

Armita Nourmohammad (she/her)

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Education	2008 - 2012	PhD in theoretical physics (summa cum laude) Thesis: <i>Evolution of regulatory complexes: a many-body system</i> Committee: Prof. Dr. Michael Lässig Prof. Dr. Joachim Krug Prof. Dr. Martin Kreitman
	2006 - 2008	Master's degree in physics at University of Cologne, Grade 1.1 (very good) Thesis: <i>From microsatellite slippage to cis-regulatory module evolution</i> Advisor: Prof. Dr. Michael Lässig
	2003 - 2006	Bachelor degree in physics at Sharif University of Technology, Tehran, Grade 16.44/20
Research activity & positions	2/2020 - present	Affiliate Investigator, Fred Hutchinson Cancer Research Center, Seattle
	10/2019 - present	Faculty of "UW Computational Molecular Biology", Seattle
	7/2019	Joliot Chair, ESPCI, Paris, France
	9/2018 - present	Assistant professor of physics, University of Washington, Seattle (on leave: 9/2018-9/2019)
	5/2018 - present	Faculty of the graduate school "Physics of biological and complex systems" (PBCS), University of Göttingen, Germany
	1/2018 - present	Faculty of the GAUSS graduate school "Theoretical and computational neuroscience" (PTCN), University of Göttingen
	9/2017 - present	Max Planck Research Group Leader, Max Planck Institute for Dynamics and Self-organization, Göttingen, Germany (non-tenure track position & 20% affiliation since 9/2019)
	9/2015 - 8/2017	Part-time lecturer in physics and the Lewis-Sigler Institute for Integrative Genomics, Princeton University, Princeton, USA
	10/2014 - 8/2017	Associate research scholar, Lewis-Sigler Institute for Integrative Genomics, Princeton University, Princeton, USA
	10/2012 - 10/2014	James S. McDonnell postdoctoral fellow, Lewis-Sigler Institute for Integrative Genomics, Princeton University, Princeton, USA
Honors & Awards	2014	Postdoctoral travel award for the annual meeting of SMBE 2014
	2012	Klaus-Liebrecht award for distinguished PhD dissertation (amount of 3,000 Euros)
	2011	James S. McDonnell Foundation (JSMF) postdoctoral fellowship for studying complex systems (amount of \$200,000)
	2011	Finalist in Harvard Society of Fellows
	2011	Student fellowship by Burroughs Wellcome Fund to attend the KITP program, <i>Microbial and viral evolution</i> , at UC Santa Barbara
	2007	Scholarship from Bonn-Cologne excellency school of physics and astronomy
	2003	Ranked 15 th nationwide in the Iranian university entrance exam among more than 300,000 participants

Research Grants	2020	Royalty Research Fund, <i>Learning the shape space of protein universe to predict function</i> , \$ 38,000
	2018	Single PI project within SFB1310 <i>Predictability in Evolution</i> , EUR 207,000
	2017	Max Planck Research Group, EUR 1,851,000

Publications *Pre-prints under review*

- **G. Isacchini**, A. Walczak[†], T. Mora[†], **A. Nourmohammad**[†] (2020) Deep generative selection models of T and B cell receptor repertoires with soNNia, arXiv:2011.03112
- **Z. Montague***, H. Lv*, J. Otwinowski, W. S. DeWitt, **G. Isacchini**, G. K. Yip, W. W. Ng, O. T. Tsang, M. Yuan, H. Liu, I. A. Wilson, M. Peiris, N. C. Wu, **A. Nourmohammad**[†], C. K. Mok[†] (2020) Dynamics of B-cell repertoires and emergence of cross-reactive responses in COVID-19 patients with different disease severity, arXiv:2007.06762
Press: The Physics of a Deadly Virus
- **O.H. Schnaack**, **A. Nourmohammad** (2020) Optimal evolutionary decision-making to store immune memory. arXiv:2007.01363

