Jean Morrison

Postdoctoral Scholar University of Chicago Department of Human Genetics 520.548.3773 morrison.jeanv@gmail.com

Education

2016 PHD BIOSTATISTICS, University of Washington

Advisor: Noah Simon, Department of Biostatistics

2009 BA MATHEMATICS, University of Chicago

Work Experience

DEPARTMENT OF BIOSTATISTICS, UNIVERSITY OF WASHINGTON

12.2013- Research Assistant

8.2016 Supervisor: Noah Simon

Statistical methods for spatially structured genomic data and high dimensional studies.

6.2014–9.2015 Research Assistant, Genetic Analysis Center

Supervisor: Cathy Laurie

Genome wide association analysis of blood cell and dental traits for the Hispanic Community Health Study/ Study of Latinos, a large multi-ethnic study of Latino

populations.

6.2011–6.2014 Research Assistant

Supervisor: Bruce Weir and Tim Thornton

Investigations of population structure and its effects in genetic association studies.

DEPARTMENT OF MEDICINE, SECTION OF GENETICS, UNIVERSITY OF CHICAGO

7.2009-6.2011 Research Assistant

Supervisor: Nancy Cox

Genome wide association analysis, quality control and imputation for the Hyper-glycemia and Adverse Pregnancy Outcomes (HAPO) study. QC and programming

support for other genetic analysis projects.

Honors, Awards

2016	ASA Section on Genetics and Genomics Student Paper Award				
	University of Washington School of Public Health Gilbert S. Omenn Award for Academic				
2016	Excellence				
	Ruth L. Kirschstein Predoctoral Individual National Research Service Award, National				
2016	Institutes of Health				
2011-2014	Biostatistics Statistical Genetics Training Grant, National Institutes of Health				

Publications

- Morrison, Jean, Daniela Witten, and Noah Simon (2016). "Joint Adaptive Differential Estimation: A tool for comparative analysis of smooth genomic data types." *Biostatistics* In press.
- Schick, Ursula M., Deepti Jain, Chani J. Hodonsky, Jean V. Morrison, James P. Davis, Lisa Brown, et al. (2016). "Genome-wide Association Study of Platelet Count Identifies Ancestry-Specific Loci in Hispanic/Latino Americans". English. *The American Journal of Human Genetics* 98, pp. 229–242.
- Morrison, Jean, Cathy C Laurie, Mary L Marazita, Anne E Sanders, Steven Offenbacher, Christian R Salazar, et al. (2015). "Genome-wide association study of dental caries in the Hispanic Communities Health Study/Study of Latinos (HCHS/SOL)." *Human Molecular Genetics* 25.4, pp. 807–816.
- Hayes, M. Geoffrey, Margrit Urbanek, Marie France Hivert, Loren L. Armstrong, Jean Morrison, Cong Guo, et al. (2013). "Identification of HKDC1 and BACE2 as genes influencing glycemic traits during pregnancy through genome-wide association studies". *Diabetes* 62.9, pp. 3282–3291.
- Morrison, Jean (2013). "Characterization and correction of error in genome-wide ibd estimation for samples with population structure". *Genetic Epidemiology* 37.6, pp. 635–641.
- Urbanek, Margrit, M. Geoffrey Hayes, Loren L. Armstrong, Jean Morrison, Lynn P. Lowe, Sylvia E. Badon, et al. (2013). "The chromosome 3q25 genomic region is associated with measures of adiposity in newborns in a multi-ethnic genome-wide association study". *Human Molecular Genetics* 22.17, pp. 3583–3596.
- Below, J. E., E. R. Gamazon, J. V. Morrison, A. Konkashbaev, A. Pluzhnikov, P. M. McKeigue, et al. (2011). "Genome-wide association and meta-analysis in populations from Starr County, Texas, and Mexico City identify type 2 diabetes susceptibility loci and enrichment for expression quantitative trait loci in top signals". *Diabetologia* 54.8, pp. 2047–2055.

Presentations

- Morrison, Jean, Noah Simon, and Daniela Witten (2016). "Simultaneous Detection and Estimation of Trait Associations with Genomic Phenotypes". In: *Joint Statistical Meetigs*. Chicago, IL.
- Morrison, Jean, Richard Sandstrom, and Noah Simon (2015). "Locally adaptive comparison of DNase I profiles to detect fine scale differences in regulatory activity." In: *American Society for Human Genetics Meetings*. Baltimore, MD.
- Morrison, Jean, Daniela Witten, and Noah Simon (2014). "JADE: A tool for comparative analysis of spatially smooth genomic data." In: *American Society for Human Genetics Meetings*. San Deigo, CA.
- Morrison, Jean (2012). "Overestimation of relatedness in admixed and ancestrally heterogeneous populations using method of moments estimation." In: *American Society for Human Genetics Meetings*. San Francisco, CA.
- Morrison, Jean, DA Scheftner, A Pluzhnikov, LP Lowe, C Ackerman, L Armstrong, et al. (2011). "SNPs near CCNL1 associated with infant adiposity in multi-ethnic mega-analysis with strongest effect observed in European ancestry newborns". In: *International Congress on Human Genetics meeting*. Montreal, Quebec.
- Morrison, Jean, Anna Pluzhnikov, MG Hayes, H Lee, David Levine, Caitlin McHugh, et al. (2010). "G6PC2 associated with fasting glucose levels in pregnant women of European ancestry." In: *American Society for Human Genetics Meetings*.

R Packages

jadeTF: https://github.com/jean997/jadeTF

Teaching Experience

	Fall 2014	Teaching Assisitant,	BIOST 536: Categorica	al Data Analysis in	Epidemiology
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Supervisor: Scott Emerson

Spring 2014 Teaching Assisitant, BIOST 540: Correlated Data Analysis

Supervisor: Ken Rice

9.2008-5.2009 Teaching Assistant for SESAME Algebra, University of Chicago

Algebra for middle grade teachers.

9.2006-9.2007 Young Scholars Program Counselor/Teaching Assistant, University of Chicago

Math enrichment program for middle schoolers.

Professional Service

9.2015–Now *Member*, Biostatistics Department Curriculum Committee

9.2014–5.2015 Member, Biostatistics Department Student, Faculty Relations Committee