

Rachael Hageman Blair, Ph.D.
Curriculum Vitae
(updated 11/2021)

Department of Biostatistics
State University of New York at Buffalo
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Education

- Ph.D. (2007) Case Western Reserve University, Cleveland, OH
Mathematics (advisor: Daniela Calvetti)
Dissertation: *Bayesian methods for large-scale parameter estimation and sensitivity analysis for myocardial metabolism.*
- M.S. (2006) Case Western Reserve University, Cleveland, OH
Mathematics (advisor: Daniela Calvetti)
Thesis: *Iterative methods for blind deconvolution.*
- B.S. (2002) State University of New York College at Fredonia, Fredonia, NY
Mathematics.

Professional Appointments

- 07/2019 – present **State University of New York at Buffalo**, Buffalo, NY
Institute for Computational and Data Sciences
Director of the MPS in Data Science and Applications Program
- 07/2018 – present **State University of New York at Buffalo**, Buffalo, NY
Department of Biostatistics
Associate Professor
- Institute for Computational and Data Sciences
Core Member
- 01/2018 – present **Baypath University**, Longmeadow, MA
Data Science Program
Adjunct Professor

08/2011 – 07/2018 **State University of New York at Buffalo**, Buffalo, NY
Department of Biostatistics
Assistant Professor

08/2011 – 08/2017 **Roswell Park Cancer Institute**, Buffalo, NY
Department of Biostatistics
Adjunct Assistant Professor

Research Experience & Training

08/2007 – 08/2011 **The Jackson Laboratory**, Bar Harbor, ME
Postdoctoral Associate (mentor: Gary Churchill)

- Major projects include statistical methodologies for causal inference, high-throughput data analyses of microarray and quantitative trait loci (QTL) data in collaborations with NIGMS Center for Genome Dynamics

08/2005 – 05/2007 **Case Western Reserve University**, Cleveland, OH
Research Fellow

- Department of Mathematics and the NIGMS Center for Modeling Integrated Metabolic Systems (MIMS).

05/2001 – 08/2001 **Worcester Polytechnic Institute**, Worcester, MA
NSF Undergraduate Research

- NSF Research Experience for Undergraduates (REU) in collaboration with DEKA Research Corporation.

Select Teaching Experience & Course Development

01/2018 – present **Baypath University**, Longmeadow, MA
Developed two new graduate courses in data mining for an online Data Science program. The course and series are supported through Canvas, and zoom technology is used to facilitate additional meetings and office hours.

07/2017 – present **University at Buffalo**, Buffalo, NY
Developing a new graduate course based on data challenges to launch in the summer 2018. The course will emphasize team-based innovation for solving biomedical problems put forth in the DREAM challenges. Course development is funded through the UB Center for Educational Innovation.

08/2012 – present **University at Buffalo**, Buffalo, NY
Instructor: Statistical Data Mining I-II (STA 545 and STA 546)
Developed a data mining course series that spans topics in supervised (STA 545) and unsupervised (STA 546) learning. The course follows a hybrid online format, which makes extensive use of video capture, discussion forums, and other organizational tools through UB learns. The course series is now a required component of the Data Science Masters in Engineering and elective in the Computational and Data Enabled Science and Engineering programs.

- 08/2007 – 08/2009 **The Jackson Laboratory**, Bar Harbor, ME
Guest lectures and mentoring for the Summer Student Program in the Center for Genome Dynamics for high-school students from various magnet schools.
- 05/2003 – 08/2007 **Case Western Reserve University**, Cleveland, OH
School of Graduate Studies
School of Medicine, Department of Mathematics, Department of Computer Science and Engineering
Instructor: Calculus I-III (undergraduate course), Applied Statistics for Engineers (graduate course), Applied Mathematics and Statistics (Summer Medical and Dental Education Program).
Taught a variety of math and statistics courses as instructor on record to both undergraduate and graduate students. Full responsibility was assumed for development and delivery of lectures, assignment of homework, exam development, grading, and holding office hours.

Professional Awards and Honors

- 2021 BayPath University, Distinguished Teaching Award
- 2017 UB School of Public Health and Health Professions - Outstanding Junior Researcher
- 2016 NLMNIH Travel Award, Pacific Symposium for Biocomputing
- 2015 SAMS I Ideas/Innovation Lab Participant
- 2014 UB Individual Development Award, Spring
- 2013 Department of Biostatistics Award for Teaching Excellence
- 2012 Department of Biostatistics Award for Teaching Excellence
- 2008 High Dimensional Statistics in Biology Workshop NIH travel award
- 2007 SIAM, ICIAM travel award
- 2007 Joint Mathematics Meeting travel award
- 2006 SIAM, Conference on the Life Sciences award
- 2006 European Mathematical Society, Mathematical Models of the Heart travel award
- 2006 Mathematical Biosciences Institute, Cardiac Workshop travel award
- 2004 Mathematical Sciences Research Institute, Imaging Workshop travel award
- 2003 Graduate Dean's Award in Instructional Excellence, Case Western University

Peer Reviewed Publications

- 1) McComb M, **Hageman Blair R**, Lysy M and Ramanathan M (in press) Machine learning-guided, big data-enabled, biomarker-based systems pharmacology: modeling the stochasticity of natural history and disease progression. *Pharmacokinetics and Pharmacodynamics* doi: <https://doi.org/10.1007/s10928-021-09786-5>
- 2) Lang JK, Karthikeyan B, Quinones-Abramana A, **Hageman Blair R**, Early AP, Levine EG, CS Umesh, Blanco JG, O'Connor T (2021) CBR3 V244M is associated with LVEF reduction in breast cancer patients treated with doxorubicin. *Cardio-Oncology* 7(1): 17. doi: 10.1186/s40959-021-00103-0
- 3) Zhu K, Browne RW, **Hageman Blair R**, Bonner MR, Tian M, Niu Z, Deng F, Farahat Z, Mu L (2021) Changes in arachidonic acid (AA)- and linoleic acid (LA)-derived hydroxy metabolites and their interplay with inflammatory biomarkers in response to drastic changes

in air pollution exposure. *Environmental Research*.

