

**SHAI CARMİ****PERSONAL**

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Born: December 25, 1981, Ramat-Gan, Israel  
Family status: married + 2  
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**CURRENT OCCUPATION**

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2015- Senior lecturer (equivalent of assistant professor)  
Braun School of Public Health and Community Medicine  
The Faculty of Medicine  
The Hebrew University of Jerusalem, Israel (HUJI)

**EDUCATION**

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2005 B.Sc., Bar-Ilan University, Israel, Physics and Computer Science, summa cum laude  
2006 M.Sc., Bar-Ilan University, Israel, Physics, accelerated track, magna cum laude  
Advisor: Prof. Shlomo Havlin.  
Thesis title: Statistical physics of complex networks  
2010 Ph.D., Bar-Ilan University, Israel, Physics, with distinction  
Advisor: Prof. Shlomo Havlin  
Thesis title: Complex systems from communication networks to proteins:  
statistical analysis and modeling

**PAST ACADEMIC POSITIONS**

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2010-2011 Post-doctoral fellow, The Faculty of Life Sciences, Bar-Ilan University  
Mentor: Dr. Erez Levanon  
Topic: RNA and DNA editing in mammalian genomes  
2011-2015 Post-doctoral scientist, Department of Computer Science, Columbia University  
Mentor: Prof. Itsik Pe'er  
Topics: Ashkenazi genetics; genetic sharing in founder populations

**TEACHING EXPERIENCE**

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2005-2010 Teaching assistant, Department of Physics, Bar-Ilan University  
Courses:

- Introductory physics laboratory
- Introductory physics for the life sciences
- Thermodynamics and statistical mechanics

2016- Lecturer, School of Public Health, The Hebrew University of Jerusalem  
Courses taught alone:

- Statistical methods for biomedical sciences
- Advanced topics in data analysis for public health and medicine
- Personal genomics

Courses taught with others:

- Statistical methods for medicine
- Statistical methods for public health

- Academic and research skills
- Courses coordinated (alone):
- Introduction to R for biostatistics

## FELLOWSHIPS, HONORS, AND AWARDS

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### Undergraduate

- 2001, 2002 The Open University president's list (as a non-degree student)  
 2003, 2004, 2005 Bar-Ilan University dean's list  
 2005 Wolf Foundation award (national)

### Graduate

- 2005 Bar-Ilan University rector's award  
 2007 Bar-Ilan University president's fellowship  
 2007 Adams fellowship of The Israel Academy of Sciences and Humanities  
 2007 'Interdisciplinary Technologies' fellowship of The Council for Higher Education of Israel (declined)  
 2008 Bar-Ilan University dean's award  
 2009 Wolf Foundation award (national)

### Post-doctoral

- 2011 Human Frontier Science Program Cross Disciplinary Fellowship (international)  
 2015 Alon fellowship of The Council for Higher Education of Israel (declined)

### Principal investigator

- 2018 Work covered in Science magazine's 10 breakthroughs of 2018  
 2019 Hebrew University Faculty of Medicine research award

## PUBLICATIONS

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Overall citations: 2600; *h-index*: 23; *i10-index*: 37 (Google Scholar)

### Refereed Journals

1. **S. Carmi**, E. Y. Levanon, S. Havlin, and E. Eisenberg. Connectivity and expression in protein networks: Proteins in a complex are uniformly expressed. *Phys. Rev. E* **73**, 031909 (2006).
2. **S. Carmi**, R. Cohen, and D. Dolev. Searching complex networks efficiently with minimal information. *Europhys. Lett.* **74**, 1102 (2006).
3. **S. Carmi**, S. Havlin, S. Kirkpatrick, Y. Shavitt, and E. Shir. A Model of Internet topology using *k*-shell decomposition, *P. Natl. Acad. Sci. USA* **104**, 11150 (2007).
4. E. Lopez, R. Parshani, R. Cohen, **S. Carmi**, and S. Havlin. Limited path percolation in complex networks. *Phys. Rev. Lett.* **99**, 188701 (2007).
5. M. Maragakis, **S. Carmi**, D. ben-Avraham, S. Havlin, and P. Argyrakis. Priority diffusion model in lattices and complex networks. *Phys. Rev. E (Rapid Communication)* **77**, 020103 (2008).
6. **S. Carmi**, Z. Wu, S. Havlin, and H. E. Stanley. Transport in networks with multiple sources and sinks. *EPL* **84**, 28005 (2008).
7. A. Kittas, **S. Carmi**, S. Havlin, and Panos Argyrakis. Trapping in complex networks. *EPL* **84**, 40008 (2008).
8. **S. Carmi**, P. L. Krapivsky, and D. ben-Avraham, Partition of networks into basins of attraction. *Phys. Rev. E* **78**, 066111 (2008).
9. **S. Carmi**, E. Y. Levanon, and E. Eisenberg. Efficiency of complex production in changing environment. *BMC Sys. Biol.* **3**:3 (2009).

