference of biological systems”.
- **MSc, April 2016.** Müge Yazıcı, “Stochastic simulations of biological networks under impulses”.
- **MSc, September 2015.** Selim Yayla (Co-advisor), “Regulatory networks studied by ellipsoidal calculus”.
- **MSc, June 2015.** Gökçe Tuncer, “Comparative analysis in deterministic simulation of biological systems”.
- **MSc, August 2014.** Bilge Sürün (Co-advisor), “Analysis of motifs in microrna-transcription factor gene regulatory networks”.
- **MSc, September 2014.** Omolola Odunsi, “Detection of the distribution and parameter estimation for the departing connectivity in biological networks”.
- **MSc, July 2014.** Duygu Varol, “Comparative statistical microarray analysis of yeast data under heat shock stress”.
- **MSc, May 2013.** Ezgi Ayyıldız, “Gaussian graphical approaches in estimation of biological systems”.
- **MSc, March 2013.** Tülay Akal, “Gene expression indices for single-channel microarrays”.

## **MEMBERSHIPS**

- International Biometric Society
- International Statistical Institute
- Advisory Board of the Journal of Biostatistics-Turkish Clinics

## **LANGUAGES**

- Turkish (native), English (fluent), French (intermediate)

## **PERSONAL INTERESTS**

- Reading, particularly, philosophical, historical, and fiction books
- Playing “kanun ” (Turkish musical instrument)
- Driving and travelling
- Watching the cartoon
- Swimming