Mehmet Serkan APAYDIN

Education:		
Bilkent University, Ankara, TURKEY ☐ B.Sc., Electrical Engineering, June 1997.		
Stanford University, Stanford, CA USA ☐ M.S., Electrical Engineering, June 1999. ☐ Ph.D., Electrical Engineering, September 2004.		
Employment:		
Stanford University, Stanford, CA USA ☐ Research and Teaching Assistant, Department of Computer Science (15/9/1997 – 30/8/2004) ☐ Teaching assistant for Motion planning and C++ classes ☐ Advisors: Prof. Jean-Claude Latombe and Prof. Doug Brutlag		
Compaq Cambridge Research Labs, Cambridge, MA USA ☐ Summer intern (15/6/2000-15/9/2000) ☐ Research on whole genome assembly algorithms, paper in Recomb conference ☐ Advisor: Prof. Simon Kasif		
Dartmouth College, Hanover, NH USA ☐ Research Associate, Computer Science Department (15/11/2004 – 30/8/2006) ☐ Protein structural bioinformatics research, paper in Recomb conference ☐ Advisor: Prof. Bruce R. Donald		
 Duke University, Durham, NC USA Research Associate, Computer Science Department (1/9/2006 – 30/8/2008) NMR protein structure-based assignment algorithms, paper in J. Biomolecular NMR Advisor: Prof. Bruce R. Donald 		
Sabanci University, Istanbul, TURKEY □ Visiting Faculty Member (1/9/2008 – 30/6/2010), Principal Investigator (1/9/2010 – 30/8/2013) Faculty of Engineering and Natural Sciences □ Taught grad level bioinformatics, algorithms for computational structural biology, motion planning Recipient of European Union Marie Curie International Reintegration Grant (75000 euros) Recipient of Tübitak research grant (92000 TRY) Research on NMR protein structural based assignments		
 Istanbul Sehir University, Istanbul, TURKEY Faculty Member (19/7/2010 − 10/2020) (On leave from 3/2018 to 6/2019) College of Engineering and Natural Sciences, Department of Computer Science. Classes on AI, deep learning, machine learning, natural language processing, computer vision, reinforcement learning, linear dynamical systems, Unix, EECS I and II, software-based startups. Bilateral research grant with CNRS Gif-sur-Yvette on NMR software development for structure-based assignments Research on traffic speed prediction using machine learning Papers on GECCO, SIU conferences, RAIRO journal Graduate student supervision 		
Inria, Sophia Antipolis, FRANCE. ☐ College de France Fellow (1/3/2018 – 6/2019) ☐ Deep learning for protein structural informatics		
Kariyer.net, Istanbul ☐ Research and Development Consultant (3/2020-April 2021) Natural language processing projects using deep learning, in Turkish.		

	Algorithms in HR domain (CV parsing, CV – job ad matching, job ad text generation), content moderation for job ads, messages.	
Istanb	Faculty Member (22/10/2010 – April 2021) College of Engineering and Natural Sciences, Department of Computer Science. Classes on operating systems, probability and random variables (135 students), deep learning for natural language processing. (all remote) Research supervision of students on applications of machine and deep learning.	
İzmir]	Institute of Technology, Izmir, TURKEY Faculty Member (April 2021-Present) Department of Electrical and Electronics Engineering.	
Awards:		
	David L. Cheriton Stanford Graduate Fellowship, 1/6/1998-31/12/2003. Ranked 1st in the Faculty of Engineering, Bilkent University, 15/6/1997. Full scholarship, Bilkent University, 1/10/1993-1/6/1997. Tubitak (Turkish National Science Foundation) fellowship, 1/10/1993-1/6/1997. Bronze medalist in International Olympiad of Informatics, Argentina, 1/10/1993.	
Grant	s:	
	Nvidia (2) Titan XP GPU grant. 15/09/2017.	
Classes taught: (5xx classes are graduate level, 1xx, 2xx,3xx and 4xx coded classes are undergraduate level classes.)		
(at Izn	nir Institute of Technology)	
(at Ista	anbul Medipol University)	
	2 nd year Engineering students, Probability and Random Variables	
(at Ista	anbul Sehir University)	
	EECS201, System Design Fundamentals EECS202, Into to EECS II EECS213, UNIX Operating System	