10. **S. Carmi**, S. Havlin, C. Song, K. Wang, and H. A. Makse. Energy-landscape network approach to the glass transition. *J. Phys. A: Math. Theor.* **42**, 105101 (2009).
11. **S. Carmi**, S. Carter, J. Sun, and D. ben-Avraham. Asymptotic behavior of the Kleinberg model. *Phys. Rev. Lett.* **102**, 238702 (2009).
12. L. Turgeman, **S. Carmi**, and E. Barkai. Fractional Feynman-Kac equation for non-Brownian functionals. *Phys. Rev. Lett.* **103**, 190201 (2009).
13. H. Goldshmidt, D. Matas, A. Kabi, **S. Carmi**, R. Hope, and S. Michaeli. Persistent ER stress induces the Spliced Leader RNA Silencing pathway (SLS), leading to programmed cell death in *Trypanosoma brucei*. *PLoS Pathog.* **6**, e1000731 (2010).
14. R. Parshani, **S. Carmi**, and S. Havlin. Epidemic threshold for the Susceptible-Infectious-Susceptible model on random networks. *Phys. Rev. Lett.* **104**, 258701 (2010).
15. N. G. Kolev, J. B. Franklin, **S. Carmi**, H. Shi, S. Michaeli, and C. Tschudi. The transcriptome of the human pathogen *Trypanosoma brucei* at single-nucleotide resolution. *PLoS Pathog.* **6**, e1001090 (2010).
16. **S. Carmi**, L. Turgeman, and E. Barkai. On distributions of functionals of anomalous diffusion paths. *J. Stat. Phys.* **141**, 1071 (2010).
17. **S. Carmi**, I. Borukhov, and E. Y. Levanon. Identification of widespread ultra-edited human RNA. *PLoS Genet.* **7**, e1002317 (2011).
18. **S. Carmi**, G. M. Church, and E. Y. Levanon. Large scale DNA editing of retrotransposons accelerates mammalian genome evolution. *Nat. Commun.* **2**, 519 (2011).
19. **S. Carmi** and E. Barkai. Fractional Feynman-Kac equation for weak ergodicity breaking. *Phys. Rev. E* **84**, 061104 (2011).
20. S. K. Gupta, **S. Carmi**, H. Waldman Ben-Asher, I. D. Tkacz, I. Naboishchikov, and S. Michaeli. Basal splicing factors regulate the stability of mature mRNAs in Trypanosomes. *J. Biol. Chem.* **7**, 4991 (2013).
21. **S. Carmi**, P. F. Palamara, V. Vacic, T. Lencz, A. Darvasi, and I. Pe'er. The variance of identity-by-descent sharing in the Wright-Fisher model. *Genetics* **193**, 911 (2013).
22. S. K. Gupta, I. Kostı, G. Plaut, A. Pivko, I. D. Tkacz, S. Cohen-Chalamish, D. K. Biswas, C. Wachtel, H. Waldman Ben-Asher, **S. Carmi**, F. Glaser, Y. Mandel-Gutfreund, and S. Michaeli. The hnRNP F/H homologue of *Trypanosoma brucei* is differentially expressed in the two life cycle stages of the parasite and regulates splicing and mRNA stability. *Nucleic Acids Res.* **41**, 6577 (2013).
23. N. Bastas, M. Maragakis, P. Argyrakis, D. ben-Avraham, S. Havlin, and **S. Carmi**. Random walk with priorities in communication-like networks. *Phys. Rev. E* **88**, 022803 (2013).
24. S. K. Gupta, V. Chikne, D. Eliaz, I. D. Tkacz, I. Naboishchikov, **S. Carmi**, H. Waldman Ben-Asher, and S. Michaeli. Two splicing factors carrying serine-arginine motifs, TSR1 and TSR1IP, regulate splicing, mRNA stability and rRNA processing in *Trypanosoma brucei*. *RNA Biol.* **11**, 715 (2014).
25. H. T. Porath, **S. Carmi**, and E. Y. Levanon. A genome-wide map of hyper-edited RNA reveals numerous new sites. *Nat. Commun.* **5**, 4726 (2014).
26. **S. Carmi**, K. Y. Hui, E. Kochav, X. Liu, J. Xue, F. Grady, S. Guha, K. Upadhyay, D. Ben-Avraham, S. Mukherjee, B. M. Bowen, T. Thomas, J. Vijai, M. Cruts, G. Froyen, D. Lambrechts, S. Plaisance, C. Van Broeckhoven, P. Van Damme, H. Van Marck, N. Barzilai, A. Darvasi, K. Offit, S. Bressman, L. J. Ozelius, I. Peter, J. H. Cho, H. Ostrer, G. Atzmon, L. N. Clark, T. Lencz, and I. Pe'er. Sequencing an Ashkenazi reference panel supports population-targeted personal genomics and illuminates Jewish and European origins. *Nat. Commun.* **5**, 4835 (2014).
27. **S. Carmi**, P. R. Wilton, J. Wakeley, and I. Pe'er. A renewal theory approach to IBD sharing. *Theor. Popul. Biol.* **97**, 35 (2014).

