## Ruben Perez-Carrasco

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/166814/publications.pdf

Version: 2024-02-01

26 papers

876 citations

687363 13 h-index 21 g-index

35 all docs 35 docs citations

35 times ranked

881 citing authors

#	Article	IF	Citations
1	Species-specific pace of development is associated with differences in protein stability. Science, 2020, 369, .	12.6	163
2	Combining a Toggle Switch and a Repressilator within the AC-DC Circuit Generates Distinct Dynamical Behaviors. Cell Systems, 2018, 6, 521-530.e3.	6.2	96
3	Catch bond drives stator mechanosensitivity in the bacterial flagellar motor. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 12952-12957.	7.1	78
4	A theoretical framework for the regulation of Shh morphogen-controlled gene expression. Development (Cambridge), 2014, 141, 3868-3878.	2.5	70
5	A Statistical Approach Reveals Designs for the Most Robust Stochastic Gene Oscillators. ACS Synthetic Biology, 2016, 5, 459-470.	3.8	61
6	Exact solution of stochastic gene expression models with bursting, cell cycle and replication dynamics. Physical Review E, 2020, 101, 032403.	2.1	61
7	Intrinsic Noise Profoundly Alters the Dynamics and Steady State of Morphogen-Controlled Bistable Genetic Switches. PLoS Computational Biology, 2016, 12, e1005154.	3.2	60
8	Effects of cell cycle variability on lineage and population measurements of messenger RNA abundance. Journal of the Royal Society Interface, 2020, 17, 20200360.	3.4	59
9	Controlling spatiotemporal pattern formation in a concentration gradient with a synthetic toggle switch. Molecular Systems Biology, 2020, 16, e9361.	7.2	42
10	Precision of tissue patterning is controlled by dynamical properties of gene regulatory networks. Development (Cambridge), 2021, 148, .	2.5	39
11	A multiscale model of epigenetic heterogeneity-driven cell fate decision-making. PLoS Computational Biology, 2019, 15, e1006592.	3.2	28
12	Memory functions reveal structural properties of gene regulatory networks. PLoS Computational Biology, 2018, 14, e1006003.	3.2	23
13	Neuronal differentiation influences progenitor arrangement in the vertebrate neuroepithelium. Development (Cambridge), 2019, 146, .	2.5	19
14	Minimum Action Path Theory Reveals the Details of Stochastic Transitions Out of Oscillatory States. Physical Review Letters, 2018, 120, 128102.	7.8	15
15	Degradation rate uniformity determines success of oscillations in repressive feedback regulatory networks. Journal of the Royal Society Interface, 2018, 15, 20180157.	3.4	13
16	Theoretical Analysis of the F1-ATPase Experimental Data. Biophysical Journal, 2010, 98, 2591-2600.	0.5	12
17	†Neighbourhood watch† $^{\text{TM}}$ model: embryonic epiblast cells assess positional information in relation to their neighbours. Development (Cambridge), 2022, 149, .	2.5	8
18	Fokker-Planck approach to molecular motors. Europhysics Letters, 2010, 91, 60001.	2.0	6

#	Article	IF	CITATIONS
19	Physics of molecular machines operated by a particle flux. Europhysics Letters, 2012, 100, 40001.	2.0	2
20	Relaxation time asymmetry in stator dynamics of the bacterial flagellar motor. Science Advances, 2022, $8$ , eabl $8112$ .	10.3	2
21	Deterministic and Stochastic Study for a Microscopic Angiogenesis Model: Applications to the Lewis Lung Carcinoma. PLoS ONE, 2016, 11, e0155553.	2.5	1
22	POWER AND EFFICIENCY OF F1-ATPase MOLECULAR MOTOR. Fluctuation and Noise Letters, 2012, 11, 1240003.	1.5	0
23	Mathematical models help explain experimental data. Response to â€Transcriptional interpretation of Shh morphogen signaling: computational modeling validates empirically established models'. Development (Cambridge), 2016, 143, 1640-1643.	2.5	0
24	de la Cruz etÂal. Reply. Physical Review Letters, 2019, 122, 059802.	7.8	0
25	Interdisciplinary approaches to dynamics in biology. Interface Focus, 2021, 11, .	3.0	0
26	Stator dynamics of the bacterial flagellar motor. Biophysical Journal, 2022, 121, 2a.	0.5	0