

Chao Tang

List of Publications by Year in descending order

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Version: 2024-02-01

130
papers

22,558
citations

50276

46
h-index

17592

121
g-index

143
all docs

143
docs citations

143
times ranked

15190
citing authors

#	ARTICLE	IF	CITATIONS
1	Cell-to-cell variability in inducible Caspase9-mediated cell death. <i>Cell Death and Disease</i> , 2022, 13, 34.	6.3	5
2	Synthetic robust perfect adaptation achieved by negative feedback coupling with linear weak positive feedback. <i>Nucleic Acids Research</i> , 2022, 50, 2377-2386.	14.5	5
3	Computable early <i>Caenorhabditis elegans</i> embryo with a phase field model. <i>PLoS Computational Biology</i> , 2022, 18, e1009755.	3.2	10
4	Human pluripotent stem-cell-derived islets ameliorate diabetes in non-human primates. <i>Nature Medicine</i> , 2022, 28, 272-282.	30.7	55
5	Chemical Pretreatment Activated a Plastic State Amenable to Direct Lineage Reprogramming. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 865038.	3.7	6
6	A Multiclassifier System to Identify and Subtype Congenital Adrenal Hyperplasia Based on Circulating Steroid Hormones. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, e3304-e3312.	3.6	4
7	Whi5 is diluted and protein synthesis does not dramatically increase in pre-Start G1. <i>Molecular Biology of the Cell</i> , 2022, 33, 1t1.	2.1	13
8	Quantitative investigation reveals distinct phases in <i>Drosophila</i> sleep. <i>Communications Biology</i> , 2021, 4, 364.	4.4	6
9	Investigating Spatio-Temporal Cellular Interactions in Embryonic Morphogenesis by 4D Nucleus Tracking and Systematic Comparative Analysis Taking Nematodes <i>C. Elegans</i> and <i>C. Briggsae</i> as Examples. , 2021, , .		2
10	Finding gene network topologies for given biological function with recurrent neural network. <i>Nature Communications</i> , 2021, 12, 3125.	12.8	19
11	Red-emitting Zinc Probes with Minimal Phototoxicity for Multiplexed Recording of Orchestrated Insulin Secretion. <i>Angewandte Chemie</i> , 2021, 133, 26050.	2.0	1
12	Short-Term Plasticity Regulates Both Divisive Normalization and Adaptive Responses in <i>Drosophila</i> Olfactory System. <i>Frontiers in Computational Neuroscience</i> , 2021, 15, 730431.	2.1	2
13	Innentitelbild: Red-emitting Zinc Probes with Minimal Phototoxicity for Multiplexed Recording of Orchestrated Insulin Secretion (<i>Angew. Chem.</i> 49/2021). <i>Angewandte Chemie</i> , 2021, 133, 25790-25790.	2.0	0
14	Volume segregation programming in a nematode's early embryogenesis. <i>Physical Review E</i> , 2021, 104, 054409.	2.1	4
15	Why and how the nematode's early embryogenesis can be precise and robust: a mechanical perspective. <i>Physical Biology</i> , 2020, 17, 026001.	1.8	9
16	Protocol for Titrating Gene Expression Levels in Budding Yeast. <i>STAR Protocols</i> , 2020, 1, 100082.	1.2	1
17	Chemicals orchestrate reprogramming with hierarchical activation of master transcription factors primed by endogenous Sox17 activation. <i>Communications Biology</i> , 2020, 3, 629.	4.4	7
18	Establishment of a morphological atlas of the <i>Caenorhabditis elegans</i> embryo using deep-learning-based 4D segmentation. <i>Nature Communications</i> , 2020, 11, 6254.	12.8	45