28. J. Zidan\*, D. Ben-Avraham\*, **S. Carmi\***, T. Maray, E. Friedman, and G. Atzmon. Genotyping of geographically diverse Druze trios reveals substructure and a recent bottleneck. *Eur. J. Hum. Genet.* **23**, 1093 (2015).
29. P. R. Wilton, **S. Carmi\***, and A. Hobolth\*. The SMC' is a highly accurate approximation to the ancestral recombination graph. *Genetics* **200**, 343 (2015).
30. B. Baskovich, S. Hiraki, K. Upadhyay, P. Meyer, **S. Carmi**, N. Barzilai, A. Darvasi, K. Offit, S. Bressman, L. Ozelius, I. Peter, J. Cho, G. Atzmon, L. Clark, T. Lencz, I. Pe'er, H. Ostrer, and C. Oddoux. Expanded genetic screening panel for the Ashkenazi Jewish Population. *Genet. Med.* **18**, 522 (2016).
31. A. Quint, M. Sagi, **S. Carmi**, H. Daum, Z. Ben Neriah, V. Meiner, O. Elpeleg, and I. Lerer. An Ashkenazi founder mutation in the PKHD1 gene. *Eur. J. Med. Genet.* **59**, 86 (2016).
32. S. Yang, **S. Carmi**, and I. Pe'er. Rapidly registering identity-by-descent across ancestral recombination graphs. *J. Comput. Biol.* **23**, 495 (2016).
33. R. Jaron, N. Rosenfeld, F. Zahdeh, **S. Carmi**, L. Beni-Adani, R. Segel, S. Zeligson, L. Carmel, P. Renbaum, and E. Levy-Lahad. Expanding the phenotype of CRB2 mutations – A new ciliopathy syndrome? *Clin. Genet.* **90**, 540 (2016).
34. R. Hope, K. Egarmina, K. Voloshin, H. Waldman Ben-Asher, **S. Carmi**, D. Eliaz, Y. Drori, and S. Michaeli. Transcriptome and proteome analyses and the role of atypical calpain protein and autophagy in the spliced leader silencing (SLS) pathway in *Trypanosoma brucei*. *Mol. Microbiol.* **102**, 1 (2016).
35. J. Vijai, S. Topka, D. Villano, V. Ravichandran, K. N. Maxwell, A. Maria, T. Thomas, P. Gaddam, A. Lincoln, S. Kazzaz, B. Wenz, **S. Carmi**, K. A. Schrader, S. N. Hart, S. M. Lipkin, S. L. Neuhausen, M. F. Walsh, L. Zhang, F. Lejbkowitz, H. Rennert, Z. K. Stadler, M. Robson, J. N. Weitzel, M. J. Daly, F. J. Couch, K. L. Nathanson, L. Norton, G. Rennert, K. Offit. A recurrent *ERCC3* truncating mutation confers moderate risk for breast cancer. *Cancer Discov.* **6**, 1267 (2016).
36. E. Gilbert, **S. Carmi**, S. Ennis, J. Wilson, and G. Cavalleri. Genomic insights into the population structure and history of the Irish Travellers. *Sci. Rep.* **7**, 42187 (2017).
37. J. Xue, T. Lencz, A. Darvasi, I. Pe'er, and **S. Carmi**. The Time and Place of European Admixture in Ashkenazi Jewish History. *PLoS Genet.* **13**, e1006644 (2017).
38. Y. Einhorn, D. Weissglas-Volkov, **S. Carmi**, H. Ostrer, E. Friedman, and N. Shomron. Differential analysis of mutations in the Jewish Population and its implications on diseases. *Genet. Res., Camb.* **99**, e3 (2017).
39. D. M. Behar, L. Saag, M. Karmin, M. G. Gover, J. D. Wexler, L. F. Sanchez, E. Greenspan, A. Kushniarevich, O. Davydenko, H. Sahakyan, L. Yepiskoposyan, A. Boattini, S. Sarno, L. Pagani, **S. Carmi**, S. Tzur, E. Metspalu, C. Bormans, K. Skorecki, M. Metspalu, S. Rootsi, and R. Villems. The genetic variation in the R1a clade among the Ashkenazi Levites' Y chromosome. *Sci. Rep.* **7**, 14969 (2017).
40. K. Y. Hui, H. Fernandez-Hernandez, J. Hu, A. Schaffner, N. Pankratz, N.-Y. Hsu, L.-S. Chuang, **S. Carmi**, N. Villaverde, X. Li, M. Rivas, A. P. Levine, X. Bao, P. R. Labrias, T. Haritunians, D. Ruane, K. Gettler, E. Chen, D. Li, E. R. Schiff, N. Pontikos, N. Barzilai, S. R. Brant, S. Bressman, A. S. Cheifetz, L. N. Clark, M. J. Daly, R. J. Desnick, R. H. Duerr, S. Katz, T. Lencz, R. H. Myers, H. Ostrer, L. Ozelius, H. Payami, Y. Peter, J. D. Rioux, A. W. Segal, W. K. Scott, M. S. Silverberg, J. M. Vance, I. Ubarretxena-Belandia, T. Foroud, G. Atzmon, I. Pe'er, Y. Ioannou, D. P. B. McGovern, Z. Yue, E. E. Schadt, J. H. Cho, and I. Peter. Functional variants in the *LRRK2* gene confer shared effects on risk for Crohn's disease and Parkinson's disease. *Sci. Transl. Med.* **10**, eaai7795 (2018).
41. D. Rothschild, O. Weissbrod, E. Barkan, A. Kurilshikov, T. Korem, D. Zeevi, P. I. Costea, S. Shilo, D. Lador, A. Godneva, I. N. Kalka, N. Bar, A. V. Vila, N. Zmora, M. Pevsner-Fischer, D. Israeli, N. Kosower, G. Malka, B. C. Wolf, T. Avnit-Sagi, M. Lotan-Pompan, A.

