

# 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	<ul style="list-style-type: none"><li>◇ <b>University of California</b>, Davis, CA (2014 - now) Ph.D. Candidate in Computer Science (Machine Learning) Advisor: Prof. Ilias Tagkopoulos</li><li>◇ <b>Wayne State University</b>, Detroit MI (graduated 2012) M.Sc in Computer Science (Data Mining) Advisor: Prof. Farshad Fotouhi</li><li>◇ <b>Sharif University of Technology</b>, Tehran, Iran (graduated 2005) B.Sc in Computer Engineering (Software)</li></ul>
WORK EXPERIENCE	<ul style="list-style-type: none"><li>◇ <b>University of California</b>, Davis, CA (2014 - now) Graduate Research Assistant, Department of Computer Science and Genome Center</li><li>◇ <b>Microsoft</b>, Redmond, WA (2008 - 2014) Software Development Engineer, Microsoft Office Team</li><li>◇ <b>Microsoft</b>, Redmond, WA (Summer 2006) Software Development Engineer Intern, Microsoft Research</li><li>◇ <b>Henry Ford Health Systems</b>, Detroit, MI (2005 - 2008) Graduate Research Assistant, Health Informatics</li></ul>
SELECTED PUBLICATIONS	<ul style="list-style-type: none"><li>◇ <b>Eetemadi, A.</b> and Tagkopoulos, I., Low-FODMAP diet and microbiome in irritable bowel syndrome, a meta-analysis. (manuscript ready).</li><li>◇ Wang X, Rai N.,Pereira B., <b>Eetemadi, A.</b> and Tagkopoulos, I., Accelerated knowledge discovery from omics data by optimal experimental design. (manuscript under review)</li><li>◇ <b>Eetemadi, A.</b>, Rai N., Pereira B., Kim M., Schmitz H. and Tagkopoulos, I., 2020, The Computational Diet: A Review of Computational Methods Across Diet, Microbiome, and Health. <i>Frontiers in Microbiology</i></li><li>◇ <b>Eetemadi, A.</b> and Tagkopoulos, I., 2019. Genetic Neural Networks: an artificial neural network architecture for capturing gene expression relationships. <i>Bioinformatics</i>.</li><li>◇ Kim, M.*, <b>Eetemadi, A.*</b> and Tagkopoulos, I., 2017. DeepPep: Deep proteome inference from peptide profiles. <i>PLoS computational biology</i>, 13(9), p.e1005661. (<b>*contributed equally</b>)</li><li>◇ <b>Eetemadi, A.</b>, 2012. Medical data analysis method for epilepsy, Master's dissertation, <i>Wayne State University</i>.</li><li>◇ <b>Eetemadi, A.</b>, Siadat, M.R., Soltanian-Zadeh, H., Fotouhi, F. and Elisevich, K., 2007, "Content-Based Support Environment (C-BASE): Data Preparation and Similarity Measurement.",<i>Proceedings of the Seventh IEEE International Conference on Data Mining (ICDM'07)</i>, pp. 145-150, Omaha, NE, USA, October 28-31,</li></ul>
SKILLS	<ul style="list-style-type: none"><li>◇ <b>Programming Technologies</b> Python, R, MATLAB, C++, C#, Java, HTML JavaScript, AngularJS, Node.js, Perl, ASP.net, PHP, HPC (Slurm, TORQUE)</li><li>◇ <b>Software Technologies</b> Supervised and Unsupervised Machine Learning (worked with Torch7, TensorFlow and scikit-learn), TCP/IP, RESTful web services and sequence analysis (DNA, RNA and Metagenomics).</li><li>◇ <b>Database Systems</b> MongoDB, PostgreSQL, Oracle (+PL/SQL), MSSQL</li></ul>
TEACHING EXPERIENCE	<ul style="list-style-type: none"><li>◇ <b>University of California</b>, Davis, CA (2014 - 2019) Teaching Assistant: ECS 171 Machine Learning, ECS 124 Bioinformatics, ECS 120 Theory of Computations, ECS 36C Data Structures and Algorithms, ECS 30 Programming&amp;Prob Solving.</li></ul>