# Matthew D. Koslovsky, PhD

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# RESEARCH INTERESTS

Theory and Methods: Bayesian modeling, variable selection, graphical models, non-parametric Bayes, statistical computing, multistate Markov models, R package development, varying-coefficient models, hidden Markov models, variational inference

Application: cancer prevention, smoking behaviors, mental health, addiction, physical activity, nutrition, microbiome, mHealth, ecological momentary assessment, intensive longitudinal data, environmental health, human health and performance in space

#### **EDUCATION**

## The University of Texas Health Science Center, Houston, TX

Doctor of Philosophy, Biostatistics, GPA: 4.0/4.0

Dec 2016

- · Minor: Health Promotions and Behavioral Sciences
- · Title: Deterministic Bayesian variable selection developments for binary outcomes
- · Advisor: Michael D. Swartz, PhD

## The University of Texas, Austin, TX

Bachelor of Science, Mathematics

Aug 2011

· Concentration: Scientific Computation

## EXPERIENCE

# Colorado State University, Fort Collins, CO

Assistant Professor

Starting Fall 2020

· Department of Statistics

# Rice University, Houston, TX

Post-Doctoral Research Associate

March 2018 - July 2020

- · NSF/RTG Post-Doctoral Fellowship in Data Science
- · Advisor: Marina Vannucci, PhD

# KBRwyle, Houston, TX

Biostatistician

July 2016 - March 2018

- · Human Health and Performance Contract
- · Johnson Space Center

## The University of Texas Health Science Center, Houston, TX

Pre-Doctoral Fellow

Jan 2015 - Dec 2016

- · National Cancer Institute Pre-Doctoral Fellowship
- $\cdot$  Cancer Education and Career Development Program

Pre-Doctoral Trainee

Aug 2013 - Jan 2015

· National Institutes of Health Pre-Doctoral Traineeship

## Science Systems and Applications, Inc., Hampton, VA

Summer Intern

May 2014 - Aug 2014

- $\cdot$  DEVELOP National Program
- $\cdot$  Langley Research Center

## National Space Biomedical Research Institute, Houston, TX

Summer Apprentice

May 2013 - Aug 2013

- · Biostatistics Laboratory
- $\cdot$  Johnson Space Center

## Cancer Prevention and Research Institute of Texas, Austin, TX

Summer Intern May 2010 - Oct 2010

- · University of Texas School of Public Health
- · Biostatistics Department

## **TEACHING EXPERIENCE**

# Colorado State University, Department of Statistics

Logistic/Survival Analysis for Epidemiology (STAR 580A1) Fall 2020

UTHealth, Department of Biostatistics and Data Science

Lecturer (Ad Hoc), Foundations of Biostatistics (PH1690) Fall 2019

· Student evaluation of overall effectiveness - 4.81/5.0

Lecturer (Ad Hoc), Foundations of Biostatistics (PH1690) Summer 2019

· Student evaluation of overall effectiveness - 4.86/5.0

Teaching Assistant, Theory of Biostatistics II (PH1911) Spring 2016 Teaching Assistant, Linear Models (PH1915) Fall 2015

Fall 2015 Teaching Assistant, Intermediate Biostatistics (PH1700)

Teaching Assistant, Applied Statistical Analysis I (PH1820) Summer 2015 Spring 2013

Teaching Assistant, Applied Statistical Analysis II (PH1821)

## **PUBLICATIONS**

# Submitted/In Progress

- 1. Koslovsky, M.D., Hébert, E.T., Businelle, M.S., & Vannucci, M. An efficient Bayesian varying-coefficient modeling approach for behavioral mHealth data. (Under Revisions)
- 2. Rosenberg, M.J., Koslovsky, M.D., Noyes, M., Reschke, M.F., & Clement, G. Tandem Walk in Simulated Martian Gravity and Visual Environment. (Under
- 3. Koslovsky, M.D., Liang, M.<sup>†</sup>, & Vannucci, M. A Bayesian hidden Markov model for accommodating social desirability bias in mHealth data. (In Progress)
- 4. Shaddox, E.<sup>†</sup>, Koslovsky, M.D., & Vannucci, M. A Spiked Dirichlet Process Prior for Joint Network Inference. (In Progress)
- 5. Hébert, E.T., Koslovsky, M.D., & Businelle, M.S. Time-varying relations for smoking behaviors captured in a novel, smartphone-based just-in-time adaptive intervention. (In Progress)
- 6. Koslovsky, M.D. and Vannucci, M. Dirichlet-Multinomial Regression Models with Bayesian Variable Selection for Microbiome Data. In S. Datta & S. Guha (Eds.), Statistical Analysis of Microbiome Data. Springer Verlag. (In Progress) † indicates PhD student in Dr. Vannucci's research group at Rice University

