

Haris Vikalo

Education

- 2003 **Ph.D. in Electrical Engineering**, *Stanford University*, Stanford, CA
- 1997 **M.Sc. in Electrical Engineering**, *Lehigh University*, Bethlehem, PA
- 1995 **B.Sc. in Electrical Engineering**, *University of Zagreb*, Zagreb, Croatia

Experience

Current and Previous Academic Positions

- 2019 - **Professor**, *ECE Department, University of Texas at Austin*, Austin
- 2013 - 2019 **Associate Professor**, *ECE Department, University of Texas at Austin*, Austin
- 2007 - 2013 **Assistant Professor**, *ECE Department, University of Texas at Austin*, Austin
- 2003 - 2007 **Associate Scientist**, *EE Department, California Institute of Technology*, Pasadena

Other Professional Experience

- 2014 - 2014 **Visiting Professor**, *Max F. Perutz Laboratories*, Vienna, Austria
- Summer 1999 **Member of Technical Staff**, *Bell Labs*, Murray Hill, NJ

Teaching in the Last Five Years

- '26, '25, '23 *Statistical Machine Learning* (graduate)
- '24 *Genomic Signal Processing and Data Science* (graduate)
- '22 *Estimation Theory* (graduate)
- '22-'25 *Digital Signal Processing* (undergraduate)

Selected Honors and Awards

- 2025 Temple Foundation Endowed Professorship No. 3
- 2021-2025 William W. Hagerty Fellowship in Engineering (UT Cockrell School of Engineering)
- 2017 Jack Kilby/Texas Instruments Endowed Faculty Fellowship in Computer Engineering (UT ECE)
- 2015 IEEE Senior Member
- 2013 Jack Kilby/Texas Instruments Endowed Faculty Fellowship in Computer Engineering (UT ECE)
- 2009 National Science Foundation CAREER Award

Memberships in Professional Societies

- Institute of Electrical and Electronics Engineers (IEEE), Senior Member
- IEEE Signal Processing Society

University and College Committee Assignments

Departmental

- 2013–present DICE Academic Track Graduate Advisor
- 2024–present Faculty Annual Review Committee

- 2025–2026 Member, ECE Junior Faculty Recruiting Committee
- 2016–2025 bioECE Academic Track Admission Coordinator
- 2024–2025 Member, ECE Junior Faculty Recruiting Committee
- 2018–2019 Member, ECE Junior Faculty Recruiting Committee
- 2017–2018 Member, ECE Junior Faculty Recruiting Committee
- 2016–2017 Member, ECE Junior Faculty Recruiting Committee
- 2014–2015 Member, ECE Junior Faculty Recruiting Committee
- 2013–2014 Member, ECE Junior Faculty Recruiting Committee
- 2007–present Member, DICE Academic Track Admission Committee
- 2014–present Annual WNCG Open House Coordinator
 - 2014 ECE Benchmarking/Directions Committee
 - 2013 ECE Recruiting Vision Committee
- 2012–2013 DICE pre-qual committee
- 2016–2017 DICE pre-qual committee
- 2012–2015 Major Sequence Appeals Committee

College

- 2025–present CSE Degrees & Courses Committee
- 2011–2012 Systems Biology faculty recruitment committee

University

- 2019–2020 Provost’s Health Translation Opportunities Committee for Oncology
- 2024–2025 VPR Office Internal Competition Google Fellowship Committee
- 2025–2026 VPR Office Internal Competition Google Fellowship Committee