	CB311 Operating Bysteins
	EECS 491/492, Global Design Project
	EECS420/ECE520, Software Based Start-ups
	ECE516, Computational Biology
	ECE515, Linear Dynamical Systems
	EECS461/ECE523, Machine Learning
	EECS464/ECE524 Computer Vision
	EECS489/ECE589 Deep Learning for Natural Language Processing
at Saba	anci University)
	ENS 491/2, Graduation Project
	CS526, Motion Planning
	CS581, Algorithms for Computational Structural Biology
	Bio511, Advanced Bioinformatics

CS31/ Operating Systems

Journal Articles

- 1. A Saracoglu, H Ozen, MS Apaydin, A Maltas. A New Approach to Determine Traffic Peak Periods to Utilize in Transportation Planning. Arabian Journal for Science and Engineering, 1-10 (2021)
- 2. S. Çetinkaya, Ş.N. Ekren, M. S. Apaydin. Progress in Nuclear Vector Replacement for NMR Protein Structure Based Assignments. RAIRO Operations Research 50(2):341-349, 2016
- 3. M. Akhmedov, B. Çatay, M.S. Apaydin. Automating unambiguous NOE data usage in NVR for NMR protein structure-based assignments. J. Bioinform. Comput. Biol. 13(6), 2015
- 4. G. Çavuşlar, B. Çatay, M. S. Apaydin. A Tabu Search Approach for the NMR Protein Structure-Based Assignment Problem. IEEE Transactions on Computational Biology and Bioinformatics 9 (6): 1621-1628, 2012
- M. S. Apaydin, B. Çatay, N. Patrick and B.R. Donald. NVR-BIP: Nuclear Vector Replacement using Binary Integer Programming for NMR Structure-Based Assignments. The Computer Journal 54 (5): 708-716, 2011
- 6. M. S. Apaydin, V. Conitzer, B.R. Donald. Structure-Based Protein NMR Assignments using Native Structural Ensembles. In Journal of Biomolecular NMR, 40(4):263-276, 2008
- 7. T.H. Chiang, M. S. Apaydin, D.L. Brutlag, D. Hsu, J.-C. Latombe. Using stochastic roadmap simulation to predict experimental quantities in protein folding kinetics: folding rates and phi-values. In Journal of Computational Biology 14(5):578-93, 2007
- 8. Apaydin, D.L. Brutlag, C. Guestrin, D.Hsu, J.-C.Latombe and C. Varma. Stochastic Roadmap Simulation: An efficient Representation and Algorithm for Analyzing Molecular Motion. In Journal of Computational Biology, 10 (3-4):257-281, 2003
- 9. Apaydin, C. Guestrin, C. Varma, D.L. Brutlag, and J.-C. Latombe. Studying Protein-Ligand Interactions with Stochastic Roadmap Simulation. In Bioinformatics, 18 Suppl. 2:S18-S26, 2002