# Statistical Methodology

7. Koslovsky, M.D., Vannucci, M. (2020) MicroBVS: Dirichlet-tree multinomial regression models with Bayesian variable selection - an R package. BMC Bioinformatics, 21(301). https://doi.org/10.1186/s12859-020-03640-0

- 8. **Koslovsky, M.D.**, Hoffman, K., Daniel, C., & Vannucci, M. (2020). A Bayesian model of microbiome data for simultaneous identification of covariate associations and prediction of phenotypic outcomes. *Annals of Applied Statistics*. (In press)
- Koslovsky, M.D., Swartz, M.D., Chan, W., Leon-Novelo, L., Wilkinson, A.V., Kendzor, D.E., & Businelle, M.S. (2018). Bayesian variable selection for multistate Markov models with interval-censored data in an ecological momentary assessment study of smoking cessation. *Biometrics*, 74(2), 636-644.
- Koslovsky, M.D., Swartz, M.D., Leon-Novelo, L., Chan, W., & Wilkinson, A.V. (2018). Using the EM algorithm for Bayesian variable selection in logistic regression models with related covariates. *Journal of Statistical Computation* and Simulation, 88(3), 575-596.

# **Applications**

- Zwart, S.R., Rice, B.L., Dlouhy, H., Shackelford, L.C., Heer, M., Koslovsky, M.D., & Smith, S.M. (2018). Dietary acid load and bone turnover during long-duration spaceflight and bed rest. The American Journal of Clinical Nutrition, 107(5), 834-844.
- Conkin, J., Sanders, R.W., Koslovsky, M.D., Wear, M.L., Kozminski, A.G., & Abercromby, A.F. (2018). A systematic review and meta-analysis of decompression sickness in altitude physiological training. *Aerospace Medicine and Human Performance*, 89(11), 941-951.
- Koslovsky, M.D., Hébert, E.T., Swartz, M.D., Chan, W., Leon-Novelo, L., Wilkinson, A.V., Kendzor, D.E. & Businelle, M.S. (2017). The time-varying relations between risk factors and smoking before and after a quit attempt. Nicotine and Tobacco Research, 20(10), 1231-1236.
- Conkin, J., Wessel, J.H., Norcross, J.R., Bekdash, O.S., Abercromby, A.F., Koslovsky, M.D., & Gernhardt, M.L. (2017). Hemoglobin oxygen saturation with mild hypoxia and microgravity. Aerospace Medicine and Human Performance, 88(6), 527-534.

# **Proceedings**

 Meyers, J., Garcia, Y., Arellano, J., Boley, L., Goodenow D., Kerstman, E., Koslovsky, M.D., Reyes, D., Saile, L., Taiym, W., & Young, M. (2018, September 16-21). Validation of the NASA Integrated Medical Model: A Space Flight Medical Risk Prediction Tool. Paper presented at *Probabilistic Safety* Assessment and Management 14, Los Angeles, CA.

