

Susanna C Manrubia

List of Publications by Year in descending order

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122
papers

4,636
citations

94433

37
h-index

123424

61
g-index

132
all docs

132
docs citations

132
times ranked

3542
citing authors

#	ARTICLE	IF	CITATIONS
1	Suppression of viral infectivity through lethal defection. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 4448-4452.	7.1	170
2	Criticality and scaling in evolutionary ecology. Trends in Ecology and Evolution, 1999, 14, 156-160.	8.7	167
3	Self-similarity of extinction statistics in the fossil record. Nature, 1997, 388, 764-767.	27.8	156
4	Evolutionary Transition toward Defective RNAs That Are Infectious by Complementation. Journal of Virology, 2004, 78, 11678-11685.	3.4	150
5	Role of Intermittency in Urban Development: A Model of Large-Scale City Formation. Physical Review Letters, 1997, 79, 523-526.	7.8	148
6	Are rainforests self-organized in a critical state?. Journal of Theoretical Biology, 1995, 173, 31-40.	1.7	146
7	Eukaryotic Community Distribution and Its Relationship to Water Physicochemical Parameters in an Extreme Acidic Environment, Rìo Tinto (Southwestern Spain). Applied and Environmental Microbiology, 2006, 72, 5325-5330.	3.1	126
8	Extinction and self-organized criticality in a model of large-scale evolution. Physical Review E, 1996, 54, R42-R45.	2.1	118
9	The turning point and end of an expanding epidemic cannot be precisely forecast. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 26190-26196.	7.1	117
10	Resistance of virus to extinction on bottleneck passages: Study of a decaying and fluctuating pattern of fitness loss. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 10830-10835.	7.1	109
11	Stochastic multiplicative processes with reset events. Physical Review E, 1999, 59, 4945-4948.	2.1	106
12	Mutual synchronization and clustering in randomly coupled chaotic dynamical networks. Physical Review E, 1999, 60, 1579-1589.	2.1	102
13	Viral Genome Segmentation Can Result from a Trade-Off between Genetic Content and Particle Stability. PLoS Genetics, 2011, 7, e1001344.	3.5	95
14	Quasispecies dynamics and RNA virus extinction. Virus Research, 2005, 107, 129-139.	2.2	93
15	Phase transitions and complex systems: Simple, nonlinear models capture complex systems at the edge of chaos. Complexity, 1996, 1, 13-26.	1.6	92
16	Vertical transmission of culture and the distribution of family names. Physica A: Statistical Mechanics and Its Applications, 2001, 295, 1-8.	2.6	90
17	The dawn of the RNA World: Toward functional complexity through ligation of random RNA oligomers. Rna, 2009, 15, 743-749.	3.5	89
18	High mutation rates, bottlenecks, and robustness of RNA viral quasispecies. Gene, 2005, 347, 273-282.	2.2	84