- Weinberger, Z. Halpern, **S. Carmi**, J. Fu, Lifelines cohort study, C. Wijmenga, A. Zhernakova, E. Elinav, and E. Segal. Environmental factors dominate over host genetics in shaping human gut microbiota composition. *Nature* **555**, 210 (2018).
42. T. Lencz, J. Yu, C. Palmer, **S. Carmi**, D. Ben-Avraham, N. Barzilai, S. Bressman, A. Darvasi, J. H. Cho, L. N. Clark, Z. H. Gümüş, V. Joseph, R. Klein, S. Lipkin, K. Offit, H. Ostrer, L. J. Ozelius, I. Peter, G. Atzmon, and I. Pe'er. High-depth whole genome sequencing of an Ashkenazi Jewish reference panel: enhancing sensitivity, accuracy, and imputation. *Hum. Genet.* **137**, 343 (2018).
  43. L. King, J. Wakeley, and **S. Carmi**. A non-zero variance of Tajima's estimator for two sequences even for infinitely many unlinked loci. *Theor. Popul. Biol.* **122**, 22 (2018).
  44. Y. Yehuda, B. Blumenfeld, N. Mayorek, K. Makedonski, O. Vardi, L. Cohen-Daniel, Y. Mansour, S. Baror-Sebban, H. Masika, M. Farago, M. Berger, **S. Carmi**, Y. Buganim, A. Koren, and I. Simon. Germline DNA replication timing shapes mammalian genome composition. *Nucleic Acids Res.* **46**, 8299 (2018).
  45. G. Rosner, N. Gluck, **S. Carmi**, D. Bercovich, N. Fliss-Issakov, M. Ben-Yehoyada, S. Aharon-Caspi, E. Kellerman, H. Strul, O. Shibolet, and R. Kariv. POLD1 and POLE Gene Mutations in Jewish Cohorts of Early-Onset Colorectal Cancer and of Multiple Colorectal Adenomas. *Dis. Colon Rectum* **61**, 1073 (2018).
  46. Y. Erlich, T. Shor, I. Pe'er, and **S. Carmi**. Identity inference of genomic data using long-range familial searches. *Science* **362**, 690 (2018).
  47. E. Granot-Hershkovitz, D. Karasik, Y. Friedlander, L. Rodriguez-Murillo, R. Dorajoo, J. Liu, A. Sewda, I. Peter, **S. Carmi\***, and H. Hochner\*. A study of Kibbutzim in Israel reveals risk factors for cardiometabolic traits and subtle population structure. *Eur. J. Hum. Genet.* **26**, 1848 (2018).
  48. A. L. Severson, **S. Carmi**, and N. A. Rosenberg. The effect of consanguinity on between-individual identity-by-descent sharing. *Genetics* **212**, 305 (2019).
  49. D. A. Zeevi, F. Zahdeh, Y. Kling, P. Renbaum, E. Levy-Lahad, **S. Carmi**, and G. Altarescu. Off the street phasing (OTSP): No hassle haplotype phasing for molecular PGD applications. *J. Assist. Reprod. Gen.* **36**, 727 (2019).
  50. R. Schweiger, Y. Erlich, and **S. Carmi**. FactorialHMM: Fast and exact inference in factorial hidden Markov models. *Bioinformatics* **35**, 2162 (2019).
  51. D. Backenroth, F. Zahdeh, Y. Kling, A. Peretz, T. Rosen, D. Kort, S. Zeligson, T. Dror, S. Kirshberg, E. Burak, R. Segal, E. Levy-Lahad, D. Zangen, G. Altarescu\*, **S. Carmi\***, and D. A. Zeevi\*. Haploseek: A 24-hour all-in-one method for preimplantation genetic diagnosis (PGD) of monogenic disease and aneuploidy. *Genet. Med.* **21**, 1390 (2019).
  52. D. Backenroth and **S. Carmi**. A test for deviations from expected genotype frequencies on the X chromosome for sex-biased admixed populations. *Heredity* **123**, 470 (2019).
  53. M. Winther, S. Shpitzen, O. Yaacov, J. Landau, L. Oren, L. Foroozan-Rosenberg, N. Lev Cohain, D. Schurr, V. Meiner, A. Szalat, **S. Carmi**, M. R. Hayden, E. Leitersdorf, and R. Durst. In search for genetic explanation for LDLc variability in an FH family: Common SNPs and a rare mutation in microsomal triglyceride transfer protein explain only part of LDL variability in an FH family. *J. Lipid Res.* **60**, 1733 (2019).
  54. The ROHGen Consortium (including A. Helavy and **S. Carmi**). Associations of autozygosity with a broad range of human phenotypes. *Nat. Commun.* **10**, 4957 (2019).
  55. H. Fridman, D. M. Behar, **S. Carmi**, and E. Levy-Lahad. Preconception carrier screening yield: effect of variants of unknown significance in partners of carriers with clinically significant variants. *Genet. Med.*, in press (2019).