## **PRESENTATIONS**

- Koslovsky, M.D.\*. Bayesian Methods for Behavioral mHealth Data. Colorado State University, Department of Statistics. Fall 2020. (Departmental Seminar)
- Koslovsky, M.D.\*. Bayesian Methods for Behavioral mHealth Data. University of Colorado Denver, Department of Biostatistics & Informatics. Fall 2020. (Departmental Seminar)
- Koslovsky, M.D.\*. Bayesian Methods for Behavioral mHealth Data. University of Missouri, Department of Statistics. Fall 2020. (Departmental Seminar)
- Koslovsky, M.D.\*. Bayesian Methods for Behavioral mHealth Data. Montana State University, Department of Mathematical Sciences. Fall 2020. (Departmental Seminar)

- Koslovsky, M.D.\*, Hoffman, K., Daniel-MacDougall, C., & Vannucci, M. "A Bayesian Model of Microbiome Data for Simultaneous Identification of Covariate Associations and Prediction of Phenotypic Outcomes." iBright, Houston, TX. Nov 2019. (contributed poster presentation)
- Koslovsky, M.D.\*, Hoffman, K., Daniel-MacDougall, C., & Vannucci, M. "A Bayesian Model of Microbiome Data for Simultaneous Identification of Covariate Associations and Prediction of Phenotypic Outcomes." Joint Statitsics Meetings, Denver, CO. Aug 2019. (contributed poster presentation)
- Koslovsky, M.D.\*, Hoffman, K., Daniel-MacDougall, C., & Vannucci, M. "A Bayesian Model of Microbiome Data for Simultaneous Identification of Covariate Associations and Prediction of Phenotypic Outcomes." BigDIA, Houston, TX. Dec 2018. (contributed poster presentation)
- Yu, D., Sedory, A.C., Mohammadi, K., Koslovsky, M.D., & Swartz, M.D.\*.
  "Trio\_RVEMVS: A fast Bayesian variable selection method for trios that identifies individual rare variants," International Genetic Epidemiology Society Meetings, San Diego, CA, Oct 2018. (platform presentation)
- Koslovsky, M.D.\*, Arellano, J., Schaefer, C., Feiveson, A., & Young, M. "CommClust: A network-based algorithm for clustering multivariate repeated measures data." NASA Human Research Program Investigators' Workshop. Galveston, TX. Jan 2018. (contributed poster presentation)
- Koslovsky, M.D.\*, "Immersive Data Analysis for NASA Biomedical Data," Rice Data Science Conference, Houston, TX, Oct 2017. (contributed oral presentation)
- Koslovsky, M.D.\*, "Immersive Data Analysis for NASA Biomedical Data," Texas Collaboration Center Data Analytics Workshop, Houston, TX, Oct 2017 (contributed oral presentation)
- Koslovsky, M.D.\*, Young, M., Schaefer, C., Arellano, J., & Feiveson, A. "A Network-based Algorithm for Clustering Multivariate Longitudinal Data." Joint Statistical Meetings. Baltimore, MD. Aug 2017. (contributed oral presentation)
- Koslovsky, M.D.\*, Swartz, M.D., Chan, W., Leon-Novelo, L., Wilkinson, A.V., Kendzor, D.E., & Businelle, M.S. "Deterministic Bayesian variable selection for multistate models, with applications to ecological momentary assessment of an attempt to quit smoking." Joint Statistical Meetings. Chicago, IL. Aug 2016. (contributed oral presentation)
- Koslovsky, M.D., Swartz, M.D., Chan, W., Leon-Novelo, L., Wilkinson, A.V., Kendzor, D.E., & Businelle, M.S., "Deterministic Bayesian Variable Selection for Binary Outcomes." Joint Statistical Meetings. Seattle, WA, Aug 2015. (contributed oral presentation)
- Burley, B., Erickson, C., Fenn, T., Hope, J., & Koslovsky, M.D.\* "New England Water Resources-Historical Tracking of Harmful Algal Blooms Using Landsat Missions from 1984-2014." DEVELOP Summer Closeout at NASA Headquarters. Washington, D.C. Aug 2014. (contributed poster presentation)
- Swartz, M.D.\*, Koslovsky, M.D., Vandewater, E.A., & Wilkinson, A.V. "A Stochastic Search Through Smoking Images in Movies, Genetic, and Psychosocial Factors Associated with Smoking Initiation in Mexican American Youths." International Genetic Epidemiology Society Meetings 2014, 23rd Annual Conference. Vienna, Austria. Aug 2014. (contributed oral presentation)
  \* indicates presenter

### AWARDS

- Dr. M. Stewart West Memorial Scholarship, 2015
- UTHealth Division of Biostatistics Travel Award, 2015
- Richard D. Remington Memorial Student Scholarship, 2014
- Robert. H Bigelow Endowed Scholarship, 2013