- 4) Tian M., **Hageman Blair, R.**, Mu L., Bonner M., Browne R. and Yu, H. (2021) A framework for stability-based module detection in correlation graphs. *Statistical Analysis and Data Mining*, 1-15. doi: [10.1002/sam.11495](https://doi.org/10.1002/sam.11495)
- 5) Farhat, Z., Hershberger, P.A., Freudenheim, J.L., Mammon, M.M., **Hageman Blair, R.**, Aga, D., and Mu, L. (2021) Types of garlic and their anticancer and antioxidant activity: A review of the epidemiologic and experimental evidence. *European Journal of Nutrition*, 1-25
- 6) Yu, H. and **Hageman Blair, R.** (2020). Scalable module detection for attributed networks with applications to breast cancer. *Journal of Applied Statistics*, 1-18. Doi: <https://doi.org/10.1080/02664763.2020.1803811>
- 7) Cejas, R. B., Wang, J., **Hageman Blair, R.**, Liu, S., & Blanco, J. G. (2020). Comparative genome-wide DNA methylation analysis in myocardial tissue from donors with and without Down syndrome. *Gene*, 764, 145099. Doi: [10.1016/j.gene.2020.145099](https://doi.org/10.1016/j.gene.2020.145099)
- 8) Bubier, J., Philip, V., Quince, C., Campbell, J., Zhou, Y., Vishnivetskaya, T., Duvvuru, S., **Hageman Blair, R.**, Ndikum, J., Donohue, K.D. Phillips, C., Foster, C.M., Mellert, D.J., Weinstock, G., Culiati, C.T., O'Hara, B.F., Weinstock, G., Culiati, C.T., O'Hara, A.V., Podar, M. and Chesler, E.J. (2020). A microbe associated with sleep revealed by a novel systems genetic analysis of the microbiome of collaborative cross mice. *Genetics* 214(3) 719-733. Doi: <https://doi.org/10.1534/genetics.119.303013>
- 9) Yu H, Moharil J, **Hageman Blair R*** (2020) BayesNetBP an R package for probabilistic reasoning in Bayesian Networks. *Journal of Statistical Software* (94) 3, <https://doi.org/10.18637/jss.v094.i03>. *Senior author - first author graduate student
- 10) Mu L, Niu Zhongzheng, **Hageman Blair R**, Yu H, Browne RW, Bonner MR, Fanter T, Deng F, Swanson M (2019) Metabolomics profiling before, during and after the Beijing Olympics: a panel study of within individual differences during periods of high and low air pollution 127(5):57010 *Environmental Health Perspectives*.doi: doi: 10.1289/EHP3705
- 11) Yu H and **Hageman Blair R*** (2019) Integration of probabilistic regulatory networks into constraint-based models of metabolism with applications to Alzheimer's disease. 20 (386) *BMC Bioinformatics* doi: <https://doi.org/10.1186/s12859-019-2872-8> *Senior author - first author graduate student
- 12) Grieshober L, Wactawski-Wende J, **Hageman Blair R**, Mu L, Liu Jingmin, Nie J, Carty C, Hale L, Kroenke C, LaCroix A, Reiner AP, Ochs-Balcom HM (2019) A cross-sectional analysis of telomere length and sleep in the Women's Health Initiative. 188(9): 1616-1626 *American Journal of Epidemiology*. Doi: 10.1093/aje/kwz134
- 13) Yu H, Chapman B, Di Florio, A, Eischen E, Gotz D, Jacob M, **Hageman Blair R*** (2018) Bootstrapping estimates of stability for clusters, observations and model selection, 1-24 *Computational Statistics*. Doi: <https://doi.org/10.1007/s00180-018-0830-y> *Senior author - first author graduate student