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19	Reconstructing the multicellular structure of a developing metazoan embryo with repulsion-attraction model and cell-cell connection atlas in vivo. <i>Journal of Physics: Conference Series</i> , 2020, 1592, 012020.	0.4	2
20	Chemical perturbations reveal that RUVBL2 regulates the circadian phase in mammals. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	25
21	Multilevel regulation of muscle-specific transcription factor hlh-1 during <i>Caenorhabditis elegans</i> embryogenesis. <i>Development Genes and Evolution</i> , 2020, 230, 265-278.	0.9	3
22	Circulating re-entrant waves promote maturation of hiPSC-derived cardiomyocytes in self-organized tissue ring. <i>Communications Biology</i> , 2020, 3, 122.	4.4	32
23	Analysis of Circulating Waves in Tissue Rings derived from Human Induced Pluripotent Stem Cells. <i>Scientific Reports</i> , 2020, 10, 2984.	3.3	4
24	Critical slowing down and attractive manifold: A mechanism for dynamic robustness in the yeast cell-cycle process. <i>Physical Review E</i> , 2020, 101, 042405.	2.1	3
25	Computational study on ratio-sensing in yeast galactose utilization pathway. <i>PLoS Computational Biology</i> , 2020, 16, e1007960.	3.2	5
26	Cell Cycle Inhibitor Whi5 Records Environmental Information to Coordinate Growth and Division in Yeast. <i>Cell Reports</i> , 2019, 29, 987-994.e5.	6.4	38
27	Network Topologies That Can Achieve Dual Function of Adaptation and Noise Attenuation. <i>Cell Systems</i> , 2019, 9, 271-285.e7.	6.2	56
28	Optimal compressed sensing strategies for an array of nonlinear olfactory receptor neurons with and without spontaneous activity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20286-20295.	7.1	14
29	Growth strategy of microbes on mixed carbon sources. <i>Nature Communications</i> , 2019, 10, 1279.	12.8	105
30	Visualization of Genomic Loci in Living Cells with BiFC and STED. <i>Current Protocols in Cell Biology</i> , 2019, 82, e78.	2.3	2
31	Bi-functional biochemical networks. <i>Physical Biology</i> , 2019, 16, 016001.	1.8	7
32	Network Motifs Capable of Decoding Transcription Factor Dynamics. <i>Scientific Reports</i> , 2018, 8, 3594.	3.3	26
33	Early-warning signals of critical transition: Effect of extrinsic noise. <i>Physical Review E</i> , 2018, 97, 032406.	2.1	10
34	Low Cell-Matrix Adhesion Reveals Two Subtypes of Human Pluripotent Stem Cells. <i>Stem Cell Reports</i> , 2018, 11, 142-156.	4.8	37
35	A systematic study of the determinants of protein abundance memory in cell lineage. <i>Science Bulletin</i> , 2018, 63, 1051-1058.	9.0	1
36	Nanog induced intermediate state in regulating stem cell differentiation and reprogramming. <i>BMC Systems Biology</i> , 2018, 12, 22.	3.0	31

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37	Single-Cell RNA-Seq Reveals Dynamic Early Embryonic-like Programs during Chemical Reprogramming. <i>Cell Stem Cell</i> , 2018, 23, 31-45.e7.	11.1	122
38	Live visualization of genomic loci with BiFC-TALE. <i>Scientific Reports</i> , 2017, 7, 40192.	3.3	12
39	Adaptation with transcriptional regulation. <i>Scientific Reports</i> , 2017, 7, 42648.	3.3	25
40	Design of Tunable Oscillatory Dynamics in a Synthetic NF- κ B Signaling Circuit. <i>Cell Systems</i> , 2017, 5, 460-470.e5.	6.2	39
41	Odor-evoked inhibition of olfactory sensory neurons drives olfactory perception in <i>Drosophila</i> . <i>Nature Communications</i> , 2017, 8, 1357.	12.8	53
42	Adaptation through proportion. <i>Physical Biology</i> , 2016, 13, 046007.	1.8	4
43	Reliable cell cycle commitment in budding yeast is ensured by signal integration. <i>ELife</i> , 2015, 4, .	6.0	67
44	The Center for Quantitative Biology at Peking University. <i>Quantitative Biology</i> , 2015, 3, 1-3.	0.5	0
45	<i>Arabidopsis</i> DET1 degrades HFR1 but stabilizes PIF1 to precisely regulate seed germination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 3817-3822.	7.1	69
46	An Atlas of Network Topologies Reveals Design Principles for <i>Caenorhabditis elegans</i> Vulval Precursor Cell Fate Patterning. <i>PLoS ONE</i> , 2015, 10, e0131397.	2.5	2
47	Community detection for networks with unipartite and bipartite structure. <i>New Journal of Physics</i> , 2014, 16, 093001.	2.9	9
48	Multiple mechanisms determine the order of APC/C substrate degradation in mitosis. <i>Journal of Cell Biology</i> , 2014, 207, 23-39.	5.2	68
49	Costs and Benefits of Mutational Robustness in RNA Viruses. <i>Cell Reports</i> , 2014, 8, 1026-1036.	6.4	49
50	Synergistic and Antagonistic Drug Combinations Depend on Network Topology. <i>PLoS ONE</i> , 2014, 9, e93960.	2.5	99
51	QB: A new inter- and multi-disciplinary forum for modeling, engineering and understanding life. <i>Quantitative Biology</i> , 2013, 1, 1-2.	0.5	7
52	Bridging cross-cultural gaps in scientific exchange through innovative team challenge workshops. <i>Quantitative Biology</i> , 2013, 1, 3-8.	0.5	0
53	Generic properties of random gene regulatory networks. <i>Quantitative Biology</i> , 2013, 1, 253-260.	0.5	15
54	Design Principles of Regulatory Networks: Searching for the Molecular Algorithms of the Cell. <i>Molecular Cell</i> , 2013, 49, 202-212.	9.7	139

