ALEXANDRE AMLIE-WOLF

https://orcid.org/0000-0002-2073-1519 https://alexamlie.github.io alexaml [at] upenn.edu

EDUCATION

University of Pennsylvania, Philadelphia, Pennsylvania, USA.

2013-Present

Ph.D. Genomics and Computational Biology, GPA: 3.86

Thesis Advisor: Li-San Wang

Oberlin College, Oberlin, Ohio, USA.

2009-2013

B.A. Computer Science with High Honors and B.A. Neuroscience, Phi Beta Kappa

GPA: Computer Science 3.93, Neuroscience 3.7, Cumulative 3.8

Thesis: "A Swarm of Salesmen: Algorithmic Approaches to Multiagent Modeling"

http://rave.ohiolink.edu/etdc/view?acc_num=oberlin1368052652

Thesis Advisor: Tom Wexler

PUBLICATIONS

*: co-first author

Manuscripts in preparation

- 1. **Amlie-Wolf A***, Wang W*, ..., Wang LS, Schellenberg GD. Progressive supranuclear palsy and genomic features of the chromosome 17 MAPT region.
- 2. Kuksa P*, **Amlie-Wolf A***, Hwang YC*, Gregory BD, Wang LS. Identifying the transcription factors mediating enhancer target gene regulation in the human genome.

Manuscripts under review

- 1. Amlie-Wolf A*, Tang M*, Way J, Dombroski B, Jiang M, Vrettos N, Chou YF, Zhao Y, Kuzma A, Mlynarski EE, Brown CD, Wang LS, Schellenberg GD. Inferring the molecular mechanisms of noncoding Alzheimer's disease-associated genetic variants. bioRxiv 2018, doi: 10.1101/401471. https://www.biorxiv.org/content/early/2018/08/27/401471
- 2. Kunkle BW, Grenier-Boley B, Sims R, Bis JC, Naj AC, Boland A, Vronskaya M, Lee SJ van der, **Amlie-Wolf A**, Bellenguez C, ..., Schellenberg GD, Lambert JC, Pericak-Vance MA (2018) Meta-analysis of genetic association with diagnosed Alzheimers disease identifies novel risk loci and implicates Abeta, Tau, immunity and lipid processing. bioRxiv, doi: 10.1101/294629. https://www.biorxiv.org/content/early/2018/04/05/294629
- 3. Wheeler JM*, McMillan P*, Strovas JT*, Liachko NF, **Amlie-Wolf A**, ..., Schellenberg GD, Kraemer B (2018) The poly(A) binding protein MSUT2 controls resistance to both pathological tau and gliosis.

Journal Articles

- Kuksa PP, Amlie-Wolf A, Katanic Z, Valladares O, Wang LS, and Leung YY (2018). DASHR 2.0: integrated database of human small non-coding RNA genes and mature products. Bioinformatics. doi: 10.1093/bioinformatics/bty709
- 2. Amlie-Wolf A, Tang M, Mlynarski EE, Kuksa PK, Valladares O, Katanic Z, Tsuang D, Brown CD, Schellenberg GD, Wang LS. (2018) INFERNO: INFERring the molecular mechanisms of NOncoding genetic variants. Nucleic Acids Research, doi: 10.1093/nar/gky686
- 3. Kuksa PP, **Amlie-Wolf A**, Katanic Z, Valladares O, Wang LS. and Leung YY (2018) SPAR: small RNA-seq portal for analysis of sequencing experiments. Nucleic Acids Research, 46, W36W42. doi: 10.1093/nar/gky330

- 4. Nativio R, Donahue G, Berson A, Lan Y, **Amlie-Wolf A**, Tuzer F, Toledo J, Gosai S, Gregory B, Torres C, Trojanowski J, Wang LS, Johnson FB, Bonini N, Berger S. (2018) Dysregulation of the epigenetic landscape of normal aging in Alzheimer's disease. Nature Neuroscience. doi: 10.1038/s41593-018-0101-9
- 5. Leung YY, Kuksa P, **Amlie-Wolf A**, Valladares O, Ungar L, Kannan S, Gregory B, Wang LS. (2016) DASHR: database of small human noncoding RNAs. Nucleic Acids Research. doi: 10.1093/nar/gkv1188
- 6. Amlie-Wolf A, Ryvkin P, Tong R, Dragomir I, Suh E, Van Deerlin VM, Gregory BD, Kwong LK, Trojanowski JQ, Lee VM, Wang LS, Lee EB. (2015) Transcriptomic Changes Due to Cytoplasmic TDP-43 Expression Reveal Dysregulation of Histone Transcripts and Nuclear Chromatin. PLoS ONE 10(10): e0141836. doi: 10.1371/journal.pone.0141836

Published Conference Abstracts

- Amlie-Wolf A, Tang M, Way J, Dombroski BA, Vrettos N, Chou YF, Mlynarski EE, Brown CD, Wang LS, Schellenberg GD. (2018) Inferring the Molecular Mechanisms of Noncoding AD-Associated Genetic Variants. Alzheimer's & Dementia: The Journal of the Alzheimer's Association.
- 2. Amlie-Wolf A, Tang M, King J, Dombroski BA, Wang LS, Schellenberg GD. (2016) Computational identification of regulatory mechanisms affected by noncoding variants associated with late-onset Alzheimer's disease. Alzheimer's & Dementia: The Journal of the Alzheimer's Association 12(7), P640-P641.