- 14) Quiñones-Lombrana A, **Hageman Blair R**, Blanco JG. (2017) Investigation of the role of DNA methylation in the expression of ERBB2 in human myocardium. 628 286-294 *Gene* doi: 10.1016/j.gene.2017.07.058
- 15) Hao F and **Hageman Blair R*** (2016) A comparative study: classification vs. user-based collaborative filtering for clinical risk prediction. 172(16) *BMC Medical Research Methodology* *Senior author - first author graduate student
- 16) Hoefer CA*, **Hageman Blair R***, Blanco JG (2016) Development of a CART model to predict the synthesis of cardiotoxic daunorubicinol in heart tissue samples from donors with and without down-syndrome. *Journal of Pharmaceutical Sciences* 105(6) 2005-8. *equal contributions
- 17) Moharil J, May P, Gaile D, **Hageman Blair R*** (2016) Belief Propagation in Genotype-phenotype networks. *Statistical Applications in Genetics and Molecular Biology* 15(1):39-53. *Senior author - first author graduate student
- 18) Hoefer CC, Quinones-Lombrana A, **Hageman Blair R**, Blanco JG (2016) Role of DNA methylation on the expression of the anthracycline metabolizing enzyme AKR7A2 in human heart. *Cardiovascular Toxicity* 16(2): 182-192.
- 19) Farasat A, Nikolaev A, Srihari S, **Hageman Blair R*** (2015) Probabilistic Graphical Models in Modern Social Networks. *Social Network Analysis and Mining* 5(1) 1-18. *Senior author - first author graduate student
- 20) Griesenhofer L., **Hageman Blair R*** (2014) Evaluating performance of link prediction in scale-free evolving networks and a Facebook community. *Social Network Analysis and Mining* 4(183). * Senior author - first author graduate student
- 21) Quinnes-Lombrana A, Ferguson D., **Hageman Blair R**, Kalabus JL, Redzematovic A, and Blanco J.G. (2014) Individual variability in the cardiac expression of anthracycline reductases in donors with- and without- Down Syndrome. *Pharmaceutical Research* 2014; (31)7: 1644-55.
- 22) Alkakna A, Choi S, Savage H, **Hageman Blair R**, Gu T, Svenson KL, Churchill GA, Hibbs M, Korstaje R. (2012) Pla2g12b and Hpn are genes identified by mouse ENU mutagenesis that affect HDL cholesterol. *PLoS One* 7(8): e43139.
- 23) **Hageman Blair R**, Trichler DL, Gaile DP. (2012) Mathematical and Statistical Modeling in Cancer Systems Biology. *Frontiers in Physiology* 3(227).
- 24) **Hageman Blair R**, Kliebenstein DJ, Churchill GA. (2012) What can causal networks tell us about metabolic pathways? *PLoS Computational Biology* 8(4): e10024258.
- 25) Leduc MS, **Hageman Blair R**, Tsaih SW, Verdugo RA, Walsh K, Churchill GA, Paigen B. (2012) Using bioinformatics and systems genetics to dissect HDL cholesterol genetics in an MRL/MpJ x SM/J intercross. *The Journal of Lipid Research* 53: 1163-1175.
- 26) Leduc MS, **Hageman RS**, Tsaih SW, Verdugo RA, Walsh K, Churchill GA, Paigen B. (2011) Integration of QTL and bioinformatic tools to identify candidate genes for

triglycerides in an MRL/MpJ x SM/J intercross. *The Journal of Lipid Research* 52: 1672-1682.

- 27) **Hageman RS**, Leduc M, Paigen B, Korstanje R, and Churchill GA. (2011) A Bayesian Framework for Inference of the Genotype-Phenotype Map for Segregating Populations. *Genetics* 187(4): 1163-1170.
- 28) **Hageman RS**, Leduc MS, Caputo CR, Tsaih SW, Paigen B, Churchill GA, and Korstanje R. (2011) Uncovering Genes and Regulatory Pathways Related to Urinary Albumin Excretion in Mice. *Journal of the American Society of Nephrology* 22: 73-81. (Selected by the Faculty of 1000 Biology.)
- 29) Leduc MS, **Hageman RS**, Meng Q, Verdugo RA, Tsaih SW, Churchill GA, Paigen B, Yuan R. (2010) Identification of genetic determinants of IGF-1 levels and longevity among mouse inbred strains. *Aging Cell* 9(5): 823-836.
- 30) **Hageman RS**, Wagener A, Hantschel C, Svenson KL, Churchill GA, and Brockmann GA. (2010) High fat diet leads to tissue specific changes reflecting risk factors for diseases in DBA/2J mice. *Physiological Genomics* 42:55-66.
- 31) *Calvetti D, ***Hageman R**, *Occhipinti R, and *Somersalo E. (2008) Dynamic Bayesian sensitivity analysis of a myocardial metabolic model. *Mathematical Biosciences* 212:1-21.
* Authors alphabetized with equal contributions
- 32) *Calvetti D, ***Hageman R**, and *Somersalo E. (2006) Large-scale Bayesian parameter estimation for a three-compartment cardiac metabolism model during ischemia. *Inverse Problems* 22:1797-1816. * - Authors alphabetized with equal contributions

Peer Reviewed Conference Proceedings

1. Yu H and **Hageman Blair R*** (2016) A framework for attribute-based community detection with applications to integrative functional genomics. *Pacific Symposium on Biocomputing* 21:69-90. *Senior author - first author graduate student (rank 3/52 - conferences in Bioinformatics and Computational Biology via Microsoft academic search)

Book Chapters

1. **Hageman Blair R**, Trichler DL, Gaile DP. (2014) Cancer Systems Biology. In *Recent Advances in Systems Biology*. NovaScience Publishers.

Software Development (open source)

1. clusterboot (<https://cran.r-project.org/web/packages/bootcluster/index.html>)
Supports stability estimation for clustering (described in Yu. et al. *Computational Statistics*, 2018)
2. BayesNetBP (<https://cran.r-project.org/web/packages/BayesNetBP/index.html>)
Supports probabilistic reasoning in Bayesian Networks that are mixed, discrete and continuous, also has a shiny application interface. (Yu *et al.*, *Journal of Statistical Software*, 2020).
3. GeneNetBP (<https://cran.r-project.org/web/packages/geneNetBP/index.html>)
Supports belief propagation in genotype-phenotype networks (Moharil *et al.*, 2016).
4. wfg (<https://cran.r-project.org/web/packages/wfg/index.html>)

Supports community detection with mixed attributes (Yu *et al.*, Journal of Applied Statistics, 2020).