Community Activities and Professional Service

- 2018 - present Associate Editor, IEEE Trans. on Molecular, Biol. and Multi-Scale Communications
 - 2026 Area Chair, The Fortieth Annual Conference on Neural Information Processing Systems (NeurIPS)
 - 2025 Technical Program Chair, 30th International Conference on Information, Communication, and Automation Technologies (ICAT 2025), Sarajevo, Bosnia and Herzegovina
 - 2023 Technical Program Chair, 29th International Conference on Information, Communication, and Automation Technologies (ICAT 2023), Sarajevo, Bosnia and Herzegovina
 - 2022 Program Committee Member, 7th International Workshop on Computational Network Biology (CNB-MAC), Chicago, IL
 - 2022 Program Chair, 28th International Conference on Information, Communication and Automation Technologies (ICAT 2022), Sarajevo, Bosnia and Herzegovina
 - 2022 NIH Proposal Panel Reviewer
 - 2022 Technical Program Committee Member, 5th International Workshop on Computational Network Biology (CNB-MAC), Chicago, IL
 - 2021 Senior Program Committee Member, the 35th AAAI Conference on Artificial Intelligence
 - 2021 NSF Proposal Panel Reviewer, CCF Program
 - 2019 Technical Program Committee Member, 6th International Workshop on Computational Network Biology (CNB-MAC), Niagara Falls, NY
 - 2018 Technical Program Committee Member, IEEE International Symposium on Information Theory, Vail, CO

- 2017 Technical Program Committee Member, the 4th International Workshop on Computational Network Biology (CNB-MAC), Boston, MA
- 2017 Session Chair (Genome Assembly and DNA Storage Systems), 2017 Information Theory and Applications Workshop, San Diego, CA
- 2017 NSF Proposal Panel Reviewer, CCF Program
- 2015 Technical Program Chair, Symposium on Signal Processing and Mathematical Modeling of Biological Processes with Applications to Cyber-Physical Systems for Precise Medicine, IEEE Global Conference on Signal and Information Processing, Orlando, FL
- 2015 Technical Program Committee Member, the 2nd International Workshop on Computational Network Biology (CNB-MAC), Atlanta, GA
- 2014 NSF Proposal Panel Reviewer, CCF Program
- 2014 Technical Program Committee Member, the 8th International Symposium on Turbo Codes and Iterative Information Processing (ISTC), Bremen, Germany
- 2014 Technical Program Committee Member, the 1st International Workshop on Computational Network Biology (CNB-MAC), Newport Beach, CA
- 2013 General Chair, Symposium on Bioinformatics and Systems Biology, IEEE Global Conference on Signal and Information Processing (GlobalSIP), Austin, TX
- 2013 Technical Program Chair, the 24th International Conference on Information, Communication, and Automation Technologies, Sarajevo, Bosnia and Herzegovina
- 2013 Student Paper Award Chair, IEEE Workshop on Genomic Signal Processing and Statistics (GENSIPS), Houston, TX
- 2011 Technical Program Committee Member, IEEE Workshop on Genomic Signal Processing and Statistics (GENSIPS), San Antonio, TX
- 2011 Technical Program Chair, the IEEE Asilomar Conference on Signals, Systems, and Computers, Pacific Grove, CA (Track F: Bio image and signal processing)
- 2010 Technical Program Chair, IEEE International Workshop on Genomic Signal Processing and Statistics (GENSIPS), Cold Spring Harbor, NY
- 2010 Guest Editor, EURASIP Journal on Advances in Signal Processing, Special Issue on Genomic Signal Processing
- 2009 Tutorial Chair, IEEE International Workshop on Genomic Signal Processing and Statistics (GENSIPS), Minneapolis, MN
- 2009 Tutorial Program Committee Member, IEEE International Workshop on Genomic Signal Processing and Statistics (GENSIPS), Minneapolis, MN
- 2008 Co-organized and presented a full-day tutorial on Bioinformatics and Computational Biology I, II at the IEEE International Conference on Acoustics, Speech, and Signal Processing, Las Vegas, NV
- 2007 Technical Program Committee Member, IEEE Workshop on Genomic Signal Processing and Statistics (GENSIPS), Tuusula, Finland

Advising

PhD In Progress

Usman Akram, Seohyeon Cha, Haoran Zhang, Dongjun Kim, John Robertson, Kutay Tire

PhD Supervisions Completed

- 2025 Huancheng Chen
- 2024 Shorya Consul
- 2023 Yiyue Chen

2022 Monica Ribero
2021 Ziqi Ke
2020 Abolfazl Hashemi
2018 Soyeon Ahn
2017 Somsubhra Barik
2016 Natalia Arzeno-Gonzalez
2015 Shreepriya Das
2014 Xiaohu Shen
2012 Manohar Shamaiah
2011 Sang-Hyun Lee (co-supervised)
[MSc Supervisions Completed](#)
2024 Ahmet Aydin
2022 Sameer Bibikar, Huancheng Chen
2020 Yiyue Chen, Monica Ribero
2019 Shorya Consul, Ziqi Ke
2016 Abolfazl Hashemi
2012 Shreepriya Das
2011 Ting Wu
2010 Mahsuni Gokdemir, Mijung Park (co-supervised), Xiaohu Shen
2009 Juhun Lee

Grants and Contracts

Total funding \$6,153,799. My share \$4,444,329.