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19	Biodiversity in model ecosystems, I: coexistence conditions for competing species. <i>Journal of Theoretical Biology</i> , 2005, 235, 521-530.	1.7	77
20	Topological Structure of the Space of Phenotypes: The Case of RNA Neutral Networks. <i>PLoS ONE</i> , 2011, 6, e26324.	2.5	72
21	Potential Benefits of Sequential Inhibitor-Mutagen Treatments of RNA Virus Infections. <i>PLoS Pathogens</i> , 2009, 5, e1000658.	4.7	68
22	Population Bottlenecks in Quasispecies Dynamics. , 2006, 299, 141-170.		67
23	Multipartite viruses: adaptive trick or evolutionary treat?. <i>Npj Systems Biology and Applications</i> , 2017, 3, 34.	3.0	62
24	Modeling Viral Genome Fitness Evolution Associated with Serial Bottleneck Events: Evidence of Stationary States of Fitness. <i>Journal of Virology</i> , 2002, 76, 8675-8681.	3.4	58
25	Pathways to extinction: beyond the error threshold. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2010, 365, 1943-1952.	4.0	57
26	Evolutionary dynamics of genome segmentation in multipartite viruses. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 3812-3819.	2.6	54
27	Shape of Ecological Networks. <i>Physical Review Letters</i> , 2001, 86, 4418-4421.	7.8	49
28	From genotypes to organisms: State-of-the-art and perspectives of a cornerstone in evolutionary dynamics. <i>Physics of Life Reviews</i> , 2021, 38, 55-106.	2.8	49
29	Intermittency model for urban development. <i>Physical Review E</i> , 1998, 58, 295-302.	2.1	48
30	On the Genealogy of a Population of Biparental Individuals. <i>Journal of Theoretical Biology</i> , 2000, 203, 303-315.	1.7	48
31	Viral evolution. <i>Physics of Life Reviews</i> , 2006, 3, 65-92.	2.8	48
32	The impact of quasispecies dynamics on the use of therapeutics. <i>Trends in Microbiology</i> , 2012, 20, 595-603.	7.7	48
33	At the Boundary between Biological and Cultural Evolution: The Origin of Surname Distributions. <i>Journal of Theoretical Biology</i> , 2002, 216, 461-477.	1.7	44
34	Modular evolution and increase of functional complexity in replicating RNA molecules. <i>Rna</i> , 2006, 13, 97-107.	3.5	44
35	Disentangling the effects of selection and loss bias on gene dynamics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E5616-E5624.	7.1	44
36	On the structural repertoire of pools of short, random RNA sequences. <i>Journal of Theoretical Biology</i> , 2008, 252, 750-763.	1.7	43

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37	Internal Disequilibria and Phenotypic Diversification during Replication of Hepatitis C Virus in a Noncoevolving Cellular Environment. <i>Journal of Virology</i> , 2017, 91, .	3.4	42
38	Self-similarity in rain forests: Evidence for a critical state. <i>Physical Review E</i> , 1995, 51, 6250-6253.	2.1	41
39	On the networked architecture of genotype spaces and its critical effects on molecular evolution. <i>Open Biology</i> , 2018, 8, .	3.6	41
40	On Forest Spatial Dynamics with Gap Formation. <i>Journal of Theoretical Biology</i> , 1997, 187, 159-164.	1.7	40
41	Statistical Properties of Genealogical Trees. <i>Physical Review Letters</i> , 1999, 82, 1987-1990.	7.8	39
42	Tempo and mode of inhibitorâ€™mutagen antiviral therapies: A multidisciplinary approach. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 16008-16013.	7.1	38
43	Extinction: bad genes or weak chaos?. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1996, 263, 1407-1413.	2.6	37
44	Collective properties of evolving molecular quasispecies. <i>BMC Evolutionary Biology</i> , 2007, 7, 110.	3.2	36
45	Foot-and-Mouth Disease Virus Evolution: Exploring Pathways Towards Virus Extinction. <i>Current Topics in Microbiology and Immunology</i> , 2005, 288, 149-173.	1.1	35
46	Stochastic extinction of viral infectivity through the action of defectors. <i>Europhysics Letters</i> , 2009, 85, 18001.	2.0	34
47	Efficient HIV-1 inhibition by a 16 nt-long RNA aptamer designed by combining in vitro selection and in silico optimisation strategies. <i>Scientific Reports</i> , 2014, 4, 6242.	3.3	34
48	Fitness Distributions in Exponentially Growing Asexual Populations. <i>Physical Review Letters</i> , 2003, 90, 188102.	7.8	33
49	Large-Scale Genomic Analysis Suggests a Neutral Punctuated Dynamics of Transposable Elements in Bacterial Genomes. <i>PLoS Computational Biology</i> , 2014, 10, e1003680.	3.2	32
50	Evolutionary dynamics on networks of selectively neutral genotypes: Effects of topology and sequence stability. <i>Physical Review E</i> , 2009, 80, 066112.	2.1	30
51	Distribution of genotype network sizes in sequence-to-structure genotypeâ€™phenotype maps. <i>Journal of the Royal Society Interface</i> , 2017, 14, 20160976.	3.4	30
52	Self-organized criticality in rainforest dynamics. <i>Chaos, Solitons and Fractals</i> , 1996, 7, 523-541.	5.1	29
53	The vast unknown microbial biosphere. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6585-6587.	7.1	29
54	Small-world behaviour in a system of mobile elements. <i>Europhysics Letters</i> , 2001, 53, 693-699.	2.0	28