Conference Proceedings

1. **Hageman Blair R**, Gaile D, Grabowski J, Kuhlmann D, Markatou M, Tritchler D, Wilding G, Yu G. (2018) *Integrative learning through the data-driven classroom: the DREAM challenge*. AMIA 2018 Informatics educators' forum, New Orleans, LA.
2. Yu H and **Hageman Blair R*** Attribute-based module detection for the elucidation of tissue-specific pathways for oncoimmunology. Topic contributed paper, Joint Statistical Meetings, Baltimore, 2017. *Senior author - first author graduate student
3. Smith RV, McGann D, Gotwals R, Symonds R, **Hageman RS**, Vedell PT, and Churchill GA. (2009) *Independent Studies in Computational Biology*. National Consortium for Specialized Secondary Schools of Math, Science and Technology Journal, 15(1):10-16.
4. Calvetti D, **Hageman RS**, Occhipinti R, and Somersalo E. (2005) *Large-scale statistical parameter estimation during circulatory occlusion*. International Conference on Computational and Mathematical Methods in Science and Engineering (CMMSE) proceedings.
5. Calvetti D, **Hageman RS**, Occhipinti R, and Somersalo E. (2005) *Large-scale statistical estimation of metabolic parameters at steady-state*. International Conference on Computational and Mathematical Methods in Science and Engineering (CMMSE) proceedings.

Select Abstracts from Presentations at Conferences and Meetings (selected from ~40)

- 2020 Naragon-Gainey, K., Park, J., McMahon, T. P., **Hageman Blair, R.**, & Yu, H. (2020, *Predictors and outcomes of perceived emotion regulation success in daily life*. Annual Meeting of the Society for Ambulatory Assessment, Melbourne, Australia.
- 2019 Yu H and **Hageman Blair R.** Regularized regression by graph propagation for genomic data analysis, JSM, Denver, CO. (poster presentation – Yu H)
- 2019 Millen A, Hall K, Kamm K, Liu Z, Krajewski B.J., Wactawski-Wende, J, Lema G, **Hageman Blair R**, Sun, Y, and McSkimming D. *Mail-based stool collection in women with and without age-related macular degeneration (AMD)*, ARVO annual meeting, Baltimore, MD. (poster presentation – Millen A)
- 2018 Asma H, Miller L, Tian M, Wintrob Z, **Hageman Blair R.** *Drosophila embryo structure from transcriptome: 3-D data recovery post derop-seq profiling using 20 genes*, ISCB-DREAM RECOMB, NY, NY. (poster presentation – students)
- 2018 Hageman Blair R. (2018) *Module detection in attributed networks for the association with outcomes minisymposium: Air pollution exposure in metabolomics.*) ISES-ISEE Joint Annual Meeting, Ottawa, CA, (oral presentation)

- 2017 Yu H and **Hageman Blair R.** BayesNetBP: Probabilistic Reasoning in Bayesian Networks. Complex Traits Consortium, Memphis, TN. (oral presentation – Yu H).
- 2017 Yu H, Chapman B, DiFlorio A, Eischen E, Jacob M, Gotz D, and **Hageman Blair R.** *Bootstrapping estimates of stability for clusters, observations and model selection.* Upstate Chapters of the American Statistical Association UPSTAT-2017. Buffalo, NY (oral presentation – Yu H) Award: “Best Young Researchers Award”, Category A: Methodology – silver award
- 2017 Grieshober L, Wactawski-Wende J, **Hageman Blair R,** Mu L, Liu K, Nie J, Carty CL, Hale L, Kroenke C, LaCroix AZ, Reiner AP, Ochs-Balcom HM. *Longer sleep duration is associated with longer telomeres: results from the Women’s Health Initiative Society for Epidemiologic Research,* Seattle, WA, 2017. (poster – Grieshober L)
- 2016 **Hageman Blair R,** Chapman B, DiFlorio A, Eischen E, Gotz D, Yu H. *Flexible bootstrapping and analytic approaches towards the clustering of complex medical data.* *F1000Research* 2016, **5**:2724 (doi: [10.7490/f1000research.1113440.1](https://doi.org/10.7490/f1000research.1113440.1)) B2DK All Hands Meeting, Bethesda MD, 2017. (poster – Hageman Blair R)
- 2017 **Hageman Blair R***, Chapman B*, DiFlorio A*, Eischen E*, Gotz D*. *Interactive ensemble clustering for medical data with application to mood disorders.* B2DK All Hands Meeting, Bethesda MD, 2017. (poster – Eischen E) *Authors alphabetized with equal contributions
- 2016 Yu H, Chapman B, DiFlorio A, Eischen E, Jacob M, Gotz D, and **Hageman Blair R.** *Bootstrapping estimates of stability for clusters, observations and model selection.* ISCB Rocky Mountain Bioinformatics Conference, Aspen, CO, 2016. (poster – Yu H)
- 2016 Moharil J and **Hageman Blair R.** *Probabilistic reasoning in genotype-phenotype networks.* ISCB RSG-DREAM Phoenix, AZ, 2016. (oral presentation – Blair RH)
- 2016 Tison GH, Nah G, Olgin, J, Vittinghoff E, Howard B, Foraker R, Allison M, Casanova R, **Hageman Blair R,** Breathett K, Klein L, Parikh N. *Identifying novel predictors for incident heart failure using statistical learning techniques in the Women’s Health Initiative (WHI) cohort.* AHA Scientific Sessions, New Orleans LA, 2016. (poster – Tison GH)
- 2016 Yu H and **Hageman Blair R.** *A framework for attribute-based community detection with applications to integrative functional genomics.* Pacific Symposium on Biocomputing, Kona, HI. (oral presentation –Hageman Blair R)
- 2015 Crowley A, Zhoulin H, **Hageman Blair R.** *Data mining major league baseball’s pace of play problem.* New England Symposium on Statistics in Sports, Boston, MA. (poster – Crowley A)
- 2014 Moharil J., May P, Gaile D., **Hageman Blair R.** *Belief Propagation in Genotype-phenotype networks.* BioMedical Engineering Society (BMES) annual meeting, San Antonio TX. (poster – Moharil J)
- 2010 **Hageman RS,** Leduc M, Paigen B, Korstanje R, and Churchill GA. *A Bayesian Framework for Inference of the Genotype-Phenotype Map for Segregating Populations.* International

