

Elhanan Borenstein

Department of Genome Sciences, University of Washington
3720 15th Ave NE, Box 355065, Seattle, WA 98195-5065

Phone: 206.685.8165 | Email: elbo@uw.edu

Web: <http://elbo.gs.washington.edu/>

Education

- 2000 – 2006 **Ph.D. with distinction, Computer Science, Tel-Aviv University**
(w/ 1 yr recess) Evolutionary Dynamics of Adaptive Populations: The Effect of Phenotypic Plasticity, Imitation and Culture on Evolution; Prof. Eytan Ruppin's Lab
GPA: 99/100
- 1994 – 1996 **B.Sc. Summa Cum Laude, Physics and Computer Science, Tel-Aviv Univ'**
Program for the Fostering of Excellence (outstanding students special program)
GPA: 97/100

Postgraduate Training

- 2007 – 2009 **Postdoctoral Fellow, Stanford University** (with Prof. Marc Feldman)
- 2007 – 2009 **Omidyar Fellow, Santa Fe Institute** (with Prof. David Krakauer)

Additional Training:

- April 2006 Evolutionary and Ecological Genomics, Ben-Gurion University of the Negev
- Aug-Sep 2005 Research residency, Philip Steinmetz Fellow, Santa Fe Institute
- Jan 2005 Biological Networks & Evolution, The Hebrew Univ', Inst' for Advanced Studies
- Jun 2004 Complex Systems Summer School, Santa Fe Institute

Faculty Positions

- 07/2014 – present **Adjunct Associate Professor**, Computer Science and Engineering, UW
- 07/2014 – present **Associate Professor, Genome Sciences, University of Washington**
- 07/2011 – present **External Professor, Santa Fe Institute**
- 05/2011 – 06/2014 **Adjunct Assistant Professor**, Computer Science and Engineering, UW
- 12/2009 – 06/2014 **Assistant Professor, Genome Sciences, University of Washington**

Additional Affiliations:

- 2017 – present Environmental Pathology/Toxicology training program, UW
- 2014 – present IGERT Program in Big Data and Data Science, UW
- 2010 – present Computational and Molecular Biology Interdisciplinary Program, UW
- 2010 – present Molecular and Cellular Biology Graduate Program, UW

Advisory Boards & Consulting

- 2017 – present **Northwest Institute for Advanced Computing**, Steering Committee
- 2016 – present **Phase Genomics**, Scientific Advisory Board
- 2016 **Celgene**, Consulting
- 2015 – present **Pacific Northwest National Laboratory**, Microbiomes in Transition (MinT), External Advisory Committee

Honors, Fellowships & Awards

2012 – 2017	NIH Director's New Innovator Award (\$1,500,000)
2016	Keynote Speaker, NIH, The 2nd HIV and the Microbiome Workshop
2011 – 2013	Alfred P. Sloan Research Fellowship in molecular biology (two-year fellowship for outstanding early-career scientists, \$50,000)
2012	UW nominee for Pew Biomedical Scholars Award
2011	Keynote speaker, NIH, Annual MSM consortium meeting
2011 – present	Faculty of 1000 Member, Genomics & Genetics
2011	UW nominee for Searle Scholars Program
2007 – 2009	Omidyar Fellowship, Santa Fe Institute
2007 – 2009	Postdoctoral Fellowship, Stanford University
2005	Philip Steinmetz Fellowship, Santa Fe Institute
2005	Best Teacher, School of Computer Science, Tel-Aviv University
2004 – 2006	Yeshaya Horowitz Association Scholarship in Complexity Science (three years research grant for Ph.D. students, \$22,000 per year)
2004	Aharon Katzir Center Training Fellowships for Ph.D., Weizmann Institute
2004	The Don & Sara Marejn Scholarship for Outstanding Ph.D. Student
2004	School of Computer Science Award for Ph.D. Student, Tel-Aviv University
2004	Travel Scholarships: ALIFE9 Student Scholarship, CSSS Scholarship (SFI)
2003	Best Teaching Assistant Prize, School of Computer Science, Tel-Aviv Univ'
1996	B.Sc. Summa Cum Laude (graduated first in the class, GPA 97)
1994 – 1996	Dean's List (3 years)
1994 – 1996	Program for the Fostering of Excellence (Three year merit-based stipend and scholarship, Tel-Aviv University)

Teaching Experience

Regularly obtaining top scores in teaching evaluations with high praises from both students and teaching colleagues. Teaching evaluations available upon request.

Courses Taught - University of Washington:

Spring 2017	GENOME 373: Genomic Informatics; [with D. Fowler] Instructor (responsible for 15 contact hrs/5 weeks)
Winter 2017	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2016	GENOME 373: Genomic Informatics; [with D. Fowler] Instructor (responsible for 15 contact hrs/5 weeks)
Winter 2016	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2015	GENOME 373: Genomic Informatics; [with D. Fowler] Instructor (responsible for 15 contact hrs/5 weeks)
Winter 2015	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)

Spring 2014	GENOME 373: Genomic Informatics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2014	GENOME 541: Intro' to Computational Molecular Biology; [with multiple faculty] Co-instructor (responsible for 3 contact hrs/1 week)
Winter 2014	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2013	GENOME 373: Genomic Informatics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks);
Spring 2013	GENOME 541: Intro' to Computational Molecular Biology; [with multiple faculty] Co-instructor (responsible for 3 contact hrs/1 week)
Winter 2013	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2012	GENOME 373: Genomic Informatics; [with J. Shendure] Instructor (responsible for 15 contact hrs/5 weeks);
Spring 2012	GENOME 541: Intro' to Computational Molecular Biology; [with multiple faculty] instructor (responsible for 3 contact hrs/1 week)
Winter 2012	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2011	GENOME 541: Intro' to Computational Molecular Biology; [with multiple faculty] instructor (responsible for 3 contact hrs/1 week)
Winter 2011	GENOME 559: Intro' to Statistical and Computational Genomics; [with J. Thomas] Instructor (responsible for 15 contact hrs/5 weeks)
Spring 2010	GENOME 541: Intro' to Computational Molecular Biology; [with multiple faculty] instructor (responsible for 3 contact hrs/1 week)

Courses Mentored - University of Washington:

2010-2013 (7 quarters)	CSE 590C: Reading & Research in Computational Biology; [with multiple faculty] Co-mentor
2010- 2011 (2 quarters)	GENOME 580: Ethics in Biomedical Research & Teaching; [with multiple faculty] Guest session leader (2hr)
2010- 2016 (6 quarters)	GENOME 599B: Journal Club Preparation Co-mentor

Courses Taught - Tel-Aviv University:

Spring 2005	Object Oriented Programming Senior Lecturer (responsible for ~42 contact hrs/14 week)
Spring 2004	Object Oriented Programming Senior Lecturer (responsible for ~42 contact hrs/14 week)
Spring 2004	Artificial Life Workshop Teaching Assistant
Summer2003	Object Oriented Programming Lecturer (responsible for ~42 contact hrs/14 week)
Spring 2003	Object Oriented Programming Lecturer (responsible for ~42 contact hrs/14 week)
Spring 2003	Artificial Life Workshop Teaching Assistant
Fall 2004	Software 1

Teaching Assistant

Fall 2003 **Software 1**
Teaching Assistant

Summer Schools, Short Courses, Workshops:

Jul 2017 **Computational Genomics Summer Institute (CGSI)**, UCLA
Lecture of "Model-Based Analysis of Multi-Omic Microbiome Data"

Jun 2017 **Complex Systems Summer School**, Santa Fe Institute
Lecture series on "Analyzing and modeling the human microbiome" (total 3 hrs)

Jun 2016 **Complex Systems Summer School**, Santa Fe Institute
Lecture series on "Model-based analysis of Microbial Communities" (total 3 hrs)

Apr 2014 **Keystone Symposium**, Montana, USA
Workshop on "Reconstructing and Analyzing Metabolic Networks"

Jun 2013 **Complex Systems Summer School**, Santa Fe Institute
Lecture series on "Models of Microbial Communities" (total 3 hrs)

Jun 2012 **Complex Systems Summer School**, Santa Fe Institute
Lecture series on "Modeling Microbial Metabolism" (total 4.5 hrs)

Mentoring and Trainees:Postdoctoral Fellows:

2015 – present **Adrian Verster**
Project: Microbial warfare in the gut microbiome
NSERC Postdoctoral Fellowship

2013 – 2016 **Ohad Manor**
Project: Model-based computational methods for comparative metagenomics

2014 – 2014 **Jody Wright** (now at an environmental stewardship nonprofit organization)
Project: Modeling microbial interaction networks in the human microbiome

2011 – 2015 **Hsuan-Chao Chiu** (now at MediaTek, Taiwan)
Project: Metabolic models of multi-species systems

2011 – 2014 **Rogan Carr** (now at Microsoft)
Project: Computational deconvolution of metagenomic data

Additional Postdoctoral Mentoring:

2014 – 2015 **Attila Kertesz-Farkas**, University of Washington (Career Advisor)

2014 – present **Eric Libby**, Omidyar fellow, Santa Fe Institute (External Advisor)

Ph.D. Students:

2014 – present **Cecilia Noecker**, Genome Sciences, University of Washington
Project: Integrative analysis of multi-meta-omic data
IGERT Big Data Fellowship (2014-present)

2014 – present **Alex Eng**, Genome Sciences, University of Washington
Project: Algorithms for synthetic microbial community design

2014 – present **Colin McNally**, Genome Sciences, University of Washington
Project: Modeling the evolution of mutualism in microbial communities
Genome Training Grant Fellowship (2014-present)

2011 – 2016 **Maximilian Press**, Genome Sciences, University of Washington
Project: Co-Evolutionary inference of microbial genes' function

- 2010 – 2014 **Sharon Greenblum**, Genome Sciences, University of Washington
Project: Systems biology analysis of the human microbiome
Departmental nominee for the UW distinguished thesis award
- 2010 – 2015 **Roie Levy**, Molecular and Cellular Biology, University of Washington
Project: *In-silico* models of species interactions in microbial communities
NSF Graduate Fellowship (2011-2014)

Doctoral and Master's Supervisory Committee:

- 2017 – present **Maria Nelson**, Medical Scientist Training Program, University of Washington
- 2017 – present **Sofiya Shevchenko**, Microbiology, University of Washington
- 2016 – present **Helena van Tol**, Biological Oceanography, University of Washington
- 2016 – present **John E. Lazar**, Genome Sciences, University of Washington
- 2016 – present **Benjamin Fu**, Epidemiology - Public Health, University of Washington
- 2016 – present **Damon H. May**, Genome Sciences, University of Washington
- 2015 – 2016 **Mahsa Khorasani**, Environmental/Forest Sciences, University of Washington
- 2015 – present **Aaron Goodman**, Department of Biological Sciences, Stanford University
- 2015 – present **Moriah Echlin**, Molecular and Cellular Biology, University of Washington
- 2015 – present **Gili Zilberman**, Department of Immunology, Weizmann Institute of Science
- 2015 – present **Timothy Durham**, Genome Sciences, University of Washington
- 2015 – 2015 **Kaitlyn LaCourse**, Microbiology (qualifying exam), University of Washington
- 2014 – present **Daniel Chee**, Genome Sciences, University of Washington
- 2014 – 2014 **Arend Voorman**, Biostatistics, University of Washington
- 2013 – 2016 **Rob Lawrence**, Molecular and Cellular Biology, University of Washington
- 2013 – 2016 **David Young**, Medical Scientist Training Program, University of Washington
- 2011 – 2015 **Katharine Marshall**, School of Oceanography, University of Washington
- 2011 – 2015 **Aaron Miller**, Genome Sciences, University of Washington
- 2011 – 2012 **Samuel Lancaster**, Epidemiology, Fred Hutchinson Cancer Research Center
- 2010 – 2014 **Jennifer Lachowiec**, Molecular and Cellular Biology, University of Washington
- 2010 – 2014 **Aaron Brooks**, Molecular and Cellular Biology, University of Washington
- 2010 – 2012 **Matthew Maurano**, Genome Sciences, University of Washington

Rotation Students:

- Winter 2017 **April Lo**, Genome Sciences, University of Washington
Project: Codon usage bias in the microbiome data
- Spring 2016 **Ken Jean-Baptiste**, Genome Sciences, University of Washington
Project: Characterizing the landscape of 16S data
- Winter 2016 **Sarah Hilton**, Genome Sciences, University of Washington
Project: Co-variation in microbiome strains copy number
- Winter 2014 **Cecilia Noecker**, Genome Sciences, University of Washington
Project: Dynamic co-occurrence network analysis in the gut microbiome
- Autumn 2013 **Seungsoo Kim**, Genome Sciences, University of Washington
Project: Genome-level comparative analysis of the human microbiome
- Autumn 2013 **Colin McNally**, Genome Sciences, University of Washington

