## CÄlfin C Guet

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/583941/publications.pdf

Version: 2024-02-01

35 papers 2,052 citations

394421 19 h-index 395702 33 g-index

47 all docs

47 docs citations

times ranked

47

2464 citing authors

#	Article	IF	CITATIONS
1	Predicting bacterial promoter function and evolution from random sequences. ELife, 2022, $11$ , .	6.0	17
2	Local genetic context shapes the function of a gene regulatory network. ELife, 2021, 10, .	6.0	15
3	Long lived transients in gene regulation. Theoretical Computer Science, 2021, 893, 1-16.	0.9	1
4	Sequential and Switchable Patterning for Studying Cellular Processes under Spatiotemporal Control. ACS Applied Materials & Samp; Interfaces, 2021, 13, 35545-35560.	8.0	1
5	Gene amplification as a form of population-level gene expression regulation. Nature Ecology and Evolution, 2020, 4, 612-625.	7.8	27
6	Molecular noise of innate immunity shapes bacteria-phage ecologies. PLoS Computational Biology, 2019, 15, e1007168.	3.2	7
7	Lack of cations in flow cytometry buffers affect fluorescence signals by reducing membrane stability and viability of Escherichia coli strains. Journal of Biotechnology, 2018, 268, 40-52.	3.8	11
8	Phage–host population dynamics promotes prophage acquisition in bacteria with innate immunity. Nature Ecology and Evolution, 2018, 2, 359-366.	7.8	56
9	Evolutionary potential of transcription factors for gene regulatory rewiring. Nature Ecology and Evolution, 2018, 2, 1633-1643.	7.8	25
10	Statistical mechanics for metabolic networks during steady state growth. Nature Communications, 2018, 9, 2988.	12.8	38
11	Leaky resistance and the conditions for the existence of lytic bacteriophage. PLoS Biology, 2018, 16, e2005971.	5.6	58
12	Biased partitioning of the multidrug efflux pump AcrAB-TolC underlies long-lived phenotypic heterogeneity. Science, 2017, 356, 311-315.	12.6	168
13	Effects of mutations in phage restriction sites during escape from restriction–modification. Biology Letters, 2017, 13, 20170646.	2.3	36
14	Shaping bacterial population behavior through computer-interfaced control of individual cells. Nature Communications, 2017, 8, 1535.	12.8	92
15	Model checking the evolution of gene regulatory networks. Acta Informatica, 2017, 54, 765-787.	0.5	12
16	Regulatory network structure determines patterns of intermolecular epistasis. ELife, 2017, 6, .	6.0	15
17	Bacterial flagella grow through an injection-diffusion mechanism. ELife, 2017, 6, .	6.0	66
18	Complex chromosomal neighborhood effects determine the adaptive potential of a gene under selection. ELife, 2017, 6, .	6.0	17

#	Article	IF	CITATIONS
19	On the mechanistic nature of epistasis in a canonical cis-regulatory element. ELife, 2017, 6, .	6.0	21
20	Intrinsic limits to gene regulation by global crosstalk. Nature Communications, 2016, 7, 12307.	12.8	63
21	Bacterial Autoimmunity Due to a Restriction-Modification System. Current Biology, 2016, 26, 404-409.	3.9	92
22	Epistatic Interactions in the Arabinose <i>Cis</i> -Regulatory Element. Molecular Biology and Evolution, 2016, 33, 761-769.	8.9	16
23	Variation of the folding and dynamics of the <i><scp>E</scp>scherichia coli</i> chromosome with growth conditions. Molecular Microbiology, 2012, 86, 1318-1333.	2.5	127
24	Structure and Dynamics of the Bacterial Chromosome in E. coli. Biophysical Journal, 2012, 102, 422a.	0.5	0
25	Noise Underlies Switching Behavior of the Bacterial Flagellum. Biophysical Journal, 2011, 101, 2336-2340.	0.5	23
26	Fine-Tuning of Chemotactic Response in E. coli Determined by High-Throughput Capillary Assay. Current Microbiology, 2011, 62, 764-769.	2.2	7
27	Interdependence of behavioural variability and response to small stimuli in bacteria. Nature, 2010, 468, 819-823.	27.8	67
28	Minimally invasive determination of mRNA concentration in single living bacteria. Nucleic Acids Research, 2008, 36, e73-e73.	14.5	47
29	Uncovering cis Regulatory Codes Using Synthetic Promoter Shuffling. PLoS ONE, 2008, 3, e2030.	2.5	42
30	Dynamical Determinants of Drug-Inducible Gene Expression in a Single Bacterium. Biophysical Journal, 2006, 90, 3315-3321.	0.5	20
31	Protein expression enhancement in efflux-deleted mutant bacteria. Protein Expression and Purification, 2006, 48, 28-31.	1.3	4
32	Real-time RNA profiling within a single bacterium. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 9160-9164.	7.1	90
33	Modeling network dynamics. Journal of Cell Biology, 2003, 161, 471-476.	5.2	195
34	Combinatorial Synthesis of Genetic Networks. Science, 2002, 296, 1466-1470.	12.6	480
35	Influence of M-phase chromatin on the anisotropy of microtubule asters Journal of Cell Biology, 1996, 133, 125-140.	5.2	86