- 2024–2025 Towards Situational Understanding at the Tactical Edge: Uncertainty-Driven Agile Machine Learning System (\$1,038,940, co-PI with Prof. G. de Veciana, my share \$519,470). Agency: Army Research Laboratory.
- 2024–2025 Federated Learning in Large-Scale Systems: Training Personalized Models on Heterogeneous Time-Series Data (\$120,000, PI). Agency: Toyota.
- 2022–2025 RINGS: Scalable and Resilient Networked Learning Systems (\$630,000, co-PI with Prof. G. de Veciana, my share \$315,000). Agency: National Science Foundation.
- 2022–2023 The Value of Side Information in Sensing (\$60,000, co-PI with Prof. G. de Veciana). Agency: Honda.
 - 2023 Federated Learning for Vehicular Systems under Communication and Computation Constraints (\$118,000, PI). Agency: Toyota.
 - 2022 Federated Learning for Vehicular Systems under Communication and Computation Constraints (\$77,100, PI). Agency: Toyota.
- 2021–2025 FET: Small: Accurate and Scalable Methods for Analysis of Complex Genomic Populations (\$500,000, PI). Agency: National Science Foundation.
- 2020–2022 RAPID: Methods for Reconstructing Disease Transmissions from Viral Genomic Data with Application to COVID-19 (\$100,000, PI). Agency: National Science Foundation.
- 2019–2022 Distributed Learning under Privacy and Resource Constraints (\$480,000, co-PI with Profs. C. Julien and A. Gerstlauer, my share \$160,000). Agency: Army Futures Command.
- 2019–2022 RF-Based Wireless Network Tomography (\$540,000, co-PI with Prof. G. de Veciana, my share \$270,000). Agency: Army Futures Command.

- 2018–2022 Visibility and Interactive Information Sharing in Collaborative Sensing Systems (\$450,000, co-PI with Prof. G. de Veciana, my share \$225,000). Agency: National Science Foundation.
- 2018–2019 Situational Awareness via Interactive Information Sharing in Collaborative Sensing Systems (\$50,000, co-PI with Prof. G. de Veciana, my share \$25,000). Agency: Qualcomm.
- 2016–2019 Reconstructing Mixtures of DNA Sequences from High-Throughput Sequencing Data (\$400,000, PI). Agency: National Science Foundation.
- 2014–2016 RAPID: Methods for Estimating Genetic Diversity of the Ebola Virus (\$200,000, PI). Agency: National Science Foundation.
- 2013–2017 Algorithms for Haplotype Assembly from Next-Generation Sequencing Data (\$400,000, PI). Agency: National Science Foundation.
- 2013–2018 WNCG Industrial Affiliates (\$71,000, PI). Agency: Industry (various).
- 2011–2013 Algorithms for Base-Calling in Sequencing-By-Synthesis (\$352,259, PI). Agency: National Institutes of Health.
- 2009–2013 Modeling, Estimation, and Coding for Biosensor Arrays (\$400,000, PI). Agency: National Science Foundation (CAREER Award).
- 2010–2011 Next Generation Sequencing: Signal Processing Perspective (\$122,500, PI). Agency: National Science Foundation.
- 2009–2013 WNCG Industrial Affiliates (\$44,000, PI). Agency: Industry (various).