- Project: Modeling the evolution of mutualism
- Winter 2013 **Daniel Chee**, Genome Sciences, University of Washington
Project: Metagenomic co-occurrence.
- Spring 2012 **Jorgen Nelson**, Genome Sciences, University of Washington
Project: Genetic co-occurrence networks across metagenomic samples.
- Autumn 2011 **Brandon Blakeley**, Computer Science and Engineering
Project: Functional complementarity in co-occurring microbes.
- Winter 2011 **Benjamin Vernet**, Genome Sciences, University of Washington
Project: Associations between bacterial vaginosis clinical and genomic data.
- Winter 2011 **Maximilian Press**, Genome Sciences, University of Washington
Project: Applying expression profile analysis to study perturbed microbiomes.
- Autumn 2010 **Josh Burton**, Genome Sciences, University of Washington
Project: A framework for metabolic modeling of multi-species systems.
- Autumn 2010 **Alexander Nuttle**, Genome Sciences, University of Washington
Project: Transitions in the developing infant gut microbiome.
- Winter 2010 **Sharon Greenblum**, Genome Sciences, University of Washington
Project: Phylogenetic and functional modularity in microbial communities.
- Winter 2010 **Roie Levy**, Molecular and Cellular Biology, University of Washington
Project: The effect of environmental switching on the evolution of modularity
- Spring 2010 **Caitlin Connelly**, Genome Sciences, University of Washington
Project: Modularity and capacitance in gene interaction networks

Undergraduate Students, Summer Students, and Outreach:

- 2014 – 2015 **Sierra Anderson**, Undergraduate Researcher
Project: A pipeline for comparative metagenomic analysis
- Summer 2014 **Jameson Boslough**, UW Amgen Scholars Program
Project: Species-focused methods for comparative metagenomics
- Spring 2014 **Lane Felker**, Undergraduate Researcher
Project: A software package for metagenomics annotation
- Summer 2013 **Clara Amorosi**, UW Amgen Scholars Program
Project: Scaling laws in the gut microbiome
- Summer 2010 **Lovenoor Aulck**, Genomics Outreach program, University of Washington
Project: Topological properties of gut dwelling microbes

National Service

Editorial Positions:

- 2017 – present **Editorial Board**, *Microbiome*
- 2016 – present **Associate Editor**, *PLOS Computational Biology*
- 2015 **Invited Editor**, *mBio*
- 2013 **Guest Editor**, *PLoS Computational Biology*

Reviewer for Journals:

Science, Cell, PNAS (Proc. Natl. Acad. Sci. USA), Nature Review Genetics, Nature Biotechnology, Nature Microbiology, Nature Methods, Nature Communications, Genetics, Cell Host and Microbe, Genome Biology, Genome Research, Genome Medicine, Cell Metabolism, PLOS Computational Biology, Evolution, Journal of Evolutionary Biology, Microbiology and Molecular Biology Review

(ASM), *International Society for Microbial Ecology Journal (ISMEJ)*, *Scientific Reports*, *Molecular Biology and Evolution*, *Journal of the Royal Society Interface*, *BMC Bioinformatics*, *BMC Systems Biology*, *BMC Evolutionary Biology*, *Evolutionary Ecology*, *Nucleic Acids Research*, *BioSystems*, *Oikos*, *Ecological Modelling*, *Artificial Intelligence*, *Neurocomputing*, *Cognitive Science*, *Artificial Life*, *Bioinformatics*, *Journal of Theoretical Biology*, *Theoretical Population Biology*

Reviewer for Conferences:

The 16th Annual International Conference on Research in Computational Molecular Biology (RECOMB 2012), RECOMB Regulatory Genomics & Systems Biology 2009, Pacific Symposium on Biocomputing (PSB 2009), The 10th Annual International Conference on Research in Computational Molecular Biology (RECOMB 2006), The 14th Annual International conference on Intelligent Systems for Molecular Biology (ISMB 2006), The 15th annual Computational Neuroscience meeting (CNS*2006), The 8th European Conference on Artificial Life (ECAL2005), The 9th International Conference on the Simulation and Synthesis of Living Systems (ALIFE9), The Computational Neuroscience meeting (CNS*2004)

Committees, Study Sections, and Others:

2017	Expert Reviewer , Novo Nordisk Foundation (NNF)
2017	NIH Panel Member . Modeling and Analysis of Biological Systems [MABS] Study Section
2014	Expert Reviewer , Novo Nordisk Foundation (NNF)
2013	Scientific Panel , NIH Human Microbiome Science: Vision for the Future
2013	Session Convener , American Society for Microbiology Meeting (ASM2013)
2013	Ad hoc grant reviewer , Provincia Autonoma di Trento, Grandi Progetti 2012
2012 – present	Authored multiple recommendations , Faculty of 1000, Genomics & Genetics
2012	Program Committee , 16th Annual International Conference on Research in Computational Molecular Biology (RECOMB2012)
2011	NIH/NSF Panel Member . DMS/NIGMS Study Section
2011 – present	Member , Population Modeling Group, Interagency Modeling and Analysis Group & Multi-Scale Modeling Consortium, NIH
2009	Program Committee , 17th annual International Conference on Intelligent Systems for Molecular Biology (ISMB) and 8th European Conference on Computational Biology (ECCB); Bioinformatics of Disease
2005	Program Committee , Memetic Theory in Artificial Systems and Societies (METAS 2005)

University and Departmental Service

2017	Organizer, Wednesdays Evenings at the Genome, Public Lecture Series
2016	Organizer, Wednesdays Evenings at the Genome, Public Lecture Series
2016	Reviewer, UW Center for Ecogenetics & Environmental Health (CEEH)
2015	Member, Curriculum Subcommittee for Graduate Course Revisions
2015	Member, Biomedical Informatics and Medical Education (BIME) Precision Medicine faculty Search
2015	Member, Genome Sciences Seminar Committee
2015	Organizer, Wednesdays Evenings at the Genome, Public Lecture Series
2014	Faculty, University of Washington Amgen Scholars Program

2014	Co-organizer, Wednesdays Evenings at the Genome, Public Lecture Series
2014	Genome Science, graduate program recruiting visits, introductory presentation
2013	Faculty recruitment for a joint GI/AID position, FHCRC
2013	Faculty, University of Washington Amgen Scholars Program
2013	Co-organizer, Genome Science Retreat
2012 – present	Member, UW Cancer Innovation Group
2012	Co-organizer, Genome Science Retreat
2011	Reviewer, University of Washington, Royalty Research Funds
2011	Member, Genome Sciences Seminar Committee
2011	Member, Genome Sciences Faculty Search Committee
2010	Member, Genome Sciences Seminar Committee
2010	Genome Science, graduate program recruiting visits, introductory presentation
2009	Co-organizer, Complex Networks Initiative, Stanford
2008	Co-organizer, CompBio Workgroup, Stanford