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55	Biodiversity in model ecosystems, II: species assembly and food web structure. <i>Journal of Theoretical Biology</i> , 2005, 235, 531-539.	1.7	28
56	River density and landscape roughness are universal determinants of linguistic diversity. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20133029.	2.6	27
57	Aftershock series of event February 18, 1996: An interpretation in terms of self-organized criticality. <i>Journal of Geophysical Research</i> , 1997, 102, 27407-27420.	3.3	25
58	Influence of Mutagenesis and Viral Load on the Sustained Low-Level Replication of an RNA Virus. <i>Journal of Molecular Biology</i> , 2011, 407, 60-78.	4.2	25
59	Criticality and unpredictability in macroevolution. <i>Physical Review E</i> , 1997, 55, 4500-4507.	2.1	24
60	TRANSIENT DYNAMICS AND SCALING PHENOMENA IN URBAN GROWTH. <i>Fractals</i> , 1999, 07, 1-8.	3.7	24
61	Statistical theory of phenotype abundance distributions: A test through exact enumeration of genotype spaces. <i>Europhysics Letters</i> , 2018, 123, 28001.	2.0	24
62	Distribution of repetitions of ancestors in genealogical trees. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2000, 281, 1-16.	2.6	23
63	Evolution on neutral networks accelerates the ticking rate of the molecular clock. <i>Journal of the Royal Society Interface</i> , 2015, 12, 20141010.	3.4	23
64	Scaling of voids and fractality in the galaxy distribution. <i>Monthly Notices of the Royal Astronomical Society</i> , 2002, 335, 977-983.	4.4	22
65	Phenotypic effect of mutations in evolving populations of RNA molecules. <i>BMC Evolutionary Biology</i> , 2010, 10, 46.	3.2	22
66	toyLIFE: a computational framework to study the multi-level organisation of the genotype-phenotype map. <i>Scientific Reports</i> , 2014, 4, 7549.	3.3	22
67	Adaptive multiscapes: an up-to-date metaphor to visualize molecular adaptation. <i>Biology Direct</i> , 2017, 12, 7.	4.6	22
68	Globally coupled logistic maps as dynamical glasses. <i>Europhysics Letters</i> , 2001, 53, 451-457.	2.0	20
69	Very long transients in globally coupled maps. <i>Europhysics Letters</i> , 2000, 50, 580-586.	2.0	19
70	Effects of Spatial Competition on the Diversity of a Quasispecies. <i>Physical Review Letters</i> , 2008, 100, 038106.	7.8	19
71	Enumerating secondary structures and structural moieties for circular RNAs. <i>Journal of Theoretical Biology</i> , 2017, 419, 375-382.	1.7	19
72	Adding levels of complexity enhances robustness and evolvability in a multilevel genotype-phenotype map. <i>Journal of the Royal Society Interface</i> , 2018, 15, 20170516.	3.4	19