56. E. Karavani\*, O. Zuk\*, D. Zeevi, N. Barzilai, N. C. Stefanis, A. Hatzimanolis, N. Smyrnis, D. Avramopoulos, L. Kruglyak, G. Atzmon, M. Lam, T. Lencz\*, and **S. Carmi\***. Screening human embryos for polygenic traits has limited utility. *Cell*, in press (2019).

### Refereed conferences

1. S. Yang, **S. Carmi**, and I. Pe'er. Rapidly registering identity-by-descent across ancestral recombination graphs. *Lect. Notes Comput. Sc.* **9029**, 340 (2015). [RECOMB 2015]
2. A. Jaffe, R. Weiss, **S. Carmi**, Y. Kluger, and B. Nadler. Learning binary latent variable models: a tensor eigenpair approach. ICML 2018.

### Conference and workshop proceedings in refereed journals

1. E. Lopez, **S. Carmi**, S. Havlin, S. Buldyrev, and H. E. Stanley. Anomalous electrical and frictionless flow conductance in complex networks. *Physica D* **224**, 69 (2006). [Proceedings of the workshop “*Dynamics on Complex Networks and Applications*”, Dresden, 2006]
2. **S. Carmi**, Z. Wu, E. Lopez, S. Havlin, and H. E. Stanley. Transport between multiple users in complex networks. *Eur. Phys. J. B* **57**, 165 (2007). [Proceedings of the 5th International Conference “*Applications of Physics in Financial Analysis*” (APFA5), Torino, 2006]

### Book chapters

1. **S. Carmi** and E. Barkai. Fractional Feynman-Kac equation for anomalous diffusion functionals. Chapter 8 in *Fractional Dynamics: Recent Advances*, Eds. J. Klafter, S. C. Lim, and R. Metzler (World Scientific, Singapore, 2011).