#### **MENTORING**

- Yefei Zhang, UTHealth, PhD Biostatistics candidate, Dissertation Committee, 01/2017-Current
- Scott Liang, Rice University, PhD Statistics student, Co-mentor, 03/2019-Current
- James Warner, Rice University, Rice Undergraduate Data Science Summer Program, 2018
- Karan Adams, Rice University, Rice Undergraduate Data Science Summer Program, 2018
- Stoyan Komitov, Rice University, Rice Undergraduate Data Science Summer Program, 2018
- Alex Aguilar, Rice University, PhD Statistics candidate, NASA Summer Intern, 2018
- Austin Vo, University of Central Florida, NASA Summer Intern, 2017
- UTHealth New Student Mentor, Fall 2013

# COMPUTER SKILLS

 $Languages \ \& \ Software: \ R, \ C++, \ Rcpp, \ Shiny, \ LaTEX, \ STATA, \ SAS, \ WinBUGS \ Annual S$ 

#### PROFESSIONAL Member

## **AFFILIATION**

• American Statistical Association, 2015 - Current

## PROFESSIONAL Reviewer

## **SERVICE**

• Biometrical Journal, Biometrics, Biostatistics, Nature Communications

## Board Member

• Johnson Space Center IRB, 2017 - 2018

#### Consultation

 Conference for Food Protection 2019: Program Standards Committee-Standard 8 Re-Evaluation of Staffing Level Model

# CONTINUING EDUCATION

- HACASA Short Course "Randomized Clinical Trials-Replacing Traditional Analyses with Better Alternatives," Houston, TX, May 2018
- Joint Statistical Meetings Short Course "Network Meta-Analysis," Baltimore, MD, Aug 2017
- Joint Statistical Meetings Short Course "Evolution of Classification," Baltimore, MD, Aug 2017
- NASA Human Research Program Investigator's Workshop "A New Dawn: Enabling Human Space Exploration," Galveston, TX, Jan 2017
- Technology Collaboration Center "Omics Workshop," Houston, TX, Spring 2017

- Tableau Conference 2016 Tableau Classroom Training- "Tableau Desktop II," Austin, TX, Fall 2016
- ENAR Short Course "An Introduction to Statistical Machine Learning," Austin, TX, Spring 2016
- ENAR Tutorial Session "Data Visualizations in R with shiny and ggplot2," Austin, TX, Spring 2016
- ENAR Tutorial Session "High Performance Computing with R," Austin, TX, Spring 2016
- ASA Biopharmaceutical Section FDA Industry Statistics Workshop "Equivalence and Similarity Testing," Washington, DC, Fall 2015
- ASA Biopharmaceutical Section FDA Industry Statistics Workshop "Designing Observational Comparative Studies Using Propensity Score Methodology in Regulatory Settings," Washington, DC, Fall 2015
- Joint Statistical Meetings "Adaptive Methods for Modern Clinical Trials," Seattle, WA, Summer 2015
- UT Summer Statistics Institute "Introduction to Mixed Models with Applications," Austin, TX, Summer 2015
- UT Summer Statistics Institute "Big Data Analytics," Austin, TX, Summer 2015

REFERENCES

Marina Vannucci, PhD Noah Harding Professor of Statistics Department of Statistics Rice University marina@rice.edu 713-348-6132

Michael D. Swartz, PhD Michael.D.Swartz@uth.tmc.edu Associate Professor 713-500-9570 Department of Biostatistics and Data Science University of Texas Health Science Center at Houston

Wenyaw Chan, PhD Wenyaw.Chan@uth.tmc.edu Professor 713-500-9321 Department of Biostatistics and Data Science University of Texas Health Science Center at Houston

 $\begin{array}{lll} \textit{Michael Businelle, PhD} & \textit{Michael-Businelle@OUHSC.edu} \\ \textit{Associate Professor} & 405\text{-}271\text{-}8001 \text{ x}50460 \\ \textit{Oklahoma Tobacco Research Center} \\ \textit{The University of Oklahoma Health Sciences Center} \end{array}$ 

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