¹: equal contribution, [†]: corresponding author, **bold face**: group members

Peer reviewed

1. **A. Nourmohammad**[†] and C. Eksin (2021) Optimal evolutionary control for artificial selection on molecular phenotypes, Phys Rev. X (*in press*)
2. J. Otwinowski, **C. LaMont**, **A. Nourmohammad** (2020) Information-geometric optimization with natural selection, Entropy, 22(9), 967
3. **G. Isacchini**, C. Olivares, **A. Nourmohammad**, A. Walczak and T. Mora (2020) SOS: Online probability estimation and generation of T and B cell receptors, Bioinformatics, btaa574
4. **G. Isacchini**, Z. Sethna, Y. Elhanati, **A. Nourmohammad**, A. Walczak and T. Mora (2020) Generative models of T-cell receptor sequences, Phys. Rev. E 101, 062414
5. N.C. Wu, J. Otwinowski, A.J. Thompson, C.M. Nycholat, J.C. Paulson, **A. Nourmohammad** and I. A. Wilson (2020) Major hemagglutinin antigenic site B of human influenza H3N2 viruses has an evolving local fitness landscape, Nature Communications 11,1233
6. S. Bradde[†], **A. Nourmohammad**[†], S. Goyal[†], V. Balasubramanian[†] (2020) The size of the immune repertoire in bacteria, Proc. Nat. Acad. Sci. 117 (10) 5144-5151
Commentary by M. Deem: 10.1073/pnas.2002746117
7. **A. Nourmohammad**[†], J. Otwinowski, M. Łuksza, T. Mora and A. M. Walczak (2019), Fierce Selection and Interference in B-Cell Repertoire Response to Chronic HIV-1, Mol Biol Evol, 36(10): 2184–2194
8. T. Hagai, X. Chen, R.J. Miragaia, R. Rostom, T. Gomes, N. Kunowska, J. Henriksson, J. Park, V. Proserpio, G. Donati, L. Bossini-Castillo, F.A.V. Braga, G. Naamati, J. Fletcher, E. Stephenson, P. Vegh, G. Trynka, I. Kondova, M. Dennis, M. Haniffa, **A. Nourmohammad**, M. Lässig and S.A. Teichmann (2018), Gene expression variability across cells and species shapes innate immunity, Nature 563: 197–202
F1000Prime Recommended

9. **A. Nourmohammad**[†], J. Rambeau, T. Held, V. Kovacova, J. Berg and M. Lässig (2017) Adaptive evolution of gene expression in Drosophila, Cell Reports 20, 1385-95
F1000Prime Recommended
10. **A. Nourmohammad**^{1†}, J. Otwinowski¹ and J. Plotkin (2016) Host-pathogen coevolution and the emergence of broadly neutralizing antibodies in chronic infections, PLoS Genet. 12(7): e1006171
11. T. Held, **A. Nourmohammad** and M. Lässig (2014) Adaptive evolution of molecular phenotypes, J. Stat. Mech, P09029
12. **A. Nourmohammad**¹, T. Held¹, M. Lässig, (2013) Universality and predictability in molecular quantitative genetics, Curr Opin Genet Dev. 23(6):684-93
13. **A. Nourmohammad**¹, S. Schiffels¹ and M. Lässig (2013) Evolution of molecular phenotypes under stabilizing selection, J. Stat. Mech. P01012
14. **A. Nourmohammad** (2012) Evolution of regulatory complexes: a many-body system, PhD Thesis, University of Cologne
15. **A. Nourmohammad** and M. Lässig (2011) Formation of regulatory modules by local sequence duplication, PLoS Comput Biol 7(10): e1002167
16. R.H. Abdolvahhab, F. Roshani, **A. Nourmohammad**, M. Sahimi and MR. Tabar (2008) Analytical and numerical studies of sequence dependence of passage times for translocation of heterobiopolymers through nanopores, J. Chem. Phys. 129(23):235102