### MEDIA COVERAGE

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- 2006 *Limited path percolation in complex networks*. Physical Review Focus.
- 2007 *A model of Internet topology*. MIT Technology Review, Science News, IEEE Spectrum, PopSci, Scientific American, Bloomberg News, Wired, and over 15 others.
- 2011 *Large scale DNA editing of retrotransposons*. Ha'aretz.
- 2014 *Sequencing an Ashkenazi reference panel*. LA Times, NBC News, Yahoo! News, The Scientist, National Geographic, Genome Web, Ha'aretz, The Jewish Daily Forward, and over 20 others, including radio and TV shows in Israel.
- 2014 *A renewal theory approach to IBD sharing*. Stanford Scope.
- 2014 *Genotyping of geographically diverse Druze trios*. Ha'aretz, Makor Rishon, NRG (Israel), Gene Expression blog.
- 2015 *Expanded Genetic Screening Panel*. The Jerusalem Post.
- 2016 40 most promising under 40. TheMarker Magazine, Israel
- 2017 *Genomic insights into the Irish Travellers*. Genome Web, Daily Mail, The Irish Times, The Irish Post, Irish Central, and others.
- 2018 *Functional variants in the LRRK2 gene*. Genome Web, Science Daily.
- 2018 *The human gut microbiota composition*. The Scientist, Science Daily, MedicalXpress, ynet, Ha'aretz, and others.
- 2018 *Identity inference of genomic data*. The New York Times, The Wall Street Journal, The Washington Post, The Atlantic, Time, Los Angeles Times, NBC News, ABC News, CNN, Bloomberg, Yahoo News, Wired, Science, Nature, Scientific American, The Scientist, GenomeWeb, IEEE Spectrum, Ha'aretz, The Marker (Israel), and others. Covered in Science Magazine's breakthroughs of 2018.
- 2018 *A 24-hour all-in-one method for preimplantation genetic diagnosis*. Doctors Only, Mako (Israel).
- 2019 *A study of Kibbutzim in Israel*. The NY Jewish Week.

- 2019 *Screening human embryos for polygenic traits*. Science, GenomeWeb, MIT Technology Review, Science News, Live Science, Inverse, Futurism, Jerusalem Post, and others
- 2018- Media appearances as an expert (in Israel). *Newspapers*: Ha'aretz, Yedioth Ahronoth, Israel Hayom, Times of Israel. *TV*: Channel 10, ynet (online). *Radio*: 103FM, Tel-Aviv radio.

## FUNDING

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- 2015-2023 Startup grant, The Faculty of Medicine (HUJI), \$200,000
- 2016- Private gift, The Barouh and Channah Berkovits Foundation, \$15,000
- 2017-2020 Abisch-Frenkel Foundation for the promotion of life sciences, "Evaluating the potential of personalized risk prediction for common diseases in the Ashkenazi Jewish population", \$55,000
- 2017 Israel Science Foundation (ISF) New Faculty Equipment Grant 932/17, "A computing and storage system for statistical and population genetics research", 202,215₪.
- 2017-2021 Israel Science Foundation (ISF) personal grant 407/17, "New methods for using genetic variation and genealogical records for inferring recent demographic events", 1,000,000₪.
- 2018-2022 Sub-award of the National Institutes of Health, "The X-factor of complex disease: Development, implementation, and extensive application of methods for analysis of the X chromosome in GWA, sequence-based association, and eQTL studies", PI: Alon Keinan, Cornell University. Amount to SC: TBD.
- 2018-2019 German-Israeli Foundation for Scientific Research and Development (GIF) I-2489-407.6/2017, "Reconstructing recent population histories using multi-individual haplotype sharing and applications to Ashkenazi Jewish genetics", €18,000.
- 2019-2022 United States – Israel Binational Science Foundation (BSF) 2017024, "Theory of consanguinity and its effects on genomic sharing within and between individuals". \$216,000. Joint PI with Noah Rosenberg, Stanford University.
- 2018 Hebrew University president award for the publication of high profile research, 10,000₪.
- 2019 Sub-award of the Israel Science Foundation (ISF) program in precision medicine, "Combined omics and electronic medical records-big data for prediction of phenotype sub-types, progression, and treatment outcome in age-related macular degeneration". Lead PI: Itay Chowers, Hadassah Medical Center. Amount to SC: TBD.

## CONFERENCES

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- 2004 Science of Complex Networks: from Biology to the Internet and WWW, Portugal (poster)
- 2005 Katzir Conference on Molecular Perspectives on Protein-Protein Interactions, Israel (poster)
- 2007 Complex Networks: from Biology to Information Technology, Italy (talk)
- 2011 Weak Chaos, Infinite Ergodic Theory, and Anomalous Dynamics, Germany (talk)
- 2012 The annual meeting of the American Society of Human Genetics, USA (poster)
- 2012 Personal Genomes and Medical Genomics, Cold Spring Harbor Laboratory, USA (talk)
- 2013 The annual meeting of the American Society of Human Genetics, USA (talk)
- 2014 Human Evolution in the Genomic Era: Origins, Populations, and Phenotypes, UK (talk)
- 2014 The Biology of Genomes, Cold Spring Harbor Laboratory, USA (poster)
- 2014 The annual meeting of the Society for Molecular Biology and Evolution, Puerto Rico (poster)
- 2014 The annual meeting of the American Society of Human Genetics, USA (poster)
- 2015 The Biology of Genomes, Cold Spring Harbor Laboratory, USA (poster)

