# Sean Whalen, Ph.D.

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#### **Current Position**

2017- Research Scientist, Gladstone Institutes, University of California, San Francisco

#### Research Interests

Machine Learning; Computational Biology & Bioinformatics; Computer Security; Cloud & High Performance Computing; Data Visualization

### **Programming Languages**

Python (numpy/scipy/pandas/scikit-learn), R (caret/dplyr/ggplot), Matlab, Java, Scala, C, C++, CUDA, OpenGL, HTML/CSS/JavaScript

### Appointments

2013-2016	Biostatistician, Gladstone Institutes, University of California, San Francisco
2012-2013	Postdoctoral Fellow, Institute for Genomics and Multiscale Biology, Mount Sinai School of
	Medicine, Advisor: Gaurav Pandey
2011-2012	Postdoctoral Researcher, DARPA Mission-oriented Resilient Clouds, Intrusion Detection
	Systems Lab, Columbia University, Advisor: Salvatore J. Stolfo
2010-2011	I3P Postdoctoral Fellow, Computational Research Division, Lawrence Berkeley Natl. Lab
2010	Researcher, Anomaly Detection in High Performance Computing, UCDavis Security Lab
2009	Teaching Assistant, Operating Systems, Instructor: Daniela Oliviera
2008-2009	Researcher, Immersive Network Visualization, William M. Keck Foundation
2007	Teaching Assistant, Artificial Intelligence, Instructor: Ian Davidson
2006	Teaching Assistant, Operating Systems, Instructor: Felix Wu
2006	Teaching Assistant, Introduction to Object Oriented Programming, Instructor: Sean Davis
2004-2006	Researcher, Vulnerability Analysis, UCDavis Security Lab

#### Education

2010	Ph.D. in Computer Science, University of California, Davis
	Advisors: Matt Bishop (Computer Security) and James P. Crutchfield (Physics)
2006	M.S. in Computer Science, University of California, Davis
2002	B.S. in Computer Science (Cum Laude), University of Nebraska, Omaha

#### Grants, Honors & Awards

2013	Center for Technology, Innovation, and Entrepreneurship Grant Recipient
2010-2011	Institute for Information Infrastructure Protection Postdoctoral Fellowship (\$150k)
2003	Graduate Assistance in Areas of National Need Doctoral Fellowship (Tuition)
1998-2002	Walter Scott Jr. Scholarship, University of Nebraska Foundation (Tuition, Board)

#### Peer-Reviewed Publications

#### **Papers**

2017	"Genomic analyses for age at menarche identify 389 independent signals and indicate
	BMI-independent effects of puberty timing on cancer susceptibility," Felix R. Day et al.,
	Nature Genetics.

- "Enhancer-promoter interactions are encoded by complex genomic signatures on looping chromatin," **Sean Whalen**, Rebecca M. Truty, and Katherine S. Pollard, *Nature Genetics*.
- "Unboxing Cluster Heatmaps," Sophie Engle, **Sean Whalen**, Alark Joshi, and Katherine S. Pollard, *Proceedings of the 6th Symposium on Biological Data Visualization (BioVis)*, 2016.
- "Prediction of Human Population Responses to Toxic Compounds by a Collaborative Competition," Federica Eduati et al., *Nature Biotechnology*.
- "Predicting Protein Function and Other Biomedical Characteristics with Heterogeneous Ensembles," **Sean Whalen**, Om Pandey, and Gaurav Pandey, *Methods*.
- "Model Aggregation for Distributed Content Anomaly Detection," **Sean Whalen**, Nathaniel Boggs, and Salvatore J. Stolfo, *Proceedings of the 7th ACM Workshop on Artificial Intelligence and Security.*
- "Enhancing the Functional Content of Eukaryotic Protein Interaction Networks," Gaurav Pandey, Sonali Arora, Sahil Manocha, and **Sean Whalen**, *PLoS ONE*.
- "A Comparative Analysis of Ensemble Classifiers: Case Studies in Genomics," **Sean Whalen** and Gaurav Pandey, *Proceedings of the 13th IEEE International Conference on Data Mining.*
- "Multiclass Classification of Distributed Memory Parallel Computations," **Sean Whalen**, Sean Peisert, and Matt Bishop, *Pattern Recognition Letters*.
- "Visualizing Distributed Memory Computations with Hive Plots," Sophie Engle and **Sean** Whalen (co-first), *Proceedings of the 9th International Symposium on Visualization for Cyber Security.*
- "Structural Drift: The Population Dynamics of Sequential Learning," James P. Crutchfield and Sean Whalen (co-first), PLoS Computational Biology.
- "Network-Theoretic Classification of Parallel Computation Patterns," Sean Whalen, Sophie

- Engle, Sean Peisert, and Matt Bishop, International Journal of High Performance Computing Applications.
- "A Taxonomy of Buffer Overflow Characteristics," Matt Bishop, Sophie Engle, Damien Howard, and **Sean Whalen**, *IEEE Transactions on Dependable and Secure Computing*.
- "This is the Remix: Structural Improvisation using Automated Pattern Discovery," **Sean Whalen** and James P. Crutchfield, *Proceedings of the 4th International Workshop on Machine Learning and Music.*
- "Network-Theoretic Classification of Parallel Computation Patterns," **Sean Whalen**, Sean Peisert, and Matt Bishop, *Proceedings of the 1st International Workshop on Characterizing Applications for Heterogeneous Exascale Systems*.
- "Hidden Markov Models for Automated Protocol Learning," **Sean Whalen**, Matt Bishop, and James P. Crutchfield, Proceedings of the 6th International ICST Conference on Security and Privacy in Communication Networks.
- "Case Studies of an Insider Framework," Matt Bishop, Sophie Engle, Carrie Gates, Sean Peisert, and **Sean Whalen**, *Proceedings of the 42nd Annual Hawaii International Conference on System Sciences*.
- "We Have Met the Enemy and He is Us," Matt Bishop, Sophie Engle, Carrie Gates, Sean Peisert, and **Sean Whalen**, *Proceedings of the 2008 New Security Paradigms Workshop*.

#### **Book Chapters**

2010 "A Risk Management Approach to the Insider Threat," Matt Bishop, Sophie Engle, Deborah A. Frincke, Carrie Gates, Frank L. Greitzer, Sean Peisert, and **Sean Whalen**, in "Insider Threats in Cybersecurity—And Beyond," Springer Verlag, Berlin.

#### Talks & Posters

- "Enhancer-promoter interactions are encoded by complex genomic signatures on looping chromatin", Biology of Genomes 2016, Cold Spring Harbor, NY.
- "Supervised ensembles to boost the predictive power of DREAM challenges", Intelligent Systems for Molecular Biology 2016, Orlando, FL.
- "The Genesips Project: an NHLBI-Sponsored induced Pluripotent Stem Cell (iPSC) Resource for the Study of Cardiovascular Diseases", American Heart Association Scientific Session 2014, Chicago, IL.
- "Predicting Protein Function with Heterogeneous Ensembles", Automated Protein Function Prediction 2014 (held in conjunction with the 22nd International Conference on Intelligent Systems in Molecular Biology), Boston, MA.
- "Enhancing the Functional Content of Protein Interaction Networks", Poster, Automated Protein Function Prediction 2014 (held in conjunction with the 22nd International Conference on Intelligent Systems in Molecular Biology), Boston, MA.
- "Measuring Access, Knowledge, and Trust: A Discussion", Dagstuhl Workshop on Organizational Processes for Supporting Sustainable Security, Dagstuhl, Germany.
- "This is the Remix: Structural Improvisation using Automated Pattern Discovery", 4th International Workshop on Machine Learning and Music (held in conjunction with the 25th Annual Conference on Neural Information Processing Systems), Sierra Nevada, Spain.

- "Hybrid Approaches for Classifying Parallel Computation," Poster, Department of Energy 2011 Applied Mathematics Meeting, Washington DC.
- "Network-Theoretic Classification of Parallel Computation Patterns", 1st International 2011 Workshop on Characterizing Applications for Heterogeneous Exascale Systems (held in conjunction with the 25th International Conference on Supercomputing), Tucson, AZ.
- "Hidden Markov Models for Automated Protocol Learning", 6th International Conference 2010 on Security and Privacy in Communication Networks, Singapore.
- "Anomaly Detection in High Performance Computing", Discovery 2015: Cloud Computing 2010 Institute, UCBerkeley, Berkeley, CA.
- "Realtime Social Network Analysis", SFI Workshop on Immersive Visualization, Santa Fe 2008 Institute, Santa Fe, NM.

#### Service

#### **Additional Teaching**

- 2-Part Lab, Intro to Machine Learning with R, Systems Pharmacology PSPG245, UCSF, 2016-2017 Sourav Bandyopadhyay & Rada Savic
- Lab, Intro to Machine Learning with R, Statistical Methods for Bioinformatics BMI206, 2015 UCSF, Katie Pollard
- Lab, Intro to Machine Learning with R, Systems Pharmacology PSPG245, UCSF, Natalia 2015 Khuri
- Lab, Intro to Machine Learning with Python, Statistical Methods for Bioinformatics BMI206, 2014 UCSF, Katie Pollard

#### **Committees**

- Program Committee, 13th International Symposium on Visualization for Cyber Security 2016 Program Committee, 12th International Symposium on Visualization for Cyber Security 2015
- Organizing Committee, 36th IEEE Symposium on Security and Privacy 2015
- Program Committee, Cyber and Information Security Research Conference 2015
- Program Committee, Cyber Security Awareness Week 2014
- Program Committee, Automated Protein Function Prediction 2014 2014
- Program Committee, 11th International Symposium on Visualization for Cyber Security 2014
- Organizing Committee, 35th IEEE Symposium on Security and Privacy 2014
- Program Committee, Cyber Security Awareness Week 2013
- Program Committee, 5th Workshop on Cyber Security Experimentation and Test 2012

#### **Reviews**

12th & 13th Annual IEEE Conference on Technologies for Homeland Security; 19th & 20th ACM SIGKDD Conference on Knowledge Discovery and Data Mining; 22nd ACM International Conference on Information and Knowledge Management; 5th European Workshop on Systems Security (ACM SIGOPS EuroSys Conference); Bioinformatics; Chromosome Research; IEEE Computer; IEEE Internet Computing; IEEE Transactions on Dependable and Secure Computing; IEEE Transactions on Systems, Man, and Cybernetics Part B; International Journal of Information Security; Journal of Applied Mathematics; Nature Genetics; Nucleic Acids Research; R Journal

## References (Alphabetical)

Prof. Matt Bishop bishop at cs.ucdavis.edu
Prof. James P. Crutchfield crutchfield at ucdavis.edu
Prof. Gaurav Pandey gaurav.pandey at mssm.edu

Prof. Sean Peisert sppeisert at lbl.gov

Prof. Katherine S. Pollard katherine.pollard at gladstone.ucsf.edu

Prof. Salvatore J. Stolfo sal at cs.columbia.edu