¹: equal contribution, [†]: corresponding author, **bold face**: group members

Invited Seminars & Conference talks

- | | |
|---------|---|
| 2/2021 | Computational Biology Section, Fred Hutchinson Cancer Research Center, Seattle, USA |
| 1/2021 | “Quantitative evolution, phylogeny and ecology: from models to data and back”, Institut Henri Poincaré, Paris (virtual meeting: online recording) |
| 12/2020 | AIRR Community Meeting V (virtual meeting: online recording) |
| 10/2020 | 4 th Annual Symposium on Physical Concepts and Computational Models in Immunology, Cologne, Germany |
| 8/2020 | “Physical Biology of the Cell Course”, Marine Biological Laboratory, Woods Hole (MA), USA (virtual seminar: online recording) |
| 1/2020 | Math Modeling Affinity Seminar Series, Fred Hutchinson Cancer Research Center, Seattle, USA |
| 12/2019 | Computational Biology Section, Fred Hutchinson Cancer Research Center, Seattle, USA |
| 12/2019 | “Adaptation symposium”, Graduate Center, CUNY, NYC |
| 8/2019 | “Out-of-equilibrium processes in evolution and ecology”, Oaxaca, Mexico |
| 7/2019 | “From Molecular Basis to Predictability and control of evolution” Nordita Institute, Stockholm, Sweden |
| 6/2019 | Gordon Research Conference “Molecular Mechanisms of Evolution”, Stonehill College, USA |
| 4/2019 | Campus seminar at Max Planck Institute for Biological Cybernetics, Tübingen, Germany |
| 3/2019 | APS March meeting, invited session “Statistical physics of large populations of Cells: from microbes to tissues”, Boston, USA |
| 10/2018 | “Stochasticity and control in the dynamics and diversity of immune repertoires”, Paris, France |
| 9/2018 | “Physical concepts and computational models in immunology”, ENS, Paris |
| 9/2018 | “Paths in Statistical Physics (in honor of Luca Peliti)”, Paris, France |
| 4/2018 | “APS Physics-Next Meeting: Physics of Living Matter”, Long Island, USA |

4/2018 “Exploring the unreasonable ineffectiveness of mathematics in biology”,
 University of Pennsylvania, Philadelphia, USA (panel speaker)
 2/2018 Evolution of Diversity, Les Houches, France
 1/2018 Regulation and inference in biological networks, Bardonecchia, Italy
 1/2018 SFB 937 seminar, Göttingen, Germany
 10/2017 EMBO conference “Quantitative Principles in Biology”,
 Heidelberg, Germany
 3/2017 Seminar at the physics department, Yale University, New Haven, USA
 3/2017 Biophysics seminar, Department of physics, UCLA, USA
 2/2017 Physics department seminar, Caltech, Pasadena, USA
 2/2017 Physics Colloquium, University of Washington, Seattle, USA
 1/2017 Seminar at the Max Planck Society, Berlin, Germany
 1/2017 Physics department seminar, ICTP-SAIFR, São Paulo, Brazil
 11/2016 Center for Studies in Physics and Biology seminar series,
 Rockefeller University, New York, USA
 10/2016 Biophysics seminar, ENS & ESPCI, Paris, France
 10/2016 CIRB seminar, Collège de France, Paris, France
 10/2016 Séminaire de LPTMS, Orsay, France
 9/2016 SFB 680 Evolution Colloquium, Cologne, Germany
 9/2016 20th evolutionary biology meeting, Marseilles, France
 6/2016 Seminar at AMOLF, Amsterdam, the Netherlands
 4/2016 Fred Hutchinson Cancer Research Center, Seattle, USA
 3/2016 Quantitative Life Sciences seminar, ICTP, Trieste, Italy
 3/2016 APS March meeting, “Quantitative Immunology” session, Baltimore, USA
 2/2016 Quantitative immunology, KITP Program, UCSB, USA
Online Talk: effective theory of immune-pathogen interactions
 7/2015 Forecasting evolution?, Gulbenkian Institute, Portugal
 10/2014 Seminar at MPI for Cell Biology and Genetics, Dresden, Germany
 9/2014 Quantitative Systems Biology seminar, IST, Austria
 7/2014 Biophysics seminar, Rutgers University, USA
 3/2011 Natural History Seminars, University of Chicago, USA
 2/2011 Microbial and Viral Evolution, KITP Program, UCSB, USA
Online Talk: evolutionary modes of regulatory sequences in eukaryotes

**Contributed talks
& group seminars**

7/2018 Annual meeting of the Society for Molecular Biology and Evolution (SMBE),
 Yokohama, Japan
 3/2016 Antibody viral co-evolution workshop, Los Alamos, USA
 1/2015 EvoGroup seminar, Princeton University, USA
 10/2014 Seminar at Isabel Gordo’s group, Instituto Gulbenkian Ciencia, Portugal
 8/2014 Seminar at Daniel Fisher’s group, Stanford, USA
 6/2014 Annual meeting of the Society for Molecular Biology and Evolution (SMBE),
 Puerto Rico
 12/2013 110th Statistical Mechanics Conference, Rutgers University, USA
 4/2013 Seminar at Joshua Plotkin’s group, University of Pennsylvania, USA
 3/2013 American Physical Society (APS) meeting, Baltimore, USA
 2/2013 Seminar at Daniel Fisher’s group, Stanford University, USA
 10/2011 RECOMB Conference on Regulatory Genomics & Systems Biology, Barcelona
Online Talk: formation and conservation of regulatory binding site complexes
 10/2011 Workshop at the Bonn-Cologne graduate school of physics and astronomy,
Application of information theory in genomics, Bonn, Germany.
 7/2011 Moscow Conference on Computational Molecular Biology, Moscow, Russia
 4/2011 Seminar at Edo Kussell’s group, NYU, New York, USA