Selected Invited Talks

- 2021 Texas A&M, Data Science Institute
“Sensing and Learning in Distributed Systems Operating under Resource Constraints”
- 2018 University of Southern California, CBIO Colloquium
“Efficient Algorithms for Haplotype Assembly and Viral Quasispecies Reconstruction”
- 2017 The Chinese University of Hong Kong
“Algorithms for Haplotype Assembly”
- 2015 USC, 2015 Molecular, Biological and Multi-Scale Communications Workshop
“Algorithms for Haplotype Assembly”
- 2015 California Institute of Technology, Department of Electrical Engineering
“Algorithms for Haplotype Assembly”
- 2015 Stanford University, Kailath Lecture and Colloquia
“Algorithms for Haplotype Assembly”
- 2014 EPFL, School of Computer and Communication Sciences
“Decoding Genetic Variations: Communications-Inspired Haplotype Assembly”
- 2014 ETHZ, Department of Information Technology and Electrical Engineering
“Decoding Genetic Variations: Communications-Inspired Haplotype Assembly”
- 2014 TU Munich, Department of Electrical Engineering
“Decoding Genetic Variations: Communications-Inspired Haplotype Assembly”
- 2014 University of Stuttgart, Department of Electrical Engineering
“Decoding Genetic Variations: Communications-Inspired Haplotype Assembly”
- 2014 Max F. Perutz Laboratories, Center for Integrative Bioinformatics, Vienna
“Decoding Genetic Variations: Communications-Inspired Haplotype Assembly”
- 2012 Korea Advanced Institute of Science and Technology, Dept. of Electrical Engineering
“Signal Processing for Next Generation Biosensing and Sequencing”

- 2012 Samsung Advanced Institute of Technology, South Korea
"Signal Processing for Next Generation Biosensing and Sequencing"
- 2012 Stanford University, Department of Electrical Engineering
"How to read your genes: Signal Processing for Next Generation Sequencing and Biosensing"
- 2012 University of California Los Angeles, Department of Electrical Engineering
"Signal Processing for Next Generation Sequencing and Biosensing"
- 2012 MIT, Department of Electrical Engineering and Computer Science
"How to Read your Genes: Signal Processing for Next Generation Sequencing and Biosensing"
- 2012 University of Illinois at Urbana-Champaign, Dept. of Electrical and Comp. Engineering
"How to Read your Genes: Signal Processing for Next Generation Sequencing and Biosensing"
- 2012 California Institute of Technology, Department of Electrical Engineering
"Signal Processing for Next Generation Sequencing and Biosensing"
- 2012 University of Southern California, Department of Electrical Engineering
"Signal Processing for Next Generation Sequencing and Biosensing"

Patents

- 2021 A. Hassibi, B. Hassibi, and H. Vikalo, "Methods for detecting analytes," U.S. patent no. 11,001,881
- 2016 A. Hassibi, B. Hassibi, and H. Vikalo, "Multiplex QPCR Arrays," U.S. patent no. 9,458,497
- 2015 B. Hassibi, H. Vikalo and A. Hassibi, "Method and apparatus for detection, identification and quantification of single-and multi-analytes in affinity-based sensor arrays," U.S. patent no. 9,223,929
- 2015 A. Hassibi, B. Hassibi, H. Vikalo and J. L. Riechmann, "Real-time microarrays," U.S. patent no. 9,133,504

Publications

In Review

1. U. Akram, F. Zhang, Y. Li and H. Vikalo, "Diffusion-Based Channel Inpainting for Robust SRS-Based CSI Acquisition," *IEEE Communication Letters*, 2026 (submitted).
2. S. Ahn, H. Chen and H. Vikalo, "FedProTIP: Task-Agnostic Federated Continual Learning via Replay-Free Gradient Projection," *Transactions on Machine Learning Research*, 2026 (submitted).
3. H. Zhang, S. Cha, H. B. Beytur, K. Chan, G. De Veciana and H. Vikalo, "Online Learning for Multi-Layer Hierarchical Inference under Partial and Policy-Dependent Feedback" (submitted).
4. S. Cha, H. Chen, D. Kim, H. Zhang, K. Chan, G. de Veciana and H. Vikalo, "CoreQ: Learning-Free Mismatch Correction and Successive Rounding for Quantization" (submitted).
5. D. Kim, A. de Wynter, H. Chen, H. Kim and H. Vikalo, "Foundation-Preserving Adaptation via Generalized Rayleigh-Quotient Optimization" (submitted).
6. J. Zhu, Y. Ro, J. Robertson, K. Wang, J. Li, H. Vikalo, A. Akella and Z. Wang, "Long-Lived AI Agents Age Too: They Quietly Decay After Deployment" (submitted).
7. D. Kim, H. Zhang, C. Wang and H. Vikalo, "Class-Incremental Learning with Zero-Shot Unlearning via Activation Disentanglement" (submitted).
8. J. Robertson, J. Zhu, H. Vikalo and Z. Wang, "When Is Rank-1 Steering Cheap? Geometry, Granularity, and Budgeted Search" (submitted).

