BIANCA DUMITRASCU

E-mail: biancad@princeton.edu

EDUCATION

2015–present Quantitative and Computational Biology,

Princeton University

Ph.D. Candidate (Advisor: Prof. Barbara E Engelhardt)

2013–2015 Princeton University

M.A. in Quantitative and Computational Biology

2009–2013 Massachusetts Institute of Technology

B.S. in Mathematics

PUBLICATIONS

1. PG-TS: Improved Thompson Sampling for Logistic Contextual Bandits

B. Dumitrascu, K. Feng, and B. E Engelhardt

To appear in the 32nd Conference on Neural Information Processing Systems (NeurIPS 2018).

- 2. Statistical tests for detecting variance effects in quantitative trait studies
 - **B. Dumitrascu**, G. Darnell, J. Ayroles, and B. E Engelhardt *Bioinformatics*, 2018.
- 3. BIISQ: Bayesian nonparametric discovery of Isoforms and Individual Specific Quantification from RNA-seq data
 - D. Aguiar, L. F. Cheng, **B. Dumitrascu**, F. Mordelet, A. Pai, and B. E Engelhardt *Nature Communications*, *9*(1), 2018.

PREPRINTS

- 1. **GT-TS:** Experimental design for maximizing cell type discovery in single-cell data **B. Dumitrascu**, K. Feng, and B. E Engelhardt *bioarXiv preprint* bioarXiv:386540.
- 2. **Sparse Multi-Output Gaussian Processes for Medical Time Series Prediction**L. F. Cheng, G. Darnell, **B. Dumitrascu**, C. Chivers, M. E Draugelis, K. Li, and B. E Engelhardt *arXiv preprint* arXiv:1703.09112.

WORKSHOP PAPERS

1. End-to-end training of deep probabilistic CCA for joint modeling of paired biomedical observations

G. Gundersen, **B. Dumitrascu**, J.T Ash, and B. E Engelhardt *NeurIPS Workshop on Bayesian Deep Learning, 2018*.

1. netNMF: an integrative network approach to clustering scRNA-seq data

R. Elyanow, **B Dumitrascu**, B. E Engelhardt, and B. Raphael [in submission]

2. A Bayesian nonparametric factor analysis model for gene co-expression under structured and unstructured noise

B. Dumitrascu, R. de Vito, C. Brown, and B. E Engelhardt [in preparation]

- 3. Identifying causal relationships among genes driving response to exposure with transcriptional time series data
 - **B. Dumitrascu**, J. Lu, B. Jo, I. C McDowell, T. Reddy, and B. E Engelhardt [in preparation]

EXTENDED ABSTRACTS AND INVITED TALKS

1. Bandits and Experimental Design

Models, Inference, and Algorithms, Broad Institute of MIT and Harvard, 2018 Invited Talk.

- 2. Mixed Bivariate Logistic Copulas for Depression Risk Factor Identification
 - **B. Dumitrascu**, R. de Vito, C. Brown, and B. E Engelhardt *International Society for Bayesian Analysis, Edinburgh, 2018. Poster Presentation.*
- 3. A Bayesian nonparametric factor analysis model for gene co-expression under structured and unstructured noise
 - **B. Dumitrascu**, R. de Vito, C. Brown, and B. E Engelhardt Women in Machine Learning Workshop, Barcelona, 2016. Poster Presentation.
- 4. Exploring the Glucocorticoid receptor network challenges in causal inference
 - **B. Dumitrascu**, J. Lu, B. Jo, I. C McDowell, T. Reddy, and B. E Engelhardt *Probabilistic Modeling in Genomics, Oxford University, 2016*Oral Presentation.
- 5. BIISQ: Bayesian nonparametric discovery of Isoforms and Individual Specific Quantification from RNA-seq data
 - D. Aguiar, L. F. Cheng, **B. Dumitrascu**, F. Mordelet, A. Pai, and B. E Engelhardt *American Society of Human Genetics, Baltimore, MD, 2015*. *Poster Presentation*.
- 6. Detection of variance controlling quantitative traits loci
 - **B. Dumitrascu**, G. Darnell, J. Ayroles, and B. E Engelhardt *New York Area Population Genomics Workshop, NYC, 2015 Oral Presentation*.
- 7. BTH: A Bayesian test to identify variance quantitative trait loci
 - **B. Dumitrascu**, G. Darnell, J. Ayroles, and B. E Engelhardt *American Human Genetics Society Annual Meeting, Baltimore, MD, 2015*. *Poster Presentation*.

TEACHING EXPERIENCE

Princeton University

Introduction to Java Programming (ISC231 - COS126), Fall 2015.

Interacting with Data (COS 424), Spring 2015.

Research Topics in Quantitative and Computational Biology (QCB 302), Fall 2014.

WORK EXPERIENCE

June - September 2017 Google Inc. Research Intern (video recommendation, embeddings)

ACADEMIC SERVICE

2018- Reviewer for Bioinformatics, JMLR, AISTATS

2017-2018 Graduate student mentor for Undergraduate Junior and Senior Thesis
2014–2016 Princeton Computer Science and Machine Learning Reading Group

PROFESSIONAL SKILLS

Programming: Python, R, MATLAB