	11/2010	SFB 680 Seminar, Institute for Genetics, University of Cologne, Germany
	8/2009	15 th Annual European Meeting of PhD Students in Evolutionary Biology, Schoorl, the Netherlands
Lectures in summer schools & workshops	8/2020	Physical Biology of the Cell at Marine Biological Laboratory, Woods Hole, Massachusetts
	7/2019	Boulder summer school on theoretical biophysics, Boulder, CO, USA
	11/2018	International Curie course on biophysics, “Multiscale Integration in Biological Systems”, Institut Curie, Paris
	1/2017	VI Southern-Summer School on Mathematical-Biology ICTP-SAIFR, São Paulo, Brazil (lecturer)

Undergrad & Graduate Teaching	Fall 2020	Classical mechanics (PHYS 121), University of Washington, Seattle
	Spring 2020	
	Winter 2020	Biophysics (PHYS 429), University of Washington, Seattle
	Spring 2018	Statistical Physics in Biology, George-August University of Göttingen, Göttingen, Germany
	Spring 2018	Special topics (biophysics) graduate course (phys 578), University of Washington, Seattle, USA
	2015-2017	<i>Integrated science curriculum</i> , Lewis-Sigler Institute for Integrative Genomics, Princeton University, Princeton, USA

Service to Biophysics & Comp. Bio. communities

☐ **Journal Referee and Editorial Activities**

Reviewing editor: *eLife*

ad-hoc reviewer: *BMC Evolutionary Biology, eLife, Genetics, Interface Focus, Journal of Statistical Mechanics, Journal of Statistical Physics, Molecular Biology and Evolution, PNAS, Physical Review Letters, Physical Review E, PLoS computational biology, Virus Evolution*

Grant reviewer: *QLife- Institut Curie*

recommandeuse: “Peer Community in”

☐ **Meeting organizer**

2/2022	BIRS workshop: “Sensing and Signaling in Immune Systems”, Banff Center, Canada
8/2019	“Out-of-equilibrium processes in evolution and ecology”, Casa Matematica Oaxaca (CMO), Oaxaca, Mexico
7/2019	“From Molecular Basis to Predictability and control of evolution”, Nordita program, Stockholm, Sweden
7/2018	“Predictability of rapid evolutionary processes”, SMBE session, Yokohoma, Japan
3/2018	“Immune-pathogen interactions”, CRC 1310 theory symposium, Cologne, Germany
4/2016	“Molecular coevolution: lessons from pathogen-immune system interactions” PCTS workshop, Princeton University, Princeton, USA
7/2015	“Beyond the equilibrium paradigm: the role of temporal processes in population genetics and evolution”, SMBE session, Vienna, Austria
2011	‘Statistical physics of biological systems”, BCGS session, Workshop of Bonn-Cologne Graduate School (BCGS), Bonn, Germany

□ **Other experiences and professional memberships:**

2019 - 2023 Scientific Committee Member, International Research Network (IRN),
 “Predictability, Adaptation, Navigation”
2018 - Consultant on Burroughs Wellcome grant “Quantitative and Statistical
 Thinking in the Life Sciences”
2012 - 2016 Member, American Physical Society
2010 Member, Society for Molecular Biology and Evolution

**Thesis
students**

Graduate advisees Michael Pun (UW, Physics),
 Zachary Montague (UW, Physics),
 Oskar Schnaack (MPIDS, Physics),
 Vincent Balardi (MPIDS; co-advised with Y. Rondelez)
 Giulio Isacchini (MPIDS; co-advised with A. Walczak & T. Mora)

Thesis Committee Jared Callaham (UW, Mechanical Engineering)
member William deWitt (UW, Genome Sciences)
 Seth Hirsh (UW, Physics)
 Zeeshawn Kazi (UW, Physics)
 Marco Molari (École Normale Supérieure, Paris)
 Sid Rath (UW, Materials Science and Engineering)
 Ying-Jen Yang (UW, Applied Mathematics)
 Hengji Wang (UW, Physics)

February 2021