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73	Diversity Patterns from Ecological Models at Dynamical Equilibrium. <i>Journal of Theoretical Biology</i> , 2001, 212, 11-34.	1.7	18
74	Variable Mutation Rates as an Adaptive Strategy in Replicator Populations. <i>PLoS ONE</i> , 2010, 5, e11186.	2.5	18
75	Theoretical approaches to disclosing the emergence and adaptive advantages of multipartite viruses. <i>Current Opinion in Virology</i> , 2018, 33, 89-95.	5.4	18
76	A trade-off between neutrality and adaptability limits the optimization of viral quasispecies. <i>Journal of Theoretical Biology</i> , 2009, 261, 148-155.	1.7	17
77	Topological properties of phylogenetic trees in evolutionary models. <i>European Physical Journal B</i> , 2009, 70, 583-592.	1.5	17
78	Struggle for Space: Viral Extinction through Competition for Cells. <i>Physical Review Letters</i> , 2011, 106, 028104.	7.8	17
79	Modelling viral evolution and adaptation: challenges and rewards. <i>Current Opinion in Virology</i> , 2012, 2, 531-537.	5.4	17
80	Populations of genetic circuits are unable to find the fittest solution in a multilevel genotype-phenotype map. <i>Journal of the Royal Society Interface</i> , 2020, 17, 20190843.	3.4	17
81	The space of genotypes is a network of networks: implications for evolutionary and extinction dynamics. <i>Scientific Reports</i> , 2017, 7, 13813.	3.3	16
82	Reconstructing evolutionary relationships from functional data: a consistent classification of organisms based on translation inhibition response. <i>Molecular Phylogenetics and Evolution</i> , 2005, 34, 371-381.	2.7	15
83	Genealogy in the Era of Genomics. <i>American Scientist</i> , 2003, 91, 158.	0.1	15
84	Motif frequency and evolutionary search times in RNA populations. <i>Journal of Theoretical Biology</i> , 2011, 280, 117-126.	1.7	14
85	A Simple Model of Large Scale Organization in Evolution. <i>International Journal of Modern Physics C</i> , 1998, 09, 1025-1032.	1.7	13
86	Fractal properties of isolines at varying altitude revealing different dominant geological processes on Earth. <i>Journal of Geophysical Research</i> , 2008, 113, .	3.3	13
87	Dynamical community structure of populations evolving on genotype networks. <i>Chaos, Solitons and Fractals</i> , 2015, 72, 99-106.	5.1	13
88	Long-range transport and universality classes in in vitro viral infection spread. <i>Europhysics Letters</i> , 2006, 74, 547-553.	2.0	12
89	Parsimonious Scenario for the Emergence of Viroid-Like Replicons De Novo. <i>Viruses</i> , 2019, 11, 425.	3.3	12
90	Long-Range Correlations in the Fossil Record and the Fractal Nature of Macroevolution. <i>International Journal of Modeling, Simulation, and Scientific Computing</i> , 1998, 01, 255-266.	1.4	11