- 2016 Understanding the Function of Human Genome Variation, Sweden (poster)
- 2016 Founder populations: Lessons from the Jewish Genome, Israel (talk)
- 2016 Probabilistic Modeling in Genomics, UK (talk)
- 2017 Revisiting the Question of Jewish Origins, USA (invited talk)
- 2017 Human Evolution: Fossils, Ancient and Modern Genomes, UK (talk)
- 2018 The Israel Statistical Association Annual Meeting, Israel (invited talk)
- 2018 Genomics – from disease prevention to treatment, Israel (invited talk)
- 2018 Genealogy and the Sciences, Israel (invited talk)
- 2019 The annual meeting of the American Society of Human Genetics, USA (talk)

## **WORKSHOPS**

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- 2006 Workshop on Networks and Complexity, Israel (talk)
- 2007 Adams fellowship annual poster session, the National Academy of Sciences, Israel
- 2008 Adams fellowship annual poster session, the National Academy of Sciences, Israel
- 2009 The Science of Complexity, Israel (talk)
- 2009 Bar-Ilan's Institute for Nanotechnology and Advanced Materials meeting, Israel (talk)
- 2010 The annual meeting of the Israel Physical Society, Israel (talk)
- 2014 Columbia University annual postdoc research symposium (poster; 4<sup>th</sup> prize)
- 2015 Columbia University Foundations of Data Science poster session
- 2015 Columbia University Data Science Institute Bi-Annual symposium poster session
- 2016 Israeli Bioinformatics Symposium (poster)
- 2018 HLA and KIR population dynamics, Israel (invited talk)
- 2018 Computational Genomics Summer Institute, UCLA, USA (invited talk)
- 2018 Workshop on "Jewish genes", The Hebrew University of Jerusalem, Israel (invited chair)
- 2019 Israel human population genetics meeting (keynote lecture)
- 2019 Innovations in Forensics and the Law, Israel (invited talk)
- 2019 Computational Genomics Summer Institute, UCLA, USA (invited talk)

## **SEMINAR TALKS**

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- 2007 The Center for Polymer Studies, Boston University
- 2008 The Center for Complex Network Research, Northeastern University
- 2009 The Center for Complex Network Research, Northeastern University
- 2010 IBM research center, Tel Aviv
- 2011 Department of Mathematics, Bar-Ilan University
- 2011 Department of Physics, Tel Aviv University
- 2012 Faculty of Life Sciences, Bar-Ilan University
- 2012 School of Computer Science, Tel-Aviv University
- 2013 School of Computer Science, Tel-Aviv University
- 2013 Broad Institute
- 2013 Department of Genetics, Harvard Medical School
- 2013 Department of Computer Science, Columbia University
- 2014 Division of Psychiatric Genomics, Icahn School of Medicine at Mount Sinai
- 2014 Department of Organismic and Evolutionary Biology, Harvard University
- 2014 Computational Biology and Bioinformatics, University of Southern California
- 2014 Department of Computer Science, University of California, Los Angeles
- 2014 Sackler Faculty of Medicine, Tel-Aviv University
- 2015 Department of Biological Statistics and Computational Biology, Cornell University
- 2015 Department of Genetics, The Faculty of Medicine, The Hebrew University of Jerusalem
- 2015 Department of Human Genetics and Metabolic Diseases, Hadassah Medical Center



2015 School of Computer Science and Engineering, The Hebrew University of Jerusalem  
 2015 Institute of Genetics, Shaare Zedek Medical Center  
 2017 Faculty of Mathematics and Computer Science, Weizmann Institute of Science  
 2018 Cancer Research Hub, The Faculty of Medicine, The Hebrew University of Jerusalem  
 2018 Faculty of Mathematics and Computer Science, Weizmann Institute of Science  
 2018 Department of Archaeogenetics, Max Planck Institute for the Science of Human History  
 2018 Institute of Genetics, Ha'Emek Medical Center  
 2018 School of Public Health, The Hebrew University of Jerusalem  
 2019 Department of Molecular Genetics, Weizmann Institute of Science  
 2019 Max Planck Institute for the Science of Evolutionary Biology  
 2019 Faculty of Mathematics and Computer Science, Weizmann Institute of Science  
 2019 The Faculty of Medicine, The Hebrew University of Jerusalem

### PEER REVIEW

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2008- An independent reviewer for:  
**Interdisciplinary:** Nature Communications, PLOS One, Scientific Reports  
**Life sciences:** eLife, American Journal of Human Genetics, Genome Biology, Genetics, Molecular Biology and Evolution, PLOS Genetics, PLOS Computational Biology, Theoretical Population Biology, RECOMB, BMC Bioinformatics, Human Genetics, Human Reproduction, Genes, Journal of Personalized Medicine, Molecular Biotechnology  
**Physics:** Physical Review Letters, Physical Review E, Europhysics Letters, New Journal of Physics, Journal of Statistical Mechanics, Physics Letters A, Journal of Physics A, Physica A, ESAIM: Mathematical Modelling and Numerical Analysis, Canadian Journal of Physics  
**Other:** Journal of Anthropological Research  
 2008 Selected as a Europhysics Letters distinguished referee.  
 2016- Grant reviewer for:  
 The Wellcome Trust, German-Israeli Foundation for Scientific Research and Development (GIF), Israel Science Foundation (ISF)