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55	Induction of Pluripotency in Mouse Somatic Cells with Lineage Specifiers. <i>Cell</i> , 2013, 153, 963-975.	28.9	272
56	Design Principles of the Yeast G1/S Switch. <i>PLoS Biology</i> , 2013, 11, e1001673.	5.6	51
57	A light-inducible organelle-targeting system for dynamically activating and inactivating signaling in budding yeast. <i>Molecular Biology of the Cell</i> , 2013, 24, 2419-2430.	2.1	90
58	Designing the Scientific Cradle for Quantitative Biologists. <i>ACS Synthetic Biology</i> , 2012, 1, 254-255.	3.8	3
59	Cell cycle synchronization by nutrient modulation. <i>Integrative Biology (United Kingdom)</i> , 2012, 4, 328.	1.3	21
60	Hierarchical Modularity and the Evolution of Genetic Interactomes across Species. <i>Molecular Cell</i> , 2012, 46, 691-704.	9.7	185
61	Designing Synthetic Regulatory Networks Capable of Self-Organizing Cell Polarization. <i>Cell</i> , 2012, 151, 320-332.	28.9	163
62	Flux Balance Analysis of Ammonia Assimilation Network in <i>E. coli</i> Predicts Preferred Regulation Point. <i>PLoS ONE</i> , 2011, 6, e16362.	2.5	9
63	Decision making of the p53 network: Death by integration. <i>Journal of Theoretical Biology</i> , 2011, 271, 205-211.	1.7	38
64	Defining Network Topologies that Can Achieve Biochemical Adaptation. <i>Cell</i> , 2009, 138, 760-773.	28.9	1,354
65	A more robust Boolean model describing inhibitor binding. <i>Frontiers of Electrical and Electronic Engineering in China: Selected Publications From Chinese Universities</i> , 2008, 3, 371-375.	0.6	0
66	Finding multiple target optimal intervention in disease-related molecular network. <i>Molecular Systems Biology</i> , 2008, 4, 228.	7.2	165
67	Robust, Tunable Biological Oscillations from Interlinked Positive and Negative Feedback Loops. <i>Science</i> , 2008, 321, 126-129.	12.6	602
68	Rationalizing translation attenuation in the network architecture of the unfolded protein response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 20280-20285.	7.1	51
69	SCUMBLE: a method for systematic and accurate detection of codon usage bias by maximum likelihood estimation. <i>Nucleic Acids Research</i> , 2008, 36, 3819-3827.	14.5	13
70	Dynamic Simulations on the Arachidonic Acid Metabolic Network. <i>PLoS Computational Biology</i> , 2007, 3, e55.	3.2	90
71	Hydrophobic interaction and hydrogen-bond network for a methane pair in liquid water. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 2626-2630.	7.1	78
72	Function constrains network architecture and dynamics: A case study on the yeast cell cycle Boolean network. <i>Physical Review E</i> , 2007, 75, 051907.	2.1	81

