Naama Barkai

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/3565925/publications.pdf

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28 papers	1,649 citations	361413 20 h-index	501196 28 g-index
37 all docs	37 docs citations	37 times ranked	2187 citing authors

#	Article	IF	Citations
1	Evolution of binding preferences among whole-genome duplicated transcription factors. ELife, 2022, 11, .	6.0	34
2	Rtt109 promotes nucleosome replacement ahead of the replication fork. Genome Research, 2022, 32, 1089-1098.	5.5	1
3	Measurement of histone replacement dynamics with genetically encoded