Ongoing

2011 – present	Participant in MCB graduate student recruitment
2011 – present	Participant in CSE graduate student recruitment and prospective students visit
2011 – present	Participant in MSTP graduate student recruitment
2011 – present	Lunch presentations to first year students and outreach summer students

Science Communication, Public Lectures, and Outreach:

2017	Public lecture , Bainbridge Island Open Mic Science, Bainbridge, USA
2014 – 2015	Content advisory Committee, Pacific Science Center
2015	Host , Ballard High school students visit
2014	Public lecture , John Von Neumann Public Lecture Series in Complexity & Computation, University of Wisconsin, Madison, USA
2014	Public panel lecture , Conversation about Diabetes, Pacific Northwest Diabetes Research Institute (PNDRI), Seattle, USA
2011	Public lecture , Wednesdays at the Genome, UW, Seattle, USA

Publications - (●): Corresponding author

Journal Papers

1. **Borenstein E** and Ruppin E. Enhancing Autonomous Agents' Evolution with Learning by Imitation. *Journal of Artificial Intelligence and the Simulation of Behavior (AISBJ)*, 1(4), 2003.
2. **Borenstein E** and Ruppin E. The Evolutionary Link between Mirror Neurons and Imitation: An Evolutionary Adaptive Agents Model. *Behavioral and Brain Sciences*, 28:2, 127-128, 2005.
3. (●) **Borenstein E** and Ruppin E. The Evolution of Imitation and Mirror Neurons in Adaptive Agents. *Cognitive Systems Research* (special issue on Epigenetic Robotics), 6(3), 229-242, 2005.
4. (●) **Borenstein E**, Kendal J, and Feldman MW. Cultural Niche Construction in a Metapopulation. *Theoretical Population Biology*, 70(1), 92-104, 2006.

5. (●) **Borenstein E**, Meilijson I, and Ruppin E. The Effect of Phenotypic Plasticity on Evolution in Multipeaked Fitness Landscapes. *Journal of Evolutionary Biology*, 19(5), 1555-1570, 2006.
6. (●) **Borenstein E** and Ruppin E. Direct Evolution of Genetic Robustness in MicroRNA. *PNAS (Proc. Natl. Acad. Sci. USA)*, 103(17), 6593-6598, 2006.
7. (●) **Borenstein E***, Shlomi T*, Ruppin E, and Sharan R (*equal contribution). Gene Loss Rate: A Probabilistic Measure for the Conservation of Eukaryotic Genes. *Nucleic Acids Research*, 35(1), e7, 2007.
8. **Borenstein E**, Feldman MW, and Aoki K. Evolution of Learning in Fluctuating Environments: When Selection Favors Both Individual and Social Learning. *Evolution*, 62 (3), 586–602, 2008.
9. Kreimer A*, **Borenstein E***, Gophna U, and Ruppin E (*equal contribution). The Evolution of Modularity in Bacterial Metabolic Networks. *PNAS (Proc. Natl. Acad. Sci. USA)*, 105(19), 6976-6981, 2008.
10. Lehmann L, Foster KR, **Borenstein E**, and Feldman MW. Social and Individual Learning of Helping in Humans and Other Species. *Trends in Ecology & Evolution*, 23(12), 664-671, 2008.
11. (●) **Borenstein E**, Kupiec M, Feldman MW, and Ruppin E. Large-Scale Reconstruction and Phylogenetic Analysis of Metabolic Environments. *PNAS (Proc. Natl. Acad. Sci. USA)*, 105(38), 14482-14487, 2008.
 - Reviewed in Genome Biology, 9:239, 2008 (see media coverage)
 - Recommended by Faculty of 1000 Biology
 - Featured in Astrobiology Magazine, Science 2.0
12. (●) **Borenstein E** and Krakauer DC. An End to Endless Forms: Epistasis, Phenotype Distribution Bias, and Non-Uniform Evolution. *PLoS Computational Biology*, 4(10), 2008.
 - Featured in Nature Reviews Genetics, 9(12), 2008 (see media coverage)
13. (●) **Borenstein E** and Feldman MW. Topological Signatures of Species Interactions in Metabolic Networks. *Journal of Computational Biology*, 16(2), 191-200, 2009.
 - Featured in Stanford Report, Feb 25, 2009 (see media coverage)
 - Reviewed in Frontiers in Ecology & the Environment, April, 2009
14. Cai J, **Borenstein E**, Chen R, and Petrov DA. Similarly Strong Purifying Selection Acts on Human Disease Genes of All Evolutionary Ages. *Genome Biology and Evolution*, 1(1), 131-144, 2009.
15. Freilich S, Kreimer A, **Borenstein E**, Yosef N, Sharan R, Gophna U, and Ruppin E. Metabolic-Network Driven Analysis of Bacterial Ecological Strategies. *Genome Biology*, 10(6), 2009.
16. Freilich S, Kreimer A, **Borenstein E**, Gophna U, Sharan R, and Ruppin E. Decoupling Environment-Dependent and Independent Genetic Robustness across Bacterial Species. *PLoS Computational Biology*, 6(2), 2010.
17. Cai J, **Borenstein E**, and Petrov DA. Broker Genes in Human Disease. *Genome Biology and Evolution*, 2, 815-825, 2010.
18. (●) Carr R and **Borenstein E**. NetSeed: A network-based reverse-ecology tool for calculating the metabolic interface of an organism with its environment. *Bioinformatics*, 28(5), 734-735, 2012.
19. Pinho R, **Borenstein E**, and Feldman MW. Most networks in Wagner's model are cycling. *PLoS ONE*, 7(4), e34285, 2012.
20. (●) Greenblum S, Turnbaugh P, and **Borenstein E**. Metagenomic Systems Biology of the Human Gut Microbiome Reveals Topological Shifts Associated with Obesity and IBD. *PNAS (Proc. Natl. Acad. Sci. USA)*, 109(2), 594-599, 2012.

