

Feng Liu

List of Publications by Year in descending order

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36
papers

1,376
citations

430874

18
h-index

345221

36
g-index

36
all docs

36
docs citations

36
times ranked

1452
citing authors

#	ARTICLE	IF	CITATIONS
1	Modularization of grid cells constrained by the pyramidal patch lattice. <i>IScience</i> , 2021, 24, 102301.	4.1	2
2	Background synaptic input modulates the visuospatial working memory. <i>Physical Review E</i> , 2021, 104, 024416.	2.1	4
3	Synaptic mechanisms for motor variability in a feedforward network. <i>Science Advances</i> , 2020, 6, .	10.3	15
4	Gene transcription in bursting: a unified mode for realizing accuracy and stochasticity. <i>Biological Reviews</i> , 2019, 94, 248-258.	10.4	37
5	Modeling the regulation of p53 activation by HIF α upon hypoxia. <i>FEBS Letters</i> , 2019, 593, 2596-2611.	2.8	39
6	Roles of cellular heterogeneity, intrinsic and extrinsic noise in variability of p53 oscillation. <i>Scientific Reports</i> , 2019, 9, 5883.	3.3	15
7	Regulation of Tip60 α -dependent p53 acetylation in cell fate decision. <i>FEBS Letters</i> , 2019, 593, 13-22.	2.8	8
8	Modulation of dynamic modes by interplay between positive and negative feedback loops in gene regulatory networks. <i>Physical Review E</i> , 2018, 97, 042412.	2.1	15
9	Cell type α -dependent bimodal p53 activation engenders a dynamic mechanism of chemoresistance. <i>Science Advances</i> , 2018, 4, eaat5077.	10.3	28
10	Modeling the crosstalk between the circadian clock and ROS in <i>Neurospora crassa</i> . <i>Journal of Theoretical Biology</i> , 2018, 458, 125-132.	1.7	5
11	Modeling the response of a tumor-suppressive network to mitogenic and oncogenic signals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 5337-5342.	7.1	24
12	Kinetics of transcription initiation directed by multiple cis-regulatory elements on the <i>lnAp2</i> promoter. <i>Nucleic Acids Research</i> , 2016, 44, 10530-10538.	14.5	7
13	Impact of time delays on oscillatory dynamics of interlinked positive and negative feedback loops. <i>Physical Review E</i> , 2016, 94, 052413.	2.1	10
14	Realization of tristability in a multiplicatively coupled dual-loop genetic network. <i>Scientific Reports</i> , 2016, 6, 28096.	3.3	15
15	Modeling the interplay between the HIF-1 and p53 pathways in hypoxia. <i>Scientific Reports</i> , 2015, 5, 13834.	3.3	60
16	Structured Synaptic Inhibition Has a Critical Role in Multiple-Choice Motion-Discrimination Tasks. <i>Journal of Neuroscience</i> , 2014, 34, 13444-13457.	3.6	5
17	Reconciling the concurrent fast and slow cycling of proteins on gene promoters. <i>Journal of the Royal Society Interface</i> , 2014, 11, 20140253.	3.4	7
18	Involvement of miR-605 and miR-34a in the DNA Damage Response Promotes Apoptosis Induction. <i>Biophysical Journal</i> , 2014, 106, 1792-1800.	0.5	20

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19	A switch-like dynamic mechanism for the initiation of replicative senescence. <i>FEBS Letters</i> , 2014, 588, 4369-4374.	2.8	2
20	Interplay between Mdm2 and HIPK2 in the DNA damage response. <i>Journal of the Royal Society Interface</i> , 2014, 11, 20140319.	3.4	13
21	Coordination between p21 and DDB2 in the Cellular Response to UV Radiation. <i>PLoS ONE</i> , 2013, 8, e80111.	2.5	14
22	Dynamic Mechanism for the Transcription Apparatus Orchestrating Reliable Responses to Activators. <i>Scientific Reports</i> , 2012, 2, 422.	3.3	14
23	Regulation of the DNA Damage Response by p53 Cofactors. <i>Biophysical Journal</i> , 2012, 102, 2251-2260.	0.5	28
24	A Two-Step Mechanism for Cell Fate Decision by Coordination of Nuclear and Mitochondrial p53 Activities. <i>PLoS ONE</i> , 2012, 7, e38164.	2.5	31
25	Two-phase dynamics of p53 in the DNA damage response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 8990-8995.	7.1	275
26	Coordination between Cell Cycle Progression and Cell Fate Decision by the p53 and E2F1 Pathways in Response to DNA Damage. <i>Journal of Biological Chemistry</i> , 2010, 285, 31571-31580.	3.4	56
27	Coordination of the Nuclear and Cytoplasmic Activities of p53 in Response to DNA Damage. <i>Biophysical Journal</i> , 2010, 99, 1696-1705.	0.5	22
28	Interlinking positive and negative feedback loops creates a tunable motif in gene regulatory networks. <i>Physical Review E</i> , 2009, 80, 011926.	2.1	95
29	Signal propagation through feedforward neuronal networks with different operational modes. <i>Europhysics Letters</i> , 2009, 85, 38006.	2.0	11
30	Cell fate decision mediated by p53 pulses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 12245-12250.	7.1	211
31	Reversible Phosphorylation Subverts Robust Circadian Rhythms by Creating a Switch in Inactivating the Positive Element. <i>Biophysical Journal</i> , 2009, 97, 2867-2875.	0.5	9
32	Robustness analysis of cellular memory in an autoactivating positive feedback system. <i>FEBS Letters</i> , 2008, 582, 3776-3782.	2.8	42
33	A Common Cortical Circuit Mechanism for Perceptual Categorical Discrimination and Veridical Judgment. <i>PLoS Computational Biology</i> , 2008, 4, e1000253.	3.2	24
34	Linking fast and slow positive feedback loops creates an optimal bistable switch in cell signaling. <i>Physical Review E</i> , 2007, 76, 031924.	2.1	41
35	Propagation of Firing Rate in a Feed-Forward Neuronal Network. <i>Physical Review Letters</i> , 2006, 96, 018103.	7.8	75
36	Resonance-enhanced signal detection and transduction in the Hodgkin-Huxley neuronal systems. <i>Physical Review E</i> , 2001, 63, 021907.	2.1	97