### OTHER PROFESSIONAL SERVICE

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#### University committees:

2016- Research committee, The Faculty of Medicine (HUJI)  
 2017- Advanced studies committee, The Faculty of Medicine (HUJI)  
 2016- PhD committee, Braun School of Public Health (HUJI)

#### Other university service:

2016-2017, 2018-2019 School seminar coordinator, Braun School of Public Health (HUJI)

#### PhD committees:

2016- Lily Agranat-Tamir, Life Sciences and Statistics (HUJI)

#### PhD thesis reading:

2017 Samantha Streicher, Yale School of Public Health

#### MSc/MPH thesis reading:

2017 Reshit Beizer, Public Health (HUJI)  
 2018 Nechama Averick, Public Health (HUJI)  
 2019 Itay Nitzan, Public Health (HUJI)  
 2019 Gabriel Azhari, Faculty of Engineering, Bar-Ilan University

#### Consulting (non-profit):

2019 Israel Science Foundation

2019 Israel Ministry of Justice  
 2017- Israel Ministry of Health national precision medicine initiative (PSIFAS)

**Consulting (industry):**

2016- My Heritage

**Memberships in societies:**

2010-2011 Israel Physical Society  
 2012-2014 The American Society of Human Genetics  
 2014 The Society for Molecular Biology and Evolution

**STUDENTS SUPERVISED**

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**Post-docs**

2016- Shay Tzur (part time)  
 2016 Avner Halevy (part time)  
 2019 Einat Granot-Hershkovitz

**PhD students**

2016- Shamam Waldman  
 2017- Hila Fridman (joint with Ephrat Levy-Lahad, Shaare Zedek Medical Center)

**Master's students**

2017-2019 Idit Lozowick-Gabay (joint with Ronen Durst and Dan Gilon, Hadassah Medical Center; rector's list)  
 2017-2019 Ehud Karavani  
 2016-2018 David Raveh-Brawer (faculty's excellent thesis award, graduated magna cum laude)

**Undergraduate research projects and research interns**

2018-2019 Roei Levy  
 Public health (HUJI)  
 Inference of sex-specific founder events with X-linked IBD sharing

2019- Saar Armon  
 Biomedical sciences (HUJI)  
 Algorithms for variant calling from low coverage sequencing data

2016- Kobi Landau  
 Medicine (HUJI), MD thesis  
 Computational methods for studying the male germline cell lineage

2018-2019 Itamar Medved  
 Biomedical sciences (HUJI)  
 Demographic inference based on multiway haplotype sharing

2016-2017 Or Yaacov  
 Medicine (HUJI)  
 Searching for genes associated with lipid levels in a large pedigree from Israel

2016 Dana Baril  
 Computer Science (HUJI)  
 Coalescence times in large-scale genealogical records

2015-2016 Noam Bar  
 Computer Science and Computational Biology (HUJI)  
 Imputation of missing variants for ultra-low coverage sequencing in Ashkenazi Jews

2015-2016 Gavriel Fialkoff  
 Computer Science and Computational Biology (HUJI)  
 Detection of copy number variants for preimplantation genetic screening with low-coverage single cell embryonic sequencing

- 2016 Adina Barchichat  
Bioinformatics, Jerusalem College of Bioinformatics  
IBD mapping on the X chromosome

**STUDENT PROJECTS SUPERVISED (BEFORE HUJI)**

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**Undergraduate computational biology research projects at Bar-Ilan University**

- 2010 Structural motifs in transcripts regulated under splicing factors depletion in *Trypanosoma brucei* (Or Garfunkel and Roy Azran)  
2011 DNA editing in polymorphic human retrotransposons (Anastasia Shapiro)  
2011 A comprehensive screen for DNA editing in mammalian genomes (Binyamin Knisbacher)

**Undergraduate/masters research projects at Columbia University**

- 2012 Quality control and variant statistics in the genomes of Ashkenazi Jews (Fillan Grady)  
2012 Imputation of Ashkenazi Jewish genomes using whole genome sequencing panels (Ethan Kochav)  
2012 Evaluating methods of local ancestry inference in closely related populations (James Xue)  
2013 Coverage of Ashkenazi Jewish genomes by long segments shared with a reference panel (James Xue)  
2014 Extracting IBD segments efficiently from genealogical trees (Shuo Yang)  
2015 Refining the parameters of the bottleneck in the Ashkenazi Jewish population history (Hannah Rosenwein)