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73	Dynamic Studies of Scaffold-Dependent Mating Pathway in Yeast. <i>Biophysical Journal</i> , 2006, 91, 3986-4001.	0.5	28
74	Robustness and modular design of the <i>Drosophila</i> segment polarity network. <i>Molecular Systems Biology</i> , 2006, 2, 70.	7.2	114
75	Stochastic model of yeast cell-cycle network. <i>Physica D: Nonlinear Phenomena</i> , 2006, 219, 35-39.	2.8	67
76	Gibbs sampling and helix-cap motifs. <i>Nucleic Acids Research</i> , 2005, 33, 5343-5353.	14.5	10
77	Specificity of Trypsin and Chymotrypsin: Loop-Motion-Controlled Dynamic Correlation as a Determinant. <i>Biophysical Journal</i> , 2005, 89, 1183-1193.	0.5	104
78	Correlation between sequence hydrophobicity and surface-exposure pattern of database proteins. <i>Protein Science</i> , 2004, 13, 752-762.	7.6	90
79	Flexibility of β -sheets: Principal component analysis of database protein structures. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004, 55, 91-98.	2.6	43
80	Designability and thermal stability of protein structures. <i>Polymer</i> , 2004, 45, 699-705.	3.8	35
81	The yeast cell-cycle network is robustly designed. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 4781-4786.	7.1	953
82	Flexibility of α -Helices: Results of a Statistical Analysis of Database Protein Structures. <i>Journal of Molecular Biology</i> , 2003, 327, 229-237.	4.2	62
83	Structure space of model proteins: A principal component analysis. <i>Journal of Chemical Physics</i> , 2003, 118, 4277-4284.	3.0	8
84	Origin of scaling behavior of protein packing density: A sequential Monte Carlo study of compact long chain polymers. <i>Journal of Chemical Physics</i> , 2003, 118, 6102-6109.	3.0	56
85	Statistical mechanics of RNA folding: Importance of alphabet size. <i>Physical Review E</i> , 2003, 68, 041904.	2.1	11
86	Designability of α -helical proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 11163-11168.	7.1	28
87	Fast tree search for enumeration of a lattice model of protein folding. <i>Journal of Chemical Physics</i> , 2002, 116, 352.	3.0	29
88	Identifying proteins of high designability via surface-exposure patterns. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 47, 295-304.	2.6	14
89	Emergence of highly designable protein-backbone conformations in an off-lattice model. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 47, 506-512.	2.6	42
90	Designability of protein structures: A lattice-model study using the Miyazawa-Jernigan matrix. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 49, 403-412.	2.6	60

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91	The designability of protein structures. <i>Journal of Molecular Graphics and Modelling</i> , 2001, 19, 157-167.	2.4	56
92	Exact solution of a stochastic directed sandpile model. <i>Physical Review E</i> , 2001, 63, 026111.	2.1	28
93	Simple models of the protein folding problem. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2000, 288, 31-48.	2.6	35
94	Symmetry and designability for lattice protein models. <i>Journal of Chemical Physics</i> , 2000, 113, 8329-8336.	3.0	37
95	$1/f$ Noise in Bak-Tang-Wiesenfeld Models on Narrow Stripes. <i>Physical Review Letters</i> , 1999, 83, 2449-2452.	7.8	34
96	Designability, thermodynamic stability, and dynamics in protein folding: A lattice model study. <i>Journal of Chemical Physics</i> , 1999, 110, 1252-1262.	3.0	82
97	Incommensurability in the frustrated two-dimensionalXYmodel. <i>Physical Review B</i> , 1999, 60, 3163-3168.	3.2	26
98	Low-energy excitations and phase transitions in the frustrated two-dimensionalXYmodel. <i>Physical Review B</i> , 1998, 58, 6591-6607.	3.2	9
99	Nature of Driving Force for Protein Folding: A Result From Analyzing the Statistical Potential. <i>Physical Review Letters</i> , 1997, 79, 765-768.	7.8	195
100	Domain Walls and Phase Transitions in the Frustrated Two-DimensionalXYModel. <i>Physical Review Letters</i> , 1997, 79, 451-454.	7.8	17
101	Nature of Phase Transitions of Superconducting Wire Networks in a Magnetic Field. <i>Physical Review Letters</i> , 1996, 76, 2989-2992.	7.8	62
102	Peak effect in superconductors: melting of Larkin domains. <i>Europhysics Letters</i> , 1996, 35, 597-602.	2.0	42
103	Correction of partial-volume effects in phase-contrast flow measurements. <i>Journal of Magnetic Resonance Imaging</i> , 1995, 5, 175-180.	3.4	50
104	Tang, Feng, and Golubovic Reply:. <i>Physical Review Letters</i> , 1995, 74, 3500-3500.	7.8	1
105	Phases of Josephson Junction Ladders. <i>Physical Review Letters</i> , 1995, 75, 3930-3933.	7.8	44
106	Dynamics of a driven single flux line in superconductors. <i>Physical Review B</i> , 1995, 51, 8457-8461.	3.2	1
107	Self-Organized Criticality in Nonconserved Systems. <i>Physical Review Letters</i> , 1995, 74, 742-745.	7.8	112
108	Dynamics and noise spectra of a driven single flux line in superconductors. <i>Physical Review Letters</i> , 1994, 72, 1264-1267.	7.8	23