- conference on Research in Computational Molecular Biology (RECOMB), Lisbon, Portugal. (poster - Hageman RS)
- 2009 **Hageman RS**, Aljakna A, Svenson KL, Churchill GA and Korstanje R. *Multiple mutations perturb a small number of pathways in an HDL mutagenesis screen*. International Symposium on Atherosclerosis, Newport, RI. (poster – Hageman RS)
- 2009 **Hageman RS**, Aljakna A, Svenson KL, Churchill GA and Korstanje R. *Multiple mutations perturb a small number of pathways in an HDL mutagenesis screen*. Complex Traits Consortium, Manchester, England, UK. (poster – Hageman RS)
- 2008 **Hageman RS**, Brockman G and Churchill GA. *Differences in gene expression in the adipose tissues of DBA/2 mice under a high fat diet*. High dimensional statistics in biology workshop, Cambridge, UK. (poster – Hageman RS)
- 2007 Calvetti D, **Hageman RS** and Somersalo E. *Dynamic Bayesian sensitivity analysis in myocardial metabolism*. International Congress of Industrial and Applied Mathematics (ICIAM), Zurich, Switzerland. (oral presentation – Hageman RS)
- 2007 Calvetti D, **Hageman RS** and Somersalo E. *Bayesian Large scale dynamic sensitivity analysis in myocardial metabolism*. Joint Mathematics Meeting, New Orleans, LA. (poster – Hageman RS)
- 2007 Calvetti D, **Hageman RS** and Somersalo E. *Bayesian sensitivity analysis in cardiac metabolism*. Biomedical Engineering Society (BMES): annual meeting, Chicago, IL. (poster – Hageman RS)
- 2006 Calvetti D, **Hageman RS** and Somersalo E. *Large scale parameter estimation for the dynamic cardiac metabolism during ischemia*. Society for Industrial and Applied Mathematics (SIAM): Conference on the Life Sciences, Raleigh, NC. (oral presentation – Hageman RS)
- 2007 **Hageman RS**, Kline K, Wilkins L, Larson C. *Modeling the dynamics of a motorized wheelchair*. Joint Mathematics Meeting, San Diego, CA. (poster – Hageman RS)

Invited Presentations

- 2021 Controlling networks and predicting optimal sets of perturbations
The Classification Society Annual Meeting
Bucknell University
- 2021 Stability analysis for clustering problems and community detection in graphical models
Penn State University, Department of Statistics
State College, PA
- 2019 A probabilistic framework for fusing genetic and metabolic networks with applications to Alzheimer’s disease
The Fields Institute for Research in Mathematical Sciences
Toronto, Canada
- 2019 Bootstrapping the stability of clusters and networks
Fall Eastern Meeting of the American Mathematical Society

- Binghamton, NY
- 2018 Probabilistic reasoning with networks
UB Department of Chemical and Biological Engineering
- 2018 Biomedical Data Science Education
AMIA Informatics Educator Forum
New Orleans, LA
- 2017 Attribute-based module detection with applications to integrated functional genomics
AbbVie, Redwood City, CA
- 2017 Reasoning with Networks
Sage Bionetworks, Seattle, WA
- 2016 Perturbing Gene Networks
Association of Women in Mathematics, local chapter
Buffalo, NY
- 2014 Belief Propagation in Genotype-Phenotype Networks
Department of Biological Statistics and Computational Biology (seminar speaker)
Cornell University, Ithaca, NY
- 2013 SPHHP Online Teaching Symposium
University at Buffalo, School of Public Health
- 2012 What can causal networks teach us about metabolism?
University at Buffalo, Pharmaceutical Sciences and Neurology.
- 2011 Modeling Complex Biological Systems.
NC State, Department of Genetics, Raleigh, NC
- 2011 Toward the in silico cell: Alliance Workshop
Case Western Reserve University, Cleveland, OH.
- 2011 Markov Chains and Blind Deconvolution
University at Buffalo, Department of Physiology and Biophysics.
- 2011 Modeling Complex Biological Systems
Roswell Park Cancer Institute, Department of Molecular and Developmental Genetics.
- 2009 Bayesian Networks in Biology
Humboldt University, School of Agriculture and Horticulture. Berlin, Germany.

