

Daniel M Weinreich

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

4,876
citations

331538

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414303

32
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47
all docs

47
docs citations

47
times ranked

3698
citing authors

#	ARTICLE	IF	CITATIONS
1	Density fluctuations, homeostasis, and reproduction effects in bacteria. <i>Communications Biology</i> , 2022, 5, 397.	2.0	2
2	Covalent docking and molecular dynamics simulations reveal the specificity-shifting mutations Ala237Arg and Ala237Lys in TEM beta-lactamase. <i>PLoS Computational Biology</i> , 2022, 18, e1009944.	1.5	0
3	Herding an evolving biological population with quantum control tools. <i>Nature Physics</i> , 2021, 17, 17-19.	6.5	5
4	Predicting the viability of beta-lactamase: How folding and binding free energies correlate with beta-lactamase fitness. <i>PLoS ONE</i> , 2020, 15, e0233509.	1.1	26
5	Title is missing!. , 2020, 15, e0233509.		0
6	Title is missing!. , 2020, 15, e0233509.		0
7	Title is missing!. , 2020, 15, e0233509.		0
8	Title is missing!. , 2020, 15, e0233509.		0
9	Migration promotes mutator alleles in subdivided populations. <i>Evolution; International Journal of Organic Evolution</i> , 2019, 73, 600-608.	1.1	4
10	Selection on mutators is not frequency-dependent. <i>ELife</i> , 2019, 8, .	2.8	6
11	The Influence of Higher-Order Epistasis on Biological Fitness Landscape Topography. <i>Journal of Statistical Physics</i> , 2018, 172, 208-225.	0.5	64
12	Patterns of musculoskeletal growth and dimensional changes associated with selection and developmental plasticity in domestic and wild strain turkeys. <i>Ecology and Evolution</i> , 2018, 8, 3229-3239.	0.8	9
13	Sign of selection on mutation rate modifiers depends on population size. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 3422-3427.	3.3	36
14	Evolutionary constraints in fitness landscapes. <i>Heredity</i> , 2018, 121, 466-481.	1.2	26
15	Enzyme efficiency but not thermostability drives cefotaxime resistance evolution in TEM-1 β -lactamase. <i>Molecular Biology and Evolution</i> , 2017, 34, msx053.	3.5	48
16	Variability in Fitness Effects Can Preclude Selection of the Fittest. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2017, 48, 399-417.	3.8	18
17	Measuring epistasis in fitness landscapes: The correlation of fitness effects of mutations. <i>Journal of Theoretical Biology</i> , 2016, 396, 132-143.	0.8	55
18	Genetically Determined Variation in Lysis Time Variance in the Bacteriophage ϕ X174. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 939-955.	0.8	8

#	ARTICLE	IF	CITATIONS
19	Adaptive Landscape by Environment Interactions Dictate Evolutionary Dynamics in Models of Drug Resistance. <i>PLoS Computational Biology</i> , 2016, 12, e1004710.	1.5	71
20	Quantitative Description of a Protein Fitness Landscape Based on Molecular Features. <i>Molecular Biology and Evolution</i> , 2015, 32, 1774-1787.	3.5	57
21	Should evolutionary geneticists worry about higher-order epistasis?. <i>Current Opinion in Genetics and Development</i> , 2013, 23, 700-707.	1.5	236
22	FISHER'S GEOMETRIC MODEL OF ADAPTATION MEETS THE FUNCTIONAL SYNTHESIS: DATA ON PAIRWISE EPISTASIS FOR FITNESS YIELDS INSIGHTS INTO THE SHAPE AND SIZE OF PHENOTYPE SPACE. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2957-2972.	1.1	32
23	GENOME STRUCTURE AND THE BENEFIT OF SEX. <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 523-536.	1.1	38
24	High-throughput identification of genetic interactions in HIV-1. <i>Nature Genetics</i> , 2011, 43, 398-400.	9.4	7
25	Stepwise acquisition of pyrimethamine resistance in the malaria parasite. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 12025-12030.	3.3	241
26	Temporal Constraints on the Incorporation of Regulatory Mutants in Evolutionary Pathways. <i>Molecular Biology and Evolution</i> , 2009, 26, 2455-2462.	3.5	12
27	Empirical fitness landscapes reveal accessible evolutionary paths. <i>Nature</i> , 2007, 445, 383-386.	13.7	510
28	Darwinian Evolution Can Follow Only Very Few Mutational Paths to Fitter Proteins. <i>Science</i> , 2006, 312, 111-114.	6.0	1,266
29	Missense meanderings in sequence space: a biophysical view of protein evolution. <i>Nature Reviews Genetics</i> , 2005, 6, 678-687.	7.7	586
30	PERSPECTIVE: SIGN EPISTASIS AND GENETIC COCONSTRAINT ON EVOLUTIONARY TRAJECTORIES. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1165-1174.	1.1	384
31	RAPID EVOLUTIONARY ESCAPE BY LARGE POPULATIONS FROM LOCAL FITNESS PEAKS IS LIKELY IN NATURE. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1175-1182.	1.1	201
32	The Rank Ordering of Genotypic Fitness Values Predicts Genetic Constraint on Natural Selection on Landscapes Lacking Sign Epistasis. <i>Genetics</i> , 2005, 171, 1397-1405.	1.2	34
33	RAPID EVOLUTIONARY ESCAPE BY LARGE POPULATIONS FROM LOCAL FITNESS PEAKS IS LIKELY IN NATURE. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1175.	1.1	58
34	PERSPECTIVE:SIGN EPISTASIS AND GENETIC CONSTRAINT ON EVOLUTIONARY TRAJECTORIES. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1165.	1.1	227
35	Perspective: Sign epistasis and genetic constraint on evolutionary trajectories. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1165-74.	1.1	401
36	Rapid evolutionary escape by large populations from local fitness peaks is likely in nature. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1175-82.	1.1	91

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37	Contrasting Patterns of Nonneutral Evolution in Proteins Encoded in Nuclear and Mitochondrial Genomes. <i>Genetics</i> , 2000, 156, 385-399.	1.2	111