Conference And Workshop Papers

- 1. B.Ciftci, M.S. Apaydin. A Deep Learning Approach to Sentiment Analysis in Turkish reviews dataset. International Conference on AI and Data Processing Conference, 2018.
- 2. U. C. Cakmak, M S. Apaydin and B. Catay. A Neural Network Approach for Predicting Speeds on Road Networks. Signal Processing Applications (SIU), 2018.
- 3. U.C. Cakmak, M. S. Apaydin and B. Catay. Traffic speed prediction with neural networks, Operations Research (OR) 2017 and Euro Working Group on Transportation Meeting (EWGT), 2017.
- 4. S. Cetinkaya and S. N. Ekren and M. S. Apaydin. Progress in Automated NMR Protein Structure-Based Assignments. European Conference for Operational Research, 2015.
- 5. S. N. Ekren and M. S. Apaydin. Incorporating Triple Resonance Experiment Data Into NVR for NMR Structure-Based Assignments. SIU, 2014.
- 6. C. Aslanov and B. Çatay and M. S. Apaydin. An ant colony optimization based approach for solving the NMR Structure-Based Assignment Problem. Genetic and Evolutionary Computation Conference (GECCO), 2013.
- 7. M. Akhmedov and B. Çatay and M. S. Apaydin. Distinguishing the type of NOE for NMR Protein Structure-Based Assignments. SIU, 2013.
- 8. H. Erdoğan and M. S. Apaydin. Incorporating HADAMAC experiment into NMR Structure-based assignments. Sixth International Symposium on Health Informatics and Bioinformatics (HIBIT), 2011.
- 9. H. Erdoğan and M. S. Apaydin. Using Amino Acid Typing to Improve the Accuracy of NMR Structure Based Assignments. HIBIT, 2010.

- 10. M. S. Apaydin, B. Çatay, N. Patrick and B.R. Donald. NVR-BIP: Nuclear Vector Replacement using Binary Integer Programming for NMR Structure-Based Assignments. In International Conference on Computer and Information Sciences (ISCIS), 2009.
- 11. T.H. Chiang, M. S. Apaydin, D.L. Brutlag, D. Hsu, and J.-C. Latombe. Predicting Experimental Quantities in Protein Folding Kinetics using Stochastic Roadmap Simulation. In Proceedings of the Tenth Annual International Conference on Research in Computational Molecular Biology (RECOMB), 2006.
- 12. M. S. Apaydin, D.L. Brutlag, C. Guestrin, D. Hsu and J.-C. Latombe. Stochastic Conformational Roadmaps for Computing Ensemble Properties of Molecular Motion. In Algorithmic Foundations of Robotics V (WAFR), 2002.
- 13. M. S. Apaydin, D.L. Brutlag, C. Guestrin, D. Hsu. and J.-C. Latombe. Stochastic Roadmap Simulation: An Efficient Representation and Algorithm for Analyzing Molecular Motion. In Proceedings of the Tenth Annual International Conference on Research in Computational Molecular Biology (RECOMB), 2002.
- 14. M. S. Apaydin, A.P. Singh, D.L. Brutlag, and J.-C. Latombe. Capturing Molecular Energy Landscapes with Probabilistic Conformational Roadmaps. In International Conference on Robotics and Automation (ICRA), 2001.
- 15. R. Beigel, N. Alon, , L. Fortnow and S. Kasif. An optimal procedure for gap closing in whole genome shotgun sequences. In Proceedings of the Fifth Annual International Conference on Research in Computational Molecular Biology (RECOMB), 2001.

Languages:

Turkish (native), English (fluent), French (fluent), German (basic).

M. Sc. Students supervised:

Madison, USA.

Format	: Student Name, Dates, Keywords, Next position (if applicable).			
@Şehir University:				
	B. Çiftçi, 2017-2019, #Deep Learning, #Sentiment Analysis, #Turkish @Software engineer at TRT world S. Cetinkaya, 1/10/2013—30/8/2015, #NMR, #Structure-Based Assignments. @Software engineer at Turkish airlines			
@Saba	nci University (co-supervised with Prof. B. Catay):			
	Umut Can Cakmak, 1/9/2015—30/8/2017, #Machine and Deep Learning, #Traffic speed prediction, Ph.D. Student @ Erasmus University, Netherlands.			
	M. Akhmedov, 1/9/2011—30/8/2013, #NMR, #Structure-Based Assignments, PhD Student at University of Lugano, Switzerland			
	G. Cavuşlar, 1/9/2009—30/8/2011, #NMR, #Tabu Search, PhD Student at University of Wisconsin-			