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91	Motif selection in a model of evolving replicators: The role of surfaces and limited transport in network topology. <i>Europhysics Letters</i> , 2003, 64, 557-563.	2.0	11
92	Spiders: Water-Driven Erosive Structures in the Southern Hemisphere of Mars. <i>Astrobiology</i> , 2006, 6, 651-667.	3.0	11
93	Differences in adaptive dynamics determine the success of virus variants that propagate together. <i>Virus Evolution</i> , 2018, 4, vex043.	4.9	11
94	Severe Hindrance of Viral Infection Propagation in Spatially Extended Hosts. <i>PLoS ONE</i> , 2011, 6, e23358.	2.5	11
95	Tipping points and early warning signals in the genomic composition of populations induced by environmental changes. <i>Scientific Reports</i> , 2015, 5, 9664.	3.3	10
96	Endemicity and prevalence of multipartite viruses under heterogeneous between-host transmission. <i>PLoS Computational Biology</i> , 2019, 15, e1006876.	3.2	10
97	ON THE FRACTAL NATURE OF ECOLOGICAL AND MACROEVOLUTIONARY DYNAMICS. <i>Fractals</i> , 2001, 09, 1-16.	3.7	9
98	Out-of-equilibrium competitive dynamics of quasispecies. <i>Europhysics Letters</i> , 2007, 77, 38001.	2.0	9
99	Fat tails and black swans: Exact results for multiplicative processes with resets. <i>Chaos</i> , 2020, 30, 033104.	2.5	9
100	Individual risk-aversion responses tune epidemics to critical transmissibility ($\langle R \rangle = 1$). <i>Royal Society Open Science</i> , 2022, 9, 211667.	2.4	9
101	Evolutionary Dynamics in the RNA Bacteriophage $Q\beta^2$ Depends on the Pattern of Change in Selective Pressures. <i>Pathogens</i> , 2019, 8, 80.	2.8	8
102	Role of Demographic Dynamics and Conflict in the Population-Area Relationship for Human Languages. <i>PLoS ONE</i> , 2012, 7, e40137.	2.5	6
103	Zanette and Manrubia Reply:. <i>Physical Review Letters</i> , 1998, 80, 4831-4831.	7.8	5
104	SHAPE MATTERS: EFFECT OF POINT MUTATIONS ON RNA SECONDARY STRUCTURE. <i>International Journal of Modeling, Simulation, and Scientific Computing</i> , 2013, 16, 1250052.	1.4	5
105	New patterns in human biogeography revealed by networks of contacts between linguistic groups. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20142947.	2.6	5
106	Getting to Know Viral Evolutionary Strategies: Towards the Next Generation of Quasispecies Models. <i>Current Topics in Microbiology and Immunology</i> , 2015, 392, 201-217.	1.1	5
107	Replica-symmetry breaking in dynamical glasses. <i>European Physical Journal B</i> , 2001, 23, 497-508.	1.5	4
108	Limited role of spatial self-structuring in emergent trade-offs during pathogen evolution. <i>Scientific Reports</i> , 2018, 8, 12476.	3.3	4

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109	Unresolved advantages of multipartitism in spatially structured environments. <i>Virus Evolution</i> , 2021, 7, veab004.	4.9	4
110	Epistasis between cultural traits causes paradigm shifts in cultural evolution. <i>Royal Society Open Science</i> , 2020, 7, 191813.	2.4	3
111	Neutral networks of genotypes: evolution behind the curtain. <i>Arbor</i> , 2010, 186, 1051-1064.	0.3	3
112	The simple emergence of complex molecular function. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2022, 380, .	3.4	3
113	Neutral networks and chemical function in RNA. <i>Physics of Life Reviews</i> , 2012, 9, 277-278.	2.8	2
114	The Uncertain Future in How a Virus Spreads. <i>Physics Magazine</i> , 2020, 13, .	0.1	2
115	SYNCHRONIZATION AND CLUSTERING IN COUPLED SAW-TOOTH MAPS. <i>International Journal of Bifurcation and Chaos in Applied Sciences and Engineering</i> , 2000, 10, 2465-2478.	1.7	1
116	Supercritical branching processes and the role of fluctuations under exponential population growth. <i>Journal of Theoretical Biology</i> , 2003, 225, 497-505.	1.7	1
117	Evolution of fast mutating replicatorsâ€™ RNA viruses and the RNA world. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2006, 371, 80-83.	2.6	1
118	Demography-based adaptive network model reproduces the spatial organization of human linguistic groups. <i>Physical Review E</i> , 2015, 92, 062811.	2.1	1
119	What's in a name?. <i>Physics of Life Reviews</i> , 2013, 10, 424-425.	2.8	0
120	EMERGENCE AND SELECTION OF BIOMODULES: STEPS IN THE ASSEMBLY OF A PROTOCELL. <i>World Scientific Lecture Notes in Complex Systems</i> , 2013, , 323-343.	0.1	0
121	Populations of RNA Molecules as Computational Model for Evolution. , 2010, , 67-79.		0
122	The long and winding road to understanding organismal construction. <i>Physics of Life Reviews</i> , 2022, 42, 19-24.	2.8	0