1. U. Akram and H. Vikalo, "Transformers as Implicit State Estimators: In-Context Learning in Dynamical Systems," *Transactions on Machine Learning Research*, April 2026, pp. 2835-8856.
2. A. Aydin, A. G. Aydin, C. Turhan and H. Vikalo, "Managing task offloading in viewport prediction via reinforcement learning," *IEEE Access*, vol. 14, pp. 44043-44054, 2026.
3. J. Robertson, S. Consul, and H. Vikalo, "NextVir: Enabling classification of tumor-causing viruses with genomic foundation models," *PLoS Computational Biology*, 21(8): e1013360, Aug 2025.
4. S. Consul, J. Robertson and H. Vikalo, "XVir: A transformer-based architecture for identifying viral reads from cancer samples," *Journal of Computational Biology*, vol. 32, no. 7, July 2025, pp: 646-658.
5. Y. Chen, A. Hashemi and H. Vikalo, "Accelerated distributed stochastic non-convex optimization over Time-Varying Directed Networks," *IEEE Transactions on Automatic Control*, vol. 70, no. 4, April 2025, pp: 2196-2211.
6. A. G. Aydin and H. Vikalo, "Viewport prediction via adaptive edge offloading," *IEEE Networking Letters*, vol. 7, no. 1, March 2025, pp: 21-25.
7. M. Ribero, H. Vikalo and G. de Veciana, "Federated learning at scale: Addressing client intermittency and resource constraints," *IEEE Journal of Selected Topics in Signal Processing*, vol. 19, no. 1, Jan. 2025, pp: 60-73.
8. S. Consul, Z. Ke and H. Vikalo, "XHap: Haplotype assembly using long-distance read correlations learned by transformers," *Bioinformatics Advances*, vol. 3, no. 1, Nov. 2023, pp: 1-11.
9. M. Ribero and H. Vikalo, "Reducing communication in federated learning via efficient client sampling," *Pattern Recognition*, vol. 148, April 2024, pp. 110122.
10. M. Sakthi, M. Arvinte and H. Vikalo, "Automotive RADAR sub-sampling via object detection networks: Leveraging prior signal information," *IEEE Open Journal of Intelligent Transportation Systems*, vol. 4, November 2023.
11. Z. Ke and H. Vikalo, "Graph-based reconstruction and analysis of disease transmission networks using viral genomic data," *Journal of Computational Biology*, 30(7):796-813, Jun 2023.
12. M. Ribeiro, H. Vikalo and G. de Veciana, "Federated learning under intermittent client availability and time-varying communication constraints," *IEEE Journal of Selected Topics in Signal Processing*, vol. 17, no. 1, January 2023.
13. A. Hashemi, A. Acharya, R. Das, H. Vikalo, S. Sanghavi, and I. Dhillon, "On the benefits of multiple gossip steps in communication-constrained decentralized federated learning," *IEEE Transactions on Parallel and Distributed Systems, Special Section on Parallel and Distributed Computing Techniques for AI, ML, and DL*, vol. 33, no. 11, November 2022.
14. Y. Chen, A. Hashemi and H. Vikalo, "Communication-efficient variance-reduced decentralized stochastic optimization over time-varying directed graphs," *IEEE Transactions on Automatic Control*, vol. 67, no. 12, December 2022.
15. M. Ribero, J. Henderson, S. Williamson, and H. Vikalo, "Federated recommendations using differentially private prototypes," *Pattern Recognition*, vol. 129, September 2022, 108746.
16. A. Hashemi, H. Vikalo, and G. de Veciana, "On the benefits of progressively increasing sampling sizes in stochastic greedy weak submodular maximization," *IEEE Transactions on Signal Processing*, vol. 70, pp. 3978-3992, July 2022.

