Andrea Parmeggiani

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8629288/publications.pdf

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41 papers

1,869 citations

430874 18 h-index 302126 39 g-index

46 all docs

46 docs citations

times ranked

46

1079 citing authors

#	Article	IF	CITATIONS
1	Relaxation time asymmetry in stator dynamics of the bacterial flagellar motor. Science Advances, 2022, 8, eabl8112.	10.3	2
2	Modelling the effect of ribosome mobility on the rate of protein synthesis. European Physical Journal E, 2021, 44, 19.	1.6	3
3	Supercoiled DNA and non-equilibrium formation of protein complexes: A quantitative model of the nucleoprotein ParBS partition complex. PLoS Computational Biology, 2021, 17, e1008869.	3.2	4
4	Modeling and live imaging of mechanical instabilities in the zebrafish aorta during hematopoiesis. Scientific Reports, 2021, 11, 9316.	3.3	3
5	Macrophage morphological plasticity and migration is Rac signalling and MMP9 dependant. Scientific Reports, 2021, 11, 10123.	3.3	10
6	Mechanical instabilities of aorta drive blood stem cell production: a live study. Cellular and Molecular Life Sciences, 2020, 77, 3453-3464.	5.4	9
7	ATP-Driven Separation of Liquid Phase Condensates in Bacteria. Molecular Cell, 2020, 79, 293-303.e4.	9.7	107
8	Physical Modeling of a Sliding Clamp Mechanism for the Spreading of ParB at Short Genomic Distance from Bacterial Centromere Sites. IScience, 2020, 23, 101861.	4.1	22
9	Stochastic modelling of collective motor protein transport through a crossing of microtubules. Journal of Theoretical Biology, 2020, 505, 110370.	1.7	6
10	Phase separation of polymer-bound particles induced by loop-mediated one dimensional effective long-range interactions. Physical Review Research, 2020, 2, .	3.6	2
11	Low-frequency phonon dynamics and related thermal properties of axially stressed single-walled carbon nanotubes. Journal of Physics Condensed Matter, 2019, 31, 425302.	1.8	2
12	Looping and clustering model for the organization of protein-DNA complexes on the bacterial genome. New Journal of Physics, 2018, 20, 035002.	2.9	9
13	A conserved mechanism drives partition complex assembly on bacterial chromosomes and plasmids. Molecular Systems Biology, 2018, 14, e8516.	7.2	53
14	Surfing on Protein Waves: Proteophoresis as a Mechanism for Bacterial Genome Partitioning. Physical Review Letters, 2017, 119, 028101.	7.8	34
15	Foci of cyclin A2 interact with actin and RhoA in mitosis. Scientific Reports, 2016, 6, 27215.	3.3	5
16	Stochastic Self-Assembly of ParB Proteins Builds the Bacterial DNA Segregation Apparatus. Cell Systems, 2015, 1, 163-173.	6.2	118
17	Variable Combinations of Specific Ephrin Ligand/Eph Receptor Pairs Control Embryonic Tissue Separation. PLoS Biology, 2014, 12, e1001955.	5.6	67
18	Motor protein traffic regulation by supply–demand balance of resources. Physical Biology, 2014, 11, 056006.	1.8	26

#	Article	IF	Citations
19	Stepping and Crowding of Molecular Motors: Statistical Kinetics from an Exclusion Process Perspective. Biophysical Journal, 2014, 107, 1176-1184.	0.5	18
20	Modelling Collective Cytoskeletal Transport andÂIntracellular Traffic. Mathematics for Industry, 2014, , 1-25.	0.4	0
21	Role of network junctions for the totally asymmetric simple exclusion process. Physical Review E, 2013, 88, 042104.	2.1	18
22	Modeling Cytoskeletal Traffic: An Interplay between Passive Diffusion and Active Transport. Physical Review Letters, 2013, 110, 098102.	7.8	63
23	Transport on a lattice with dynamical defects. Physical Review E, 2013, 87, 012705.	2.1	25
24	Exclusion processes on networks as models for cytoskeletal transport. New Journal of Physics, 2013, 15, 085005.	2.9	58
25	Centrosomal targeting of Syk kinase is controlled by its catalytic activity and depends on microtubules and the dynein motor. FASEB Journal, 2013, 27, 109-122.	0.5	7
26	Taming Nonequilibrium Statistics. Physics Magazine, 2012, 5, .	0.1	5
27	Totally Asymmetric Simple Exclusion Process on Networks. Physical Review Letters, 2011, 107, 068702.	7.8	84
28	Long-Range Protein Coupling Mediated by Critical Low-Energy Modes of Tubular Lipid Membranes. Physical Review Letters, 2010, 105, 028102.	7.8	12
29	Understanding totally asymmetric simple-exclusion-process transport on networks: Generic analysis via effective rates and explicit vertices. Physical Review E, 2009, 80, 041128.	2.1	53
30	Non-Equilibrium Collective Transport on Molecular Highways. , 2009, , 667-677.		3
31	HEX-TASEP: dynamics of pinned domains for TASEP transport on a periodic lattice of hexagonal topology. Journal of Physics Condensed Matter, 2008, 20, 295213.	1.8	12
32	Form-finding of complex tensegrity structures: application to cell cytoskeleton modelling. Comptes Rendus - Mecanique, 2006, 334, 662-668.	2.1	25
33	Dynamic correlation functions and Boltzmann-Langevin approach for driven one-dimensional lattice gas. Physical Review E, 2005, 72, 036123.	2.1	38
34	Renewal processes and fluctuation analysis of molecular motor stepping. Physical Biology, 2005, 2, 207-222.	1.8	9
35	Totally asymmetric simple exclusion process with Langmuir kinetics. Physical Review E, 2004, 70, 046101.	2.1	218
36	Internal Motility in Stiffening Actin-Myosin Networks. Physical Review Letters, 2004, 93, 268101.	7.8	37

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37	Micromechanics of Molecular Motors: Experiments and Theory. , 2004, , 151-176.		2
38	Collective phenomena in intracellular processes. Genome Informatics, 2004, 15, 46-55.	0.4	17
39	Phase Coexistence in Driven One-Dimensional Transport. Physical Review Letters, 2003, 90, 086601.	7.8	408
40	Detachment of molecular motors under tangential loading. Europhysics Letters, 2001, 56, 603-609.	2.0	34
41	Energy transduction of isothermal ratchets: Generic aspects and specific examples close to and far from equilibrium. Physical Review E, 1999, 60, 2127-2140.	2.1	235