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109	Accuracy of phase-contrast flow measurements in the presence of partial-volume effects. Journal of Magnetic Resonance Imaging, 1993, 3, 377-385.	3.4	276
110	SOC and the Bean critical state. Physica A: Statistical Mechanics and Its Applications, 1993, 194, 315-320.	2.6	47
111	Patterns and scaling properties in a ballistic deposition model. Physical Review Letters, 1993, 71, 2769-2772.	7.8	18
112	Earthquakes as a Complex Phenomenon. Woodward Conference, 1992, , 209-220.	0.3	0
113	A forest-fire model and some thoughts on turbulence. Physics Letters, Section A: General, Atomic and Solid State Physics, 1990, 147, 297-300.	2.1	388
114	Droplet model for autocorrelation functions in an Ising ferromagnet. Physical Review A, 1989, 40, 995-1003.	2.5	27
115	Comment on "Relaxation at the Angle of Repose". Physical Review Letters, 1989, 62, 110-110.	7.8	6
116	A physicist's sandbox. Journal of Statistical Physics, 1989, 54, 1441-1458.	1.2	52
117	Earthquakes as a self-organized critical phenomenon. Journal of Geophysical Research, 1989, 94, 15635-15637.	3.3	1,020
118	Are Earthquakes, Fractals, and $1/f$ Noise Self-organized Critical Phenomena?. Springer Series in Synergetics, 1989, , 274-279.	0.4	2
119	Mean field theory of self-organized critical phenomena. Journal of Statistical Physics, 1988, 51, 797-802.	1.2	151
120	Critical Exponents and Scaling Relations for Self-Organized Critical Phenomena. Physical Review Letters, 1988, 60, 2347-2350.	7.8	360
121	Self-organized criticality. Physical Review A, 1988, 38, 364-374.	2.5	3,730
122	Scale Invariant Spatial and Temporal Fluctuations in Complex Systems. , 1988, , 329-335.		5
123	Self-organized Critical Phenomena. Series on Directions in Condensed Matter Physics, 1988, , 238-256.	0.1	0
124	Phase organization. Physical Review Letters, 1987, 58, 1161-1164.	7.8	98
125	Critical wave functions and a Cantor-set spectrum of a one-dimensional quasicrystal model. Physical Review B, 1987, 35, 1020-1033.	3.2	662
126	Self-organized criticality: An explanation of the $1/f$ noise. Physical Review Letters, 1987, 59, 381-384.	7.8	6,415

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127	Viscous flows in two dimensions. <i>Reviews of Modern Physics</i> , 1986, 58, 977-999.	45.6	674
128	Global scaling properties of the spectrum for a quasiperiodic schrödinger equation. <i>Physical Review B</i> , 1986, 34, 2041-2044.	3.2	165
129	Diffusion-limited aggregation and the Saffman-Taylor problem. <i>Physical Review A</i> , 1985, 31, 1977-1979.	2.5	181
130	Localization Problem in One Dimension: Mapping and Escape. <i>Physical Review Letters</i> , 1983, 50, 1870-1872.	7.8	1,018