17. A. Hashemi, R. Shafipour, H. Vikalo, and G. Mateos, "Towards accelerated greedy sampling and reconstruction of bandlimited graph signals," *Signal Processing*, vol. 195, June 2022.
18. Z. Ke and H. Vikalo, "Real-Time Radio Technology and Modulation Classification via an LSTM Auto-Encoder," *IEEE Transactions on Wireless Communications*, vol. 21, no. 1, January 2022.
19. N. M. Arzeno and H. Vikalo, "Evolutionary clustering via message passing," *IEEE Transactions on Knowledge and Data Engineering*, vol. 33, no. 6, June 2021, pp. 2452 - 2466.
20. A. Hashemi, M. Ghasemi, H. Vikalo and U. Topcu, "Randomized greedy sensor selection: Leveraging weak submodularity," *IEEE Trans. on Automatic Control*, vol. 66, no. 1, January 2021, pp. 199-212.
21. A. Sankararaman, H. Vikalo, and F. Baccelli, "ComHapDet: A spatial community detection algorithm for haplotype assembly," *BMC Genomics* vol. 21 (Suppl 9), September 2020, pp. 586:1-14.
22. S. Barik and H. Vikalo, "Matrix completion and performance guarantees for single individual haplotyping," *IEEE Trans. on Signal Processing*, vol. 67, no. 18, September 2019, pp: 4782-4794.
23. A. Hashemi and H. Vikalo, "Evolutionary self-expressive models for subspace clustering," *IEEE Journal of Selected Topics in Signal Processing, Special Issue on Data Science: Robust Subspace Learning and Tracking*, vol. 12, no. 6, December 2018, pp. 1534-1546.
24. S. Barik, S. Das, and H. Vikalo, "Viral quasispecies reconstruction via correlation clustering," *Genomics*, vol. 110, no. 6, November 2018, pp. 375-381.
25. A. Hashemi and H. Vikalo, "Accelerated orthogonal least-squares for large-scale sparse reconstruction," *Digital Signal Processing*, vol. 82, no. 11, November 2018, pp. 91-105.
26. S. Ahn, Z. Ke and H. Vikalo, "Viral quasispecies reconstruction via tensor factorization with successive removal," *Bioinformatics*, vol. 34, no. 13, July 2018, pp. i23-i31.
27. A. Hassibi, A. Manickam, R. Singh, S. Bolouki, R. Sinha, K. Jirage, M. McDermott, B. Hassibi, H. Vikalo, G. Mazarei, L. Pei, L. Bousse, M. Miller, M. Heshami, M. Savage, M. Taylor, N. Gamini, N. Wood, P. Mantina, P. Grogan, P. Kuimelis, P. Savalia, S. Conradson, Y. Li, R. Meyer, E. Ku, J. Ebert, B. Pinsky, G. Dolganov, T. Van, K. Johnson, P. Naraghi-Arani, R. Kuimelis, G. Schoolnik, "Multiplexed identification, quantification and genotyping of infectious agents using a semiconductor biochip," *Nature Biotechnology*, 36, 2018, pp. 738-745.
28. S. Ahn and H. Vikalo, "aBayesQR: A Bayesian method for reconstruction of viral populations characterized by low diversity," *J. of Computational Biology*, vol. 25, no. 7, July 2018, pp: 637-648.
29. H. Yang, J. Chun, and H. Vikalo, "Cyclic block coordinate minimization algorithms for DOA estimation in co-prime arrays," *Signal Processing*, vol. 145, no. 4, April 2018, pp. 272-284.
30. A. Hashemi, B. Zhu and H. Vikalo, "Sparse tensor decomposition for haplotype assembly of diploids and polyploids," *BMC Genomics*, 19(Suppl 4):191, March 2018.
31. H. Si, H. Vikalo, and S. Vishwanath, "Information-theoretic analysis of haplotype assembly," *IEEE Transactions on Information Theory*, vol. 63, no. 7, July 2017, pp: 3468-3479.
32. S. Das and H. Vikalo, "Optimal haplotype assembly via a branch-and-bound algorithm," *IEEE Trans. on Molecular, Biological, and Multi-Scale Comm.*, vol. 3, no. 1, March 2017, pp: 1-12.
33. E. O'Reilly, F. Baccelli, G. de Veciana, and H. Vikalo, "End-to-end optimization of high-throughput DNA sequencing" *Journal of Computational Biology*, 23(10): 789-800, October 2016.
34. C. Cai, S. Sanghavi, and H. Vikalo, "Structured low-rank matrix factorization for haplotype assembly," *IEEE Journal of Selected Topics in Signal Processing, Special Issue on Structured Matrices in Signal and Data Processing*, vol. 10, no. 4, August 2016, pp: 647-657.