- Featured in This Week in PNAS, "Obesity, IBD-related topologies in the human gut microbiome"
 - Featured in Nature Reviews Microbiology, 10(674), 2012, Genome Watch
 - Featured in UW Today
 - Featured in Microbe Magazine, American Society for Microbiology
 - Featured in The Seattle Times
21. Kreimer A, Doron A, **Borenstein E**, and Freilich S. NetCmpt: A network-based tool for calculating the metabolic competition between bacterial species. *Bioinformatics*, 28 (16), 2195-2197, 2012.
22. (●) Levy R and **Borenstein E**. Reverse Ecology: From systems to environments and back. *Advances in Experimental Medicine and Biology*, 751 (Evolutionary Systems Biology), 751, 329-345, 2012.
23. O'Roak BJ, Vives L, Girirajan S, Karakoc E, Krumm N, Coe BP, Levy R, Ko A, Lee C, Smith JD, Turner EH, Stanaway IB, Vernot B, Malig M, Baker C, Reilly B, Akey JM, **Borenstein E**, Rieder MJ, Nickerson DA, Bernier R, Shendure J, and Eichler EE. Sporadic autism exomes reveal a highly interconnected protein network of de novo mutations. *Nature*, 485, 246-250, 2012.
- Featured in Nature Reviews Genetics - see Fruits of exome sequencing for autism
 - Featured in The New York Times front page (see media coverage)
 - Featured in UW Today, "Autism mutations, scattered across genes, merge into network of interactions"
 - Recommended by Faculty of 1000 Biology
24. (●) **Borenstein E**. Computational systems biology and in silico modeling of the human microbiome. *Briefings in Bioinformatics*, 13(6), 769-780, 2012.
- Recommended by Faculty of 1000 Biology
25. (●) Neph S, Stergachis AB, Reynolds A, Sandstrom R, **Borenstein E**^{*}, and Stamatoyannopoulos JA[‡] (‡ co-corresponding authors). Circuitry and dynamics of human transcription factor regulatory networks. *Cell*, 150(6), 1274-1286, 2012.
26. O'Roak BJ, Vives L, Fu W, Egertson JD, Stanaway IB, Phelps IG, Carvill G, Kumar A, Lee C, Ankenman K, Munson J, Hiatt JB, Turner EH, Levy R, O'Day DR, Krumm N, Coe BP, Martin BK, **Borenstein E**, Nickerson DA, Mefford HC, Doherty D, Akey JM, Bernier R, Eichler EE, and Shendure J. Multiplex Targeted Sequencing Identifies Recurrently Mutated Genes in Autism Spectrum Disorders. *Science*, 338, 1619-1622, 2012.
27. (●) Greenblum S, Chiu HC, Levy R, Carr R, and **Borenstein E**. Towards a Predictive Systems-Level Model of the Human Microbiome: Progress, Challenges, and Opportunities. *Current Opinion in Biotechnology*, 24, 810-820, 2013.
28. (●) Press MO^{*}, Li H^{*}, Creanza N, Kramer G, Queitsch C, Sourjik V, and **Borenstein E**. Genome-Scale Co-Evolutionary Inference Identifies Functions and Clients of Bacterial Hsp90. *PLoS Genetics*, 9(7), 2013.
29. (●) Levy R, and **Borenstein E**. Metabolic Modeling of Species Interaction in the Human Microbiome Elucidates Community-Level Assembly Rules, *PNAS (Proc. Natl. Acad. Sci. USA)*, 110(31), 12804-12809, 2013.
- Featured in UW Today
 - Recommended by Faculty of 1000 Biology
 - Highlighted in Nature Medicine (Microbiome models, on computers and in lab dishes, see progress)

30. (●) Carr R, Shen-Orr SS, and **Borenstein E**. Reconstructing the Genomic Content of Microbiome Taxa through Shotgun Metagenomic Deconvolution, *PLoS Computational Biology*, 9(10), 2013.
 - Recommended by Faculty of 1000 Biology
31. Hoffman LR, Pope CE, Hayden HS, Levy R, McNamara S, Jacobs MA, Rohmer L, Radey M, Heltshel SL, Ramsey BW, Brittnacher MJ, **Borenstein E**, and Miller SI. Escherichia Coli Dysbiosis Correlates with Gastrointestinal Dysfunction in Children with Cystic Fibrosis. *Clinical Infectious Diseases*, 58(3), 396-399, 2014.
32. (●) Levy R and **Borenstein E**. Metagenomic systems biology and metabolic modeling of the human microbiome: From species composition to community assembly rules, *Gut Microbes*, 5(2), 2014.
33. (●) Chiu HC, Levy R, and **Borenstein E**. Emergent Biosynthetic Capacity in Simple Microbial Communities. *PLoS Computational Biology*, 10(7), 2014.
 - Featured in Santa Fe Institute News
34. (●) Manor O, Levy R, and **Borenstein E**. Mapping the inner workings of the microbiome: Genomic- and metagenomic-based study of metabolism and metabolic interactions in the human microbiome. *Cell Metabolism*, 20(5), 742-752, 2014.
 - Featured in *Genetic Engineering & Biotechnology News* (GEN)
35. (●) Carr R and **Borenstein E**. Comparative Analysis of Functional Metagenomic Annotation and the Mappability of Short Reads. *PLoS ONE*, 9(8), 2014.
36. Stergachis A, Neph A, Sandstrom R, Haugen E, Reynolds AP, Zhang M, Byron R, Canfield T, Stehling-Sun S, Lee K, Thurman RE, Vong S, Bates D, Neri F, Diegel M, Giste E, Dunn D, Vierstra J, Hansen R, Johnson AK, Sabo PJ, Wilken MS, Reh T, Treuting PM, Kaul R, Groudine M, Bender MA, **Borenstein E**, and Stamatoyannopoulos JA. Conservation of trans-acting networks during mammalian regulatory evolution. *Nature*, 515, 365-370, 2014.
 - Featured in Nature News & Views
 - Featured in Science Daily
37. (●) Lachowiec J, Lemus T, **Borenstein E**^{*}, and Queitsch C^{*} (¥ co-corresponding authors). Hsp90 promotes kinase evolution. *Molecular Biology and Evolution*, 32(1), 91-99, 2015.
38. Waldor MK, Tyson G, **Borenstein E**, Ochman H, Moeller A, Finlay BB, Kong HH, Gordon JL, Nelson KE, Dabbagh K, and Smith H. Where next for microbiome research? *PLoS Biology*, 13(1), 2015.
 - Featured in *Genetic Engineering & Biotechnology News* (GEN)
39. Hormozdiari F, Penn O, Borenstein E, and Eichler E. The discovery of integrated gene networks for autism and related disorders. *Genome Research*, 25(1), 142-154, 2015.
40. (●) Manor O, and **Borenstein E**. MUSiCC: A marker genes based framework for metagenomic normalization and accurate profiling of gene abundances in the microbiome. *Genome Biology*, 16:53, 2015.
 - Featured in *Genetic Engineering & Biotechnology News* (GEN)
41. (●) Greenblum S, Carr R, and **Borenstein E**. Extensive strain-level copy number variation across human gut microbiome species. *Cell*, 160, 583-594, 2015.
 - Main feature on **Cell cover**
 - Altmetric score: 108 (in 99th percentile compared to all articles; in 98th percentile compared to all articles in Cell)
 - Featured in UW NewsBeat
 - Featured in *Genomeweb*

- Featured in Science Daily
 - Featured in Biomedical Computation Review (Computing the Gut)
 - Featured in IEEE Pulse Magazine (Deciphering the Mysterious Microbiome)
42. (●) Levy R, Carr R, Kreimer A, Freilich S, and **Borenstein E**. NetCooperate: A network-based tool for inferring host-microbe and microbe-microbe cooperation. *BMC Bioinformatics*, 16:164, 2015.
43. (●) Noecker C and **Borenstein E**. Getting personal about nutrition. *Trends in Molecular Medicine*, 22(2), 83-85, 2016.
44. (●) Noecker C, Eng A, Srinivasan S, Theriot CM, Young VB, Jansson JK, Fredricks DN, and **Borenstein E**. Metabolic model-based integration of microbiome taxonomic and metabolomic profiles elucidates mechanistic links between ecological and metabolic variation. *mSystems*, 1:1, e00013-15, 2016.
- mSystems Editor's Pick
 - Featured in IEEE Pulse Magazine (Deciphering the Mysterious Microbiome)
45. (●) Manor O, Levy R, Pope CE, Hayden HS, Brittnacher MJ, Carr R, Radey MC, Hager KR, Heltshel SI, Ramsey BW, Miller SI, Hoffman LR*, and **Borenstein E*** (¥ co-corresponding authors). Metagenomic evidence for taxonomic and functional dysbiosis in children with cystic fibrosis. *Scientific Reports*, 6:22493, 2016.
46. (●) Eng A and **Borenstein E**. An algorithm for designing minimal microbial communities with desired metabolic capacities. *Bioinformatics*. 32(13), 2016.
- Featured in IEEE Pulse Magazine (Deciphering the Mysterious Microbiome)
47. (●) Press M, Queitsch C, **Borenstein E**. Evolutionary assembly patterns of prokaryotic genomes. *Genome Research*. 26(6), 826-833, 2016.
48. May DH, Timmins-Schiffman E, Mikan MP, Harvey HR, **Borenstein E**, Nunn BL, and Noble WS. An alignment-free 'metapeptide' strategy for metaproteomic characterization of microbiome samples using shotgun metagenomic sequencing. *Journal of Proteome Research*. 15(8), 2697-2705, 2016.
49. (●) Noecker C, McNally C, Eng A, and **Borenstein E**. High-Resolution and Accurate Characterization of the Human Microbiome. *Translational Research*. 179, 7-23, 2017.
50. Snijders AM, Langley SA, Kim YM, Brislawn CJ, Noecker C, Zink EM, Fansler SJ, Casey CP, Miller DR, Huang Y, Karpen GH, Celniker SE, Brown JB, **Borenstein E**, Jansson JK, Metz TO, and Mao JH. Influence of early life exposure, host genetics and diet on the mouse gut microbiome and metabolome. *Nature Microbiology*. 2, 16221, 2017.
- Featured in Science Daily
 - Features in PNNL News
51. Mosites E, Sammons M, Otiang E, Eng A, Noecker C, Manor O, Hilton S, Mwangi T, Onyango C, Garland-Lewis G, Call DR, Njenga MK, Wasserheit J, Zambriski J, Walson J, Palmer G, Montgomery J, **Borenstein E**, Omere R, and Rabinowitz P. Microbiome Sharing Between Children, Livestock and Household Surfaces in Western Kenya. *PLOS ONE*, 12(2), 2017.
52. (●) Manor O, and **Borenstein E**. Revised computational metagenomic processing uncovers hidden and biologically meaningful functional variation in the human microbiome. *Microbiome*. 5:19, 2017.
53. (●) Manor O and **Borenstein E**. Systematic characterization and analysis of taxonomic drivers of functional shifts in the human microbiome. *Cell Host and Microbe*. 21, 254-267, 2017.
- Main feature on **Cell Host and Microbe cover**
 - Featured in **Science**, 356,6339, 2017; Mining microbes: Creating genomic tools to fight disease

- Features in UW The Daily
- Featured in UW NewsBeat
- Featured in TechCrunch
- Featured in Business Standard
- Featured in FierceBiotech
- Featured in UW Accelerate
- Recommended by Faculty of 1000 Biology

54. Whitney JC, Peterson SB, Kim J, Pazos M, Verster AJ, Radey MC, Kulasekara HD, Ching MQ, Bryant D, Goo YA, Surette MG, **Borenstein E**, Vollmer W, and Mougous JD. A Broadly Distributed Toxin Family Mediates Contact-Dependent Antagonism Between Gram-positive Bacteria. *eLife*. 6:e26938, 2017.
55. (●) Verster AJ, Ross BD, Radey MC, Bao Y, Goodman A, Mougous JD, and **Borenstein E**. The Landscape of Type VI Secretion across Human Gut Microbiomes Reveals its Role in Community Composition. *Cell Host and Microbe*. In press.
- Featured in This Week in Microbiology (ASM Podcast), Covert Pathogenesis

Submitted

1. (●) McNally C and **Borenstein E**. Metabolic Model-Based Analysis of the Emergence of Bacterial Cross-Feeding through Extensive Gene Loss.

In Preparation (selected)

1. (●) Chiu HC and **Borenstein E**. A gut microbiome-scale metabolic model for designing precise dietary interventions.
2. (●) Eng A and **Borenstein E**. Taxa-function relationship and the function-level effects of taxonomic perturbations in microbial communities.
3. (●) Eng A, Noecker C, McNally C, and **Borenstein E**. An integrative microbiome browser for linking community composition and function.
4. (●) Verster A and **Borenstein E**. Clade-specific assembly rules of the microbiome.
5. (●) Noecker C and **Borenstein E**. A model-based analysis of taxonomic drivers of metabolomic shifts in the microbiome.

Conference Papers and Other Publications

1. **Borenstein E** and Ruppín E. Enhancing Autonomous Agents' Evolution with Learning by Imitation. *Proceedings of the AISB 2003 Convention: Cognition in Machines and Animals*, 2003.
2. **Borenstein E** and Ruppín E. Evolving Imitating Agents and the Emergence of a Neural Mirror System. *Proceedings of the Ninth International Conference on the Simulation and Synthesis of Living Systems (ALife IX)*, MIT Press, 146-151, 2004.
3. **Borenstein E** and Cline E. Cellular automata model of cystogenesis and tubulogenesis. *Proceedings of the Santa-Fe Institute Complex Systems Summer School*, June 2004.
4. **Borenstein E**, Kupiec M, Feldman MW and Ruppín E. Large-Scale Reconstruction and Analysis of Growth Environments. *Proceedings of the Eighth International Conference on Systems Biology (ICSB 2007)*, 2007.
5. **Borenstein E** and Feldman MW. Topological Signatures of Species Interactions in Metabolic Networks. *4th Annual RECOMB Satellite on Systems Biology*, 2008.

Abstracts and Posters

Upon request

Invited Talks

- Jun 2018
(upcoming) The International Human Microbiome Congress, Killarney, Ireland
- Apr 2018
(upcoming) Microbiome: Hype and Hope, Fred Hutchinson Cancer Research Center, Seattle, USA
- Mar 2018
(upcoming) The Microbiome Seminar Series, NYU Langone Medical Center, NY, USA
- Mar 2018
(upcoming) Quantitative & Computational Biology, Princeton's Lewis-Sigler Institute for Integrative Genomics, Princeton, USA
- Dec 2017
(upcoming) Biomedical Big Data Science: Vision, Promises and Challenges, Haifa, Israel
- Oct 2017
(upcoming) 116th International TITISEE Conference, "From pathogen evolution to microbiome dynamics", Black Forest, Germany
- Aug 2017 **NIH special workshop**, "The Human Microbiome: Emerging Themes at the Horizon of the 21st Century", Bethesda, USA
- Aug 2017 Ecological Society of America (ESA) meeting, Investigating structure-function relationships in microbial communities, Portland, USA
- Jul 2017 Society for Industrial Microbiology and Biotechnology Annual Meeting, Denver, USA
- June 2017 The Barcelona Debates on the Human Microbiome. From Microbes to Medicines, Barcelona, Spain
- May 2017 HIV Mucosal Immunology Group Annual Meeting, Seattle, USA
- May 2017 Rush University, Department of Immunity and Emerging Pathogens, Chicago, USA
- May 2017 The Novo Nordisk Foundation conferences, "Data-driven Biotechnology", Copenhagen, Denmark
- Mar 2017 Bainbridge Island's Open Mic Science, Bainbridge, USA
- Nov 2016 **Keynote Speaker:** The 2nd HIV and the Microbiome Workshop, NIH, Bethesda, USA
- Oct 2016 Individualizing Medicine 2016 Conference, Mayo Clinic, Rochester, MN, USA
- Jun 2016 Vanderbilt Genetics Institute, Vanderbilt University, Nashville, USA
- May 2016 Bioinformatics for the Microbiome Symposium, Stanford University, Stanford, USA
- Apr 2016 Vancouver Bioinformatics Users Group (VanBUG) Seminar Series, Vancouver, Canada
- Mar 2016 Genome Sciences Combi Seminars, University of Washington, Seattle, USA
- Jan 2016 Gut Microbiome Symposium, FHCRC, Seattle, USA
- Nov 2015 25th Annual Beckman Symposium on Microbiota in Health in Disease, Duarte, USA
- Oct 2015 SRI International, Bioinformatics Research Group, Menlo Park, USA
- Sep 2015 Cell Symposium on *Human Immunity and the Microbiome in Health and Disease*, Montreal, Canada
- Sep 2015 Multi-omics for Microbiomes Conference, Pacific Northwest National Laboratory, Kennewick, USA
- Jun 2015 CF Seminar Series, University of Washington, Seattle, USA
- Jun 2015 Gladstone Institutes, UCSF, San Francisco, USA
- May 2015 Center for Bioinformatics and Computational Biology, University of Maryland, College

Park, USA

- Mar 2015 COBRA Workshop on Modelling Microbial Communities, Luxemburg
- Mar 2015 The American Society for Biochemistry and Molecular Biology (ASBMB) Annual Meeting, Boston, USA
- Mar 2015 Weizmann Institute of Science Systems Biology Conference, “two2many”, Rehovot, Israel
- Feb 2015 AAAS Symposium on Obesity and Microbiome, Jan Jose, CA, USA
- Feb 2015 Math Model Affinity Group, Fred Hutchinson Cancer Research Center, Seattle, USA
- Dec 2014 ECOFECT international colloquium on modeling infectious diseases (EMOTIONS 2014), Lyon, France
- Nov 2014 Conversation about Diabetes, Public Panel Talk, Pacific Northwest Diabetes Research Institute (PNDRI), Seattle, USA
- Nov 2014 Urban Ecology Research Laboratory, Department of Urban Design and Planning, University of Washington, Seattle, USA
- Oct 2014 Diabetes and the Microbiome Research Symposium, organized by ADA and JDRF, Washington, DC, USA (canceled)
- Sep 2014 ASM Conference on Beneficial Microbes, Washington, DC, USA
- Sep 2104 **John Von Neumann Public Lecture Series in Complexity & Computation**, University of Wisconsin, Madison, USA
- Aug 2014 META Center Symposium on Host-Microbe Systems Biology, *Modeling Our Microbial Selves*, University of Oregon, Eugene, USA
- May 2014 **Keynote Speaker:** Inland Northwest Genomics Research Symposium (INWGRS), Moscow, ID, USA
- May 2014 UW Diabetes and Metabolism Symposium, Seattle, USA
- Apr 2014 Keystone Symposium, Exploiting and Understanding Chemical Biotransformations in the Human Microbiome, Montana, USA
- Feb 2014 Lecture Series in Biomedical & Health Informatics, University of Washington, Seattle, USA
- Feb 2014 **The Henk van Verveeld lecture, Nature of Life (NOL) lecture series**, Department of Ecological Sciences, VU Amsterdam University
- Feb 2014 Genome Sciences Combi Seminars, University of Washington, Seattle, USA
- Jan 2014 University of Missouri, Department of Biological Sciences, Columbia, USA
- Nov 2013 Northwest Branch of the American Society for Microbiology meeting, Seattle, USA
- Aug 2013 Cystic Fibrosis Annual Retreat, University of Washington, Seattle, USA
- Jul 2013 IUPS Satellite Meeting on Multiscale Systems Biology, Buckinghamshire, UK
- May 2013 Mathematical tools for evolutionary systems biology, Banff International Research Station, Banff, Canada
- May 2013 American Society for Microbiology General Meeting (ASM2013), Denver, USA
- May 2013 Cell Symposium on *Microbiome and Host Health*, Lisbon, Portugal
- Feb 2013 Stanford University, Stanford, USA
- Feb 2013 VA Palo Alto Health Care System, Palo Alto, USA
- Nov 2012 META Center for Systems Biology, University of Oregon, Eugene, USA
- Jul 2012 Hopkins Marine Station Microbiology Course, Stanford University, Pacific Grove, USA

Jun 2012	Santa Fe Institute, Santa Fe, USA
May 2012	Diabetes and metabolism seminar series, Diabetes Research Center, UW, Seattle, USA
Nov 2011	Institute of Ecology and Evolution (IE ²), University of Oregon, Eugene, USA
Oct 2011	Keynote Speaker: Multiscale Modeling Consortium Meeting, NIH, Rockville, USA
Jul 2011	<i>Wednesdays at the Genome</i> (Public Lecture Series), UW, Seattle, USA
May 2011	The 5th Computational Molecular Biology Spring Symposium, UW, Seattle USA
Mar 2011	AACR-NCI Conference on Systems Biology: Confronting the Complexity of Cancer, La Jolla, USA
Dec 2010	Next Generation Sequencing seminar, Department of Biology, UW, Seattle USA
Jun 2010	Computational Biology Seminar Series, Fred Hutchinson Cancer Research Center, Seattle, USA
May 2010	Genome Sciences Seminars, University of Washington, Seattle, USA
Jul 2009	Complex Networks Initiative, Stanford University, USA
Apr 2008	Workshop on <i>The Role of Variation in Cultural Change</i> , Santa-Fe Institute, USA
Jul 2007	Santa-Fe Institute, Santa Fe, USA
Nov 2006	Department of Computer Science (<i>Barash Group</i>), Ben-Gurion University, Israel
Jun 2006	Department of Biological Chemistry (<i>Tawfik Group</i>), Weizmann Institute of Science, Israel
Apr 2006	<i>Camp Evolution II: Evolutionary and Ecological Genomics</i> , Blaustein Institute for Desert Research, Ben-Gurion University of the Negev, Israel

Contributed Talks

Apr 2015	International Human Microbiome Congress, Luxemburg
Mar 2011	International Human Microbiome Congress, Vancouver, Canada
Nov 2010	6th Annual RECOMB Satellite on Systems Biology, New York, USA
Dec 2009	5th Annual RECOMB Satellite on Systems Biology, Boston, USA
Oct 2008	4th Annual RECOMB Satellite on Systems Biology, Boston, USA
Oct 2007	Biomedical Computation at Stanford (BCATS), Stanford, USA
Oct 2007	Plenary talk: The 8 th Int'l Conference on Systems Biology, Long Beach, USA
Sep 2004	The 9th Int'l Conference on the Simulation and Synthesis of Living Systems, Boston, USA
Apr 2003	The AISB 2003 Convention: Cognition in Machines and Animals, Aberystwyth, Wales

Media Coverage

- Mining microbes: Creating genomic tools to fight disease, *Science*, 356, 6339, May 2017
- Covert Pathogenesis, *This Week in Microbiology* (ASM Podcast), May 2017
- FishTaco and Your Microbiome, *UW Accelerate*, February 2017
- Gut bacteria studies could combat associated imbalances like diabetes, *UW The Daily*, February 2017
- What's Up Bainbridge (podcast): Open Mic Science at the Treehouse with UW Professor Elhanan Borenstein on the Microbial Zoo, February 2017

- FishTaco can analyze your microbiome before or after you eat a fish taco, TechCrunch, January 2017
- A new method reveals roles different bacteria play in microbiome imbalances linked to disease, UW News Beat, January 2017
- Genes, early environment sculpt the gut microbiome, Science Daily, November 2016
- Deciphering the Mysterious Microbiome, IEEE Pulse Magazine, September 2016
- External faculty highlight - Matrix, The newsletter for Santa Fe Institute researchers, May 2016
- Computing the gut, Biomedical Computation Review, April 2016
- Metagenomics Digs Up Microbiome Riches, Genetic Engineering and Biotechnology News (GEN), October 2015
- Microbiome models, on computers and in lab dishes, see progress, Nature Medicine, June 2015
- Analysis Highlights Strain Diversity in Gut Microbiome, GenomeWeb, January 2015
- Among gut microbes, strains, not just species, matter, UW News Beat, January 2015
- Mice in the ENCODE spotlight, Nature News & Views, November 2014
- New view of mouse genome finds many similarities, striking differences with human genome, Science Daily, November 2014
- When microbes join forces, useful new compounds emerge, Santa Fe Institute News, July 2014
- Ecological forces structure your body's personal mix of microbes, UW Today, July 2013
- Metagenomics with guts, Genome Watch, Nature Reviews Microbiology, 10(674), Oct 2012
- Digestive microbes work differently in fat, lean, The Seattle Times, Feb 2012
- Fruits of exome sequencing for autism, Nature Review Genetics, 13(6), 2012
- Autism mutations, scattered across genes, merge into network of interactions, UW Today, 2012
- Scientists link gene mutation to autism risk, The New York Times (front page), April 2012
- Audio Interview, *Microbe Magazine*, March 2012
- Gut microbe networks differ from norm in obese people, systems biology approach reveals, UW Today & Science Daily, 2012
- Reverse Ecology, Astrobiology Magazine, March, 2009
- Reconstructing Bacterial Environments From Millions Of Years Ago, Science 2.0, Feb 2009
- Reverse ecology uses genes to predict environment, *Frontiers in Ecology & the Environment*, 2009
- Reversing ecology reveals ancient environments, *Stanford Report & Science Daily*, 2009
- Network-based approaches for linking metabolism with environment, *Genome Biology*, 9:239, 2008
- Modelling the evolutionarily possible, *Nature Reviews Genetics*, 9(11), 2008
- Development puts an end to the evolution of endless forms, *Science Daily*, 2008
- Santa Fe Institute researcher figures out factors that make bacteria more modular, *Medical News Today*, 2008
- Factors that make bacteria more modular detailed, *Science Daily*, 2008

Professional Industry Positions

Extensive professional experience in the IDF and hi-tech industry, holding top professional and management positions.

2000 – 2001	Co-Founder and Chief Technology Officer, Gate42 Technologies Ltd.
1997 – 1999	Vice President Research & Development, Veon Ltd. (acquired by Philips Electronics in 2000)
1995 – 1996	Software Project Leader, Veon Ltd.
1993 – 1994	Software engineer, Efrat Technologies (now Comverse)
Military service	Lieutenant, Section head, HM"N Talpiot (an elite R&D unit, Israel Defense Forces)

Patents (with other co-inventors)

- System and method and linking information to a video (US6570587)
- Linking information to and accessing information from a video (EP0922259,WO1998004984)
- System and method for guided media sequences (WO0203247; application)
- Streaming hypervideo and dynamic hypervideo (WO9910822, EP1005680)