Professional Service and Development

External Professional Service

Reviewer for: *AWM Edge Program, Bioinformatics, BMC Medical Research Methodology, BMC Systems Biology, British Journal of Applied Science and Technology, CRC Press, Taylor & Francis Group: Applications in Data Mining Techniques for Oral Cancer Detection and Prevention, IEEE/ACM Transactions on Computational Biology and Bioinformatics, Journal of the American Statistical Association, Molecular BioSystems, Pacific Symposium on Biocomputing, Physiological Genomics, PLoS One, PLoS Computational Biology and Science.*

- 2019-2021 Program Committee NERCCS 2020
- 2011-2012 Review Editor *Frontiers in Genomic Physiology*
- 2013-2017 Associate Editor *Advances in Systems Biology*
- 2016 -2022 Review Panel NSF GRFP (Graduate Research Fellowship Program)
- 2015- 2020 Review Panel NSF DMS Mathematical Biology Program
- 2014 -2017 Early Career Reviewer Program – NIH Center for Scientific Review
- 2016 Review Panel DoD Extramural Medical Research: Combat Casualty Care

2015, 2021 Review Panel NSF/NIGMS Interface of the Biological and Mathematical Sciences
 2017 Ad hoc reviewer for Center for CTSI Penn State Pilot Study program
 2017 Review Panel NSF/Simons Research Foundation - Centers for Mathematics of
 Complex Biological Systems
 2018 Reviewer NSF PGRP Program
 2019 Ad hoc reviewer BMRD study section
 2019 Review Panel NSF CAREER Mathematical Biology Program

Panel Invitations declined:

2017 Review Panel DoD Therapeutics and Treatment Gulf War Illness program
 2018 Review Panel DoD Detection and Diagnosis program

UB Internal Professional Service

2021-present Search Committee, Chair Position for UB Department of Biostatistics
 2021-present Research Initiatives Committee, UB School of Public Health and Health Professions
 2019-present Tenure and Promotions Committee, UB School of Public Health and Health
 Professions
 2019-present Director MPS Data Science, UB Institute for Computational and Data Science
 2018-present CDSE Days Organizing committee
 2019 Undergraduate Curriculum committee, UB Department of Biostatistics
 2019 Research Assistant Professor search committee, UB Department of Biostatistics
 2019 Data Science and Engineering MS and MPS admissions committee
 2018-2019 Seminar Organizer, UB Department of Biostatistics
 2018-2019 Search Committee Research Assistant Professor, UB Department of Biostatistics
 2018 UB Institute for Computing and Data Sciences committee - SPHHP representative
 2017-2018 Faculty Search Committee UB Department of Chemical and Biological Engineering
 2016-2017 Faculty Search Committee UB Department of Chemical and Biological Engineering
 2016-2017 Search Committee for the Chair of UB Department of Biostatistics
 2015-present Admissions Committee, UB Department of Biostatistics
 2015-present Computational Data Enabled Science and Engineering, participating faculty.
 2012-present Genetics, Genomics, and Bioinformatics Program, participating faculty.
 2012-2017 Department of Biostatistics vision statement committee.
 2012-2014 Faculty council
 2013-2015 Communications committee
 2016 UB IMPACT internal funding, review panel for the life sciences.
 2016 UB GEM Biostatistics consultation panel for pilot projects
 2016 UB IMPACT Biostatistics consultation panel for pilot projects
 2013-2016 Biostatistics graduate admissions committee
 2018
 2013-2014 Faculty Search Committee Biostatistics/Epidemiology and Environmental Health
 2014 SPHHP SPEC Committee
 2014 Perry poster day judge
 2015 Faculty search committee Biostatistics

Community Service

2019 Letters to a pre-scientist “pen pal” program
 2019 Inheritance of complex trait through dragon breeding – Sweet Home high school
 2016-2018 Mind your microbiome week, Buffalo Public Schools – Sponsored by UB GEM
 2011-2017 Faculty Mentor – Association for Women in Mathematics (AWM)
 2016 Genome Day 2016 for Buffalo Public Schools – Sponsored by UB GEM

2015 Faculty Mentor – UB LSAMP undergraduate research program for minorities

Professional Memberships

2021-present Society of Classification

2019-present American Mathematical Society (AMS)

2012-present American Statistical Association (ASA)

2008-present International Society for Computational Biology (ISCB)

2006-2017 Association for Women in Mathematics (AWM)

2010-2012 American Association for the Advancement of Science (AAAS)

Development (led) and synergistic teaching activities

2016-2020 Guest lecture “statistical considerations for the analysis of omics data”

UB Department of Pharmaceutical Sciences

2013-2017 Introduction to Pathway Analysis (annual lecture)

Roswell Park Cancer Institute, graduate program in Molecular Biophysics (BPR 503)
graduate course in Pharmacogenomics in pharmaceutical science (PHC 509)

2015 Introduction to R programming workshop (with Drs. Gaile and Miecznikowski)

UB Computational and Data Enabled Science (CDSE) Days

2014 Bioinformatics guest lecture (STA 525), Department of Biostatistics

2008-2010 Short Course on Systems Genetics (Formerly short course on Complex Traits)

Guest lectures and computer lab development.

The Jackson Laboratory, guest lecture and computer lab development..

Development (attended) and continuing education and training

2019 UB ICDS, Jupyter and OpenEd workshop

2018 AMIA workshop – introduction to jupyter notebooks for data science

2011-2017 UB Biostatistics, Applied genomics journal club/reading group

2016 International Genetic Epidemiology Society (IGES) Big data phenotyping workshop

University of Toronto, Toronto, Canada.

2016 UB Workshop: Network on enriched academic relationships and mentoring

2016 UB CEI Workshop: How to help students learn (to work) in teams

2012 UB Online Teaching Training Course

2008-2011 Annual Short Course on Systems Genetics,

The Jackson Laboratory, Bar Harbor ME.

2007 High Dimensional Statistics in Biology

Isaac Newton Institute for Mathematical Sciences, Cambridge, UK,

2006 Mathematical Models of the Heart

European Mathematical Society, Svalbard, Norway.