35. Z. Puljiz and H. Vikalo, "Decoding genetic variations: Communications-inspired haplotype assembly," *IEEE/ACM Trans. on Comput. Biology and Bioinformatics*, vol. 13, no. 3, June 2016, pp: 518-530.
36. N. M. Arzeno and H. Vikalo, "A novel mortality risk prediction score based on nonlinear feature transformations," *Journal of Biomedical Informatics*, vol. 56, August 2015, pp: 145-156.
37. S. Ahn and H. Vikalo, "Joint haplotype assembly and genotype calling via sequential Monte Carlo algorithm," *BMC Bioinformatics*, 16:223, July 2015, doi:10.1186/s12859-015-0651-8.
38. S. Das and H. Vikalo, "SDhaP: Haplotype assembly for diploids and polyploids via semi-definite programming," *BMC Genomics*, 16:260, April 2015, doi:10.1186/s12864-015-1408-5.
39. N. M. Arzeno and H. Vikalo, "Semi-supervised affinity propagation with soft instance-level constraints," *IEEE Trans. on Pattern Analysis and Machine Intelligence*, vol. 37, no. 5, May 2015, pp: 1041-1052.
40. X. Shen, M. Shamaiah, and H. Vikalo, "Iterative learning of a DNA consensus sequence from high-throughput short reads: Algorithms and limits of performance," *IEEE Transactions on Signal Processing*, vol. 62, no. 17, September 2014, pp: 4425-4435.
41. S. Barik and H. Vikalo, "Sparsity-aware sphere decoding: Algorithms and complexity analysis," *IEEE Transactions on Signal Processing*, vol. 62, no. 9, May 2014, pp: 2212-2225.
42. M. Park, M. Nassar, and H. Vikalo, "Bayesian active learning for drug combinations," *IEEE Transactions on Biomedical Engineering*, 60(11), November 2013, pp: 3248-3255.
43. S. Das and H. Vikalo, "Base calling for high-throughput short-read sequencing: Dynamic programming solutions," *BMC Bioinformatics*, 2013, 14:129 doi:10.1186/1471-2105-14-129.
44. S.-H. Lee, M. Shamaiah, H. Vikalo, and S. Vishwanath, "Message-passing algorithms for coordinated spectrum sensing in cognitive radio networks," *IEEE Communication Letters*, vol. 17, no. 4, April 2013, pp: 812-815.
45. M. Shamaiah, S.-H. Lee, S. Vishwanath, and H. Vikalo, "Distributed algorithms for spectrum access in cognitive radio relay networks," *IEEE Journal on Sel. Areas in Communications - Cognitive Radio Series*, vol. 30, no. 10, November 2012, pp: 1947-1957.
46. T. Wu and H. Vikalo, "Joint parameter estimation and base-calling for pyrosequencing systems," *IEEE Transactions on Signal Processing*, vol. 60, no. 8, August 2012, pp: 4376-4386.
47. M. Shamaiah, S. Banerjee, and H. Vikalo, "Greedy sensor selection under channel uncertainty," *IEEE Wireless Communications Letters*, vol. 1, no. 4, August 2012, pp: 376-379.
48. S. Das and H. Vikalo, "OnlineCall: Fast online parameter estimation and base calling for Illumina's next-generation sequencing," *Bioinformatics*, 2012, doi:10.1093/bioinformatics/bts256.
49. X. Shen and H. Vikalo, "ParticleCall: A particle filter for base calling in next-generation sequencing systems," *BMC Bioinformatics*, vol. 13, no. 160, July 2012.
50. M. Shamaiah, X. Shen, and H. Vikalo, "Estimating parameters of sampled diffusion processes in affinity biosensors," *IEEE Transactions on Signal Processing*, vol. 60, no. 6, June 2012, pp: 3228-3239.
51. M. Shamaiah, S.-H. Lee, and H. Vikalo, "Graphical models and inference on graphs in genomics," *IEEE Signal Processing Magazine*, vol. 29, no. 1, January 2012, pp: 51-65.
52. M. Shamaiah and H. Vikalo, "Estimating time-varying sparse signals under communication constraints," *IEEE Transactions on Signal Processing*, vol. 59, no. 6, June 2011, pp: 2961-2964.
53. H. Vikalo and M. Gokdemir, "An MCMC algorithm for estimation in real-time biosensor arrays," *EURASIP Journal on Advances in Signal Processing, Special Issue on Genomic Signal Processing*, 2010, doi:10.1155/2010/736301.

54. X. Shen and H. Vikalo, "Inferring parameters of gene regulatory networks via particle filtering," *EURASIP Journal on Advances in Signal Processing, Special Issue on Genomic Signal Processing*, 2010, doi:10.1155/2010/204612.
55. H. Vikalo, B. Hassibi, and A. Hassibi, "Limits of performance of quantitative polymerase chain reaction systems," *IEEE Transactions on Information Theory, Special Issue on Molecular Biology and Neuroscience*, vol. 56, no. 2, February 2010, pp: 1-8.
56. M. El-Khamy, H. Vikalo, B. Hassibi, and R. J. McEliece, "Bounds on the performance of sphere decoding of linear block codes," *IEEE Transactions on Communications*, vol. 57, no. 10, October 2009, pp: 2940-2950.
57. A. Hassibi, H. Vikalo, J.-L. Reichmann, and B. Hassibi, "Real-time DNA microarray analysis," *Nucleic Acids Research*, vol. 37, no. 20, 2009, e132:1-12.
58. S. Das, H. Vikalo, and A. Hassibi, "On scaling laws of biosensors," *Journal of Applied Physics*, vol. 105, no. 10, pp. 102021-102021-7, May 2009.
59. H. Vikalo, B. Hassibi, and A. Hassibi, "Modeling and estimation for real-time microarrays," *IEEE Journal of Selected Topics in Signal Processing, Special Issue on Genomic and Proteomic Signal Processing*, vol. 2, no. 3, June 2008, pp: 286-296.
60. F. Parvaresh, H. Vikalo, S. Misra, and B. Hassibi, "Recovering sparse signals using sparse measurement matrices in compressed DNA microarrays," *IEEE Journal of Sel. Topics in Signal Processing, Special Issue on Genomic and Proteomic Signal Processing*, vol. 2, no. 3, June 2008, pp: 275-285.
61. M. Stojnic, H. Vikalo, and B. Hassibi, "An H-infinity design approach to improve the speed of the sphere decoding algorithm," *IEEE Transactions on Signal Processing*, vol. 56, no. 2, Feb. 2008, pp. 712-726.
62. A. Hassibi, H. Vikalo, and A. Hajimiri, "On noise processes and limits of performance in biosensors," *Journal of Applied Physics*, vol. 102, no. 1, July 2007, pp. 014909-014909-12.
63. H. Vikalo and B. Hassibi, "On joint detection and decoding of linear block codes on Gaussian vector channels," *IEEE Transactions on Signal Processing*, vol. 54, no. 9, September 2006, pp. 3330-3342.
64. M. Stojnic, H. Vikalo, and B. Hassibi, "Maximizing the sum-rate of multi-antenna broadcast channels using linear preprocessing," *IEEE Trans. on Wireless Comm.*, vol. 5, no. 9, Sept. 2006, pp. 2338-42.
65. H. Vikalo, B. Hassibi, and P. Stoica, "Joint ML channel estimation and signal detection," *IEEE Transactions on Wireless Communications*, vol. 5, no. 7, July 2006, pp. 1838-1845.
66. H. Vikalo, B. Hassibi, and U. Mitra, "Sphere-constrained ML detection for frequency-selective channels," *IEEE Transactions on Communications*, vol. 54, no. 7, July 2006, pp. 1179-1183.
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Short Bio

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