Ameen Eetemadi

Title: Ph.D. Candidate

Research Area: Applied Machine Learning

Citizenship: U.S. Citizen email:eetemadi@ucdavis.edu Department of Computer Science and

UC Davis Genome Center,

Kemper Hall 2063,

University of California, Davis

EDUCATION	 ♦ University of California, Davis, CA Ph.D. Candidate in Computer Science (Machine Learning) Advisor: Prof. Ilias Tagkopoulos 	(2014 - now)
	 ♦ Wayne State University, Detroit MI M.Sc in Computer Science (Data Mining) Advisor: Prof. Farshad Fotouhi 	(graduated 2012)
	 Sharif University of Technology, Tehran, Iran B.Sc in Computer Engineering (Software) 	(graduated 2005)
Work experience	♦ University of California , Davis, CA Graduate Research Assistant, Department of Computer Science	(2014 - now) and Genome Center
	♦ Microsoft, Redmond, WA Software Development Engineer, Microsoft Office Team	(2008 - 2014)
	♦ Microsoft, Redmond, WA Software Development Engineer Intern, Microsoft Research	(Summer 2006)
	♦ Henry Ford Health Systems, Detroit, MI Graduate Research Assistant, Health Informatics	(2005 - 2008)
TEACHING EXPERIENCE	♦ University of California, Davis, CA Lead Teaching Assistant: ECS 171 - Machine Learning	(Fall 2014)
	Graded homeworks and exams; led the discussion section for aro	und 100 students
SKILLS	> Programming Languages MATLAB, C++, C#, SQL, HTML, Java, JavaScript, Perl, ASP.net, PHP, Ruby, R	
	♦ Software Technologies TCP/IP, RESTful web services, Parallel Programming, DNA/RNA Sequence Analysis	
CURRENT PROJECTS	♦ Deep learning architectures for multi-omics data Investigation of machine learning techniques for predictive modeling. Part of a team to	

SELECTED

♦ Ameen, Eetemadi, Mohammad-Reza Siadat, Hamid Soltanian-Zadeh, Farshad Fotouhi, Publications and Kost Elisevich. "Content-Based Support Environment (C-BASE): Data Preparation and Similarity Measurement.", Proceedings of the Seventh IEEE International Conference on Data Mining (ICDM'07), pp. 145-150, Omaha, NE, USA, October 28-31, 2007.

♦ Biomarker discovery using genome-level analysis of patients after ischemic episode Analysis of transcriptional profiling data (RNA-Seq) from patients with an ischemic episode. Identification of distinct patient groups, prediction of optimal treatment and outcome.

develop a data-driven, genome-scale model of the bacterium Escherichia coli.

♦ Siadat, Mohammad-Reza, Hamid Soltanian-Zadeh, Farshad Fotouhi, and Ameen Eetemadi. "Data modeling for content-based support environment (C-BASE): Application on epilepsy data mining." Proceedings of the Seventh IEEE International Conference on Data Mining (ICDM'07), pp. 181-188, Omaha, NE, USA, October 28-31, 2007.