PRESENTATIONS

Invited and Conference Talks

1. American Society of Human Genetics Conference Platform Talk	10/20/2018 San Diego, CA
2. Alzheimer's Disease Genetics Consortium University of Pennsylvania	01/30/2018 $Philadelphia, PA$
3. Genomics and Computational Biology Retreat University of Pennsylvania	06/08/2017 Philadelphia, PA
4. Alzheimer's Disease Genetics Consortium Case Western Reserve University	02/16/2017 Cleveland, OH

Refereed Conference Posters

- 1. Amlie-Wolf A, Qu L, Mlynarski EE, Brown CD, Schellenberg DG, Wang LS. (2017) The regulatory landscape of genetic variants associated with psychiatric disorders and neurodegenerative diseases. The 67th Annual Meeting of the American Society of Human Genetics, Oct 17-21, Orlando, Florida.
- 2. Kuksa PP, Leung YY, **Amlie-Wolf A**, Valladares O, Wang LS. (2017) DASHR 2.0 database of small non-coding RNAs in normal human tissues and cell types. The 67th Annual Meeting of the American Society of Human Genetics, Oct 17-21, Orlando, Florida.
- 3. Leung YY, Kuksa PP, **Amlie-Wolf A**, Wang LS. (2017) The landscape of short RNAs in human cell types and tissues. The 67th Annual Meeting of the American Society of Human Genetics, Oct 17-21, Orlando, Florida.
- 4. Amlie-Wolf A, Tang M, King J, Dombroski BA, Chou YF, Mlynarski EE, Schellenberg DG, Wang LS. (2017) Integrative analysis identifies immune-related enhancers and lncRNAs perturbed by genetic variants associated with Alzheimer's disease. UPenn Institute on Aging Retreat, May 23, Philadelphia, Pennsylvania.

- 5. Amlie-Wolf A, Tang M, Kuksa PP, Leung YY, Slaff B, King J, Dombroski BA, Schellenberg DG, Wang LS. (2016) INFERNO INFERring the molecular mechanisms of NOncoding genetic variants. The 66th Annual Meeting of the American Society of Human Genetics, Oct 18-22, Vancouver, Canada and MidAtlantic Bioinformatics Conference, Oct 26, Philadelphia, PA, USA.
- 6. Leung YY, Kuksa PP, Amlie-Wolf A, Wang LS. (2016) The landscape of regulatory post-transcriptionally derived small noncoding RNAs in the human transcriptome. The 66th Annual Meeting of the American Society of Human Genetics, Oct 18-22, Vancouver, Canada.
- Amlie-Wolf A, Tang M, King J, Dombroski BA, Wang LS, Schellenberg GD. (2016) Computational identification of regulatory mechanisms affected by noncoding variants associated with lateonset Alzheimer's disease. Alzheimer's Association International Conference, July 24-28, Toronto, Canada.
- 8. Kuksa PP, Leung YY, **Amlie-Wolf A**, Gregory BD, Wang LS. (2015) SPAR Sequencing-based pipeline for annotating novel small non-coding RNAs. The 65th Annual Meeting of the American Society of Human Genetics, Oct 6-10, Baltimore, USA.
- 9. Leung YY, Kuksa PP, **Amlie-Wolf A**, Gregory BD, Wang LS. (2015) DASHR Database of small human noncoding RNA. The 65th Annual Meeting of the American Society of Human Genetics, Oct 6-10, Baltimore, USA.
- 10. Freedman J, **Amlie-Wolf A**, Wittenberg R, Shoham O, Aronson S, Loose MD. (2014) Identifying the relative influence of multiple prior events on predictions of a probabilistic future: An artificial neural network analysis. Society for Neuroscience 2014, Nov 15-19, Washington D.C., USA.
- 11. Aronson S, **Amlie-Wolf A**, Loose MD. (2012) Making predictions in a stochastic environment: Strategies underlying the probability matching phenomenon and corresponding event-related potentials. Society for Neuroscience 2012, Oct 13-17, New Orleans, USA.
- 12. Loose MD, Aronson S, Scott H, Mehta T, Wang A, **Amlie-Wolf A**, Welch K. (2011) Salience is not always salient: Neither magnitude nor valence of expected outcome contributes to the N2/P3 event-related potential complex in a Go/NoGo task. Society for Neuroscience 2011, Nov 12-16, Washington D.C., USA.

ACADEMIC AND PROFESSIONAL HONORS

2017: 1st place poster award in basic science category at UPenn Institute on Aging Retreat

2015 - Present: Predoctoral Trainee in Age Related Neurodegenerative Diseases, NIH/NIA T32 AG00255

2013: Nancy Robell Prize in Neuroscience, Oberlin College

2013: Graduated with High Honors in Computer Science

2013: Inducted into Phi Beta Kappa

2009-2013: John F. Oberlin Merit Scholarship

2009-2013: National Merit Scholarship Finalist

2005-2009: Lower Merion High School Honor Roll

RESEARCH EXPERIENCE

Graduate Student January 2014-Present

Department of Pathology and Laboratory Medicine, University of Pennsylvania

Advisor: Li-San Wang Philadelphia, PA

Developing integrative computational methods to characterize the regulatory effects of noncoding genetic variation in Alzheimer's disease and related neurodegenerative diseases

Rotation Student March 2014-June 2014

Advisor: Edward B. Lee Philadelphia, PA

Rotation Student September 2013-December 2013

Advisor: Yoseph Barash Philadelphia, PA

Computer Science Honors Student Department of Computer Science, Oberlin College	Fall 2012-Spring 2013
Advisor: Tom Wexler	$Oberlin,\ OH$
DAAD RISE Scholar Department of Computer Science, Humboldt-Universität zu Berlin	May 2012-August 2012
Advisors: Michael Frey, Mesut Günes, Joachim Fischer	Berlin, Germany
Research Assistant Department of Neuroscience, Oberlin College Advisor: Michael Loose	Spring 2011-Spring 2013
	Oberlin, OH
Research Assistant Department of Neuroscience, Oberlin College	Fall 2011-Spring 2012
Advisor: Keith Downing	$Oberlin,\ OH$
Summer Undergraduate Research Assistant	Summer 2011
Center for Neurodegenerative Disease Research, University of Pennsylvania Advisors: Virginia Lee and Li-San Wang	$Philadelphia,\ PA$
Summer Undergraduate Research Assistant	Summer 2010
Center for Neurodegenerative Disease Research, University of Pennsylvania Advisor: Virginia Lee	$Philadelphia,\ PA$
ACADEMIC SERVICE	
Member of committee to update individual development plans (II Biomedical Graduate Studies, University of Pennsylvania	DP) 2016-2017 Philadelphia, PA
Reviewer for APBC and BIBM Conferences	2017-2018
Organizer for student-invited Penn Bioinformatics Forum talks Institute for Biomedical Informatics, University of Pennsylvania	Fall 2014-Spring 2016 Philadelphia, PA
TEACHING EXPERIENCE	
TA and course development for GCB535: Introduction to Bioinfo Biomedical Graduate Studies, University of Pennsylvania	rmatics Spring 2016 Philadelphia, PA
Tutor for BIOM555: Regulation of the Genome Biomedical Graduate Studies, University of Pennsylvania	Spring 2015 Philadelphia, PA
Grader for Computer Science 275: Programming Abstractions Department of Computer Science, Oberlin College	Spring 2013 Oberlin, OH
Grader for Computer Science 151: Principles of Computer Science Department of Computer Science, Oberlin College	e II Fall 2012 Oberlin, OH
Grader for Computer Science 150: Principles of Computer Science Department of Computer Science, Oberlin College	e I Spring 2012 Oberlin, OH
Tutor for Computer Science 151: Principles of Computer Science Department of Computer Science, Oberlin College	II Fall 2011 Oberlin, OH
RESEARCH SKILLS	

Programming languages:

Strong - R, Python, bash, Java, C/C++, LATEX

Moderate - Mathematica, Assembly, Perl, MATLAB, Scheme

Wet lab skills:

Molecular cloning, cell culture, transient transfection, luciferase assay, protein biochemistry

SOCIETY MEMBERSHIPS

American Society of Human Genetics, Alzheimer's Association, Phi Beta Kappa (Zeta of Ohio Chapter), Sigma Xi, The Pledge of the Computing Professional, Society for Neuroscience

EXTRACURRICULAR ACTIVITIES

Avid Musician: Oboe, Bass Guitar, Guitar, Piano, Drums, Vocals, Double Bass, Mandolin.

https://alexamliewolf.bandcamp.com https://soundcloud.com/alexamlie/

https://www.youtube.com/c/AlexAmlieWolf

'Honorary' Member of Oberlin Conservatory Oboe Studio, played with the Oberlin Chamber Orchestra (usually open only to Conservatory music students, extremely rare for a wind player)

Member of the Competitive Computer Programming Club at Oberlin, competed at the Denison Programming Contest and the ACM International Collegiate Programming Contest

Languages: English (native), French (conversational)