2006 Cardiac Mechanics and Remodeling Workshop

Mathematical Biosciences Institute, Columbus, OH.

2004 Statistical and Numerical Methods for Inverse Problems Workshop

University of Bologna, Italy.

2005 Image Processing Workshop

Mathematical Sciences Research Institute, Berkeley, CA.

Mentoring

Masters-level students committee chair

2020 Rachael Tylock, MS Bioinformatics and Biostatistics

2018 Shuo Qian, MS Bioinformatics and Biostatistics
 2018 Manjit Kaur, MS Bioinformatics and Biostatistics
 2018 Sindhuja Sridharan, MS Bioinformatics and Biostatistics
 2018 Lina Spokauskaite, MA Biostatistics
 2017 Krithika Krishnan, MS Bioinformatics and Biostatistics
 2017 Suruchi Jaikumar Ahunja, MS Bioinformatics and Biostatistics
 2017 Aishwarya Mandava, MS Bioinformatics and Biostatistics
 2017 Rui Cheng, MS Bioinformatics and Biostatistics
 2017 Krithika Krishnan, MS Bioinformatics and Biostatistics
 2016 Nan Nan, MA Biostatistics
 2015 Joselle O'Brien, MA Biostatistics
 2015 Janhavi Moharil, MA Biostatistics
 2015 Fang Hao, MA Biostatistics
 2015 Aaron Crowley, MA Biostatistics
 2015 Matthew Miller, MA Biostatistics

Ph.D. students committee chair

2017 Han Yu, Biostatistics
 In progress Tiange Shi, Biostatistics
 in progress Krithika Krishnan, Computational and Data-Enabled Science and Engineering
 in progress Tianmou Liu, Computational and Data-Enabled Science and Engineering

Masters-level students committee membership

2020 Zachary Wintrob, MA Biostatistics
 2020 Ki Chen, MPH Biostatistics concentration
 2020 Hua-Hsin, MA Biostatistics
 2019 Yufei Wu, MA Biostatistics
 2019 Chang Xu, MA Biostatistics
 2019 Justin Mu, MA Biostatistics
 2018 Kaifeng Yu, MA Biostatistics
 2017 Jieya Lin, MA Biostatistics
 2017 Zeyu Yang, MA Biostatistics
 2017 Jiangwang Chen, MA Biostatistics
 2016 Emily Schiller, MS Bioinformatics and Biostatistics
 2016 Kayla Morrell, MS Bioinformatics and Biostatistics
 2016 Sarah Kauss, MA Biostatistics
 2015 Michael Edinger, MPH
 2015 Mark Heiler, MA Biostatistics
 2014 Jie Gong, MA Biostatistics
 2013 Darryl George, MA Biostatistics
 2013 Chunyuan Diao, MA Biostatistics
 2013 Carolyn Horwitz, MPH
 2013 Jeffery Rathbun, MA Biostatistics

Ph.D. students committee membership

2021 Lorin Miller, Department of Biostatistics
 (advisor: Dr. Jeff Miecznikowski)

 2020 Zeinab Farhat, Epidemiology and Environmental Health
 (advisor: Dr. Lina Mu)

- 2019 Fan Zhang, Biostatistics
(advisor: Dr. Jeffrey Miecznikowski)
- 2019 Boris Boutkov, Computational and Data-Enabled Science and Engineering
(advisor: Dr. Paul Bauman)
- 2019 Vineet Payappalli, Industrial and Systems Engineering
(advisor: Dr. Jun Zhuang)
- 2018 Michael Vaiana, Computational and Data-Enabled Science and Engineering
(advisor Dr. Sarah Muldoon)
- 2017 Dr. Laurie Griesenhofer, Epidemiology & Environmental Health
(advisor: Dr. Heather Ochs-Balcom)
- 2015 Dr. Iakovos Toumazis, Industrial & Systems Engineering PhD
(advisor: Dr. Changyun Kwon)
- in progress, Renette Jones, Computational and Data-Enabled Science and Engineering
(advisor: Dr. Abani Patra)
- in progress Kexin Zhu, Epidemiology and Environmental Health
(advisor: Dr. Lina Mu)

Summer/Short Term Mentoring

- 2018 Eric Kieshel, UB Genetics, Genomics and Bioinformatics Program -4 week lab rotation
- 2016 Renee Meinhold, RIT undergraduate mentored with Dr. Susan McCann (Roswell Park)
- 2015 Gabriel Skugor, Department of Mathematics, LSMAP undergraduate summer research
- 2012 Laurie Griesenhofer, formerly in the Department of Mathematics

Research Support

Ongoing Research Support

- US Army Medical Research 888358/ Ramanathan (PI) 09/2020-08/2022
Project title: *"The metabolomics-neurofilaments-neurodegeneration nexus in multiple sclerosis"*
This project examines the associations between diet, metabolism and neurodegeneration..
Role on Project: Co-I
- BrightFocus Foundation M2020227 / Millen (PI) 09/2020-08/2022
Project title: *"Interplay of diet and gut microbiome in age related macular degeneration"*
This project examines the diet and gut microbiome of patients in various stages of AMD.
Role on Project: Co-I
- NIH/NIAID R21AI134370 / Ambruster & Ghamen (PI) 06/2020-05/2022
Project title: *Genetic susceptibility to mucosal infections with aging*
This project examines genetics of mucosal infections with aging in collaborative cross mice.
Role on Project: Co-I
- NIH/NIMH R01MH118218 / Gainey (PI) 12/2019-05/2024
Project title: *"Emotion regulation in distress disorders: elucidating the role of cognitive processes and person-situation fit in the laboratory and daily life"*
This project examines mood interventions and the impact on daily life and happiness
Role on Project: Co-I
- NIH/NIDDK R01 DK123158 / Ambruster (PI) 09/2019-08/2023

Project title: *"Conserved Proteus mirabilis genetic requirements for colonization of the catheterized urinary tract"*

This project leverages examines genetic and clinical factors related to UTIs.

Role on Project: Co-I

NCI: T32 in cancer epidemiology / Freudenheim and Moyish (PIs) 09/2017-08/2022

This grant is collaboration between the University of Buffalo and Roswell Park Cancer Institute for the training of graduate students and postdoctoral fellows using interdisciplinary mentoring and curriculum.

Role on Project: training mentor

Completed Research Support

NIH/NIGMS R01 GM073646 Blanco (PI) 08/2019-07/2020

Project title: *"Defining the Contribution of Cellular Hypoxia to the Cardiotoxicity of Anticancer Anthracyclines and Trastuzumab"*

This project examines a variety of data related to modeling the pharmacodynamics of anthracycline drugs in down-syndrome and healthy patients.

Role on Project: Co-I

NIH/NIEHS R21 ESO26429/ Mu (PI) 08/2016-07/2020

Project title: *"Metabolomics profiling of biological responses to air pollution"*

This project leverages metabolomics profiling via Mass Spectrometry to assess exposure to air pollution.

Role on Project: Co-I

UB GEM Educational and Engagement Innovation (PI: Hageman Blair) 09/2018-12/2019

Project title: *"Introduction to modern genetics through dragon breeding in the high-school classroom"*

This project supports the design and implementation of an outreach workshop that partners with Sweet Home High School (title 1).

Role on Project: PI

UB: Center of Excellence /Nowak and Surtees (PIs) 06/2015-08/2018

Project title: *The Genome, Environment, and Microbiome.*

This project integrates investigators across disciplines at UB for the purpose of examining the genome, environment, and microbiome, and the complex interplay between them.

Role on Project: Co-I

NIH/NIGMS 2R01GM073646 Blanco (PI) 03/2015-02/2019

Project title: *"Contribution of CBRs and AKRs to the Pharmacodynamics of Anthracycline Drugs"*

This project examines a variety of data related to modeling the pharmacodynamics of anthracycline drugs in down-syndrome and healthy patients.

Role on Project: Co-I

UB CEI Seed Grant (PIs: Hageman Blair & Gaile) 06/2017-12/2018

Project title: *"Integrative learning through the data-driven classroom: the DREAM challenge"*

This project develops an interdisciplinary new course that leverages integrative learning for the ISCB DREAM challenge.

Role on Project: PI

NSF/DMS-1557589 / Hageman Blair (PI) 09/2015- 08/2017
Project title: "*QuBBD: Collaborative Research: Interactive ensemble clustering for mixed data with application to mood disorders*"

This planning grant is a joint initiative between NIH and NSF to advance methodologies related to biomedical big data. Related awards for collaborators: DMS-1557642, DMS-1557593, DMS-1557576, and DMS-1557668, total amount \$100,000.

Role on Project: PI

NSF DMS-1312250/ Hageman Blair (PI) 08/2013-07/2017
Mathematical Biology

Project title: "*Integrated Bayesian deterministic and probabilistic modeling of phenotype-genotype networks in the metabolic system*".

This project examines genotype-phenotype networks and metabolic systems, their integration, with applications to disease phenotypes.

Role on Project: PI

UB: IMPACT funding/ Millen (PI) 06/2015-08/2016

Project title: "*The neonatal microbiome study: a pilot study in Meru County; Kenya*"

This project examines microbiome sample collection method to determine field feasibility. The project was discontinued due to pulled funding on a parent study.

Role on Project: Co-I

UB: IMPACT funding/ Millen (PI) 09/2016-08/2017

Project title: "*Role of the gut microbiome in age-related macular degeneration*"

This project is a pilot study for mail-based stool collection in aging women

Total amount of award: \$23,682

Effort: n/a (faculty salary not an allowable cost)

Role on Project: Co-I

UB CTSI/ Hageman Blair, Mieczkowski, Odunsi, Szyperski (PIs) 11/2015-03/2017

Project title: "*Multiclass modeling of metabonomics data for the detection of early stage ovarian cancer*"

This project develops predictive models from metabonomics profiles and biomarkers for ovarian cancer. My role was to develop the predictive models.

Total amount of award: \$75,000

Effort: n/a (faculty salary not an allowable cost) and partial graduate student support

Role on Project: PI

UB/STOR-1111511/ Hageman Blair and Szyperski (PIs) 05/2014-08/2014

Project title: "*Predictive models to diagnose ovarian cancer*"

This project provided seed money for research development for predictive modeling of ovarian cancer using metabonomics data. The project supported two summer graduate students in Biostatistics to aid in the development of the statistical models.

Total amount of award: \$35,000

Effort: n/a (faculty salary not an allowable cost)

Role on Project: PI

F32 NIH/NHLBI HL095240/ Hageman (PI) 12/2008-08/2011

Project title: “*Bayesian dynamic genome scale modeling of HDL cholesterol transport*”
This project developed mathematical models to understand HDL metabolism in *mus musculus*.
Total amount of award: \$150,000
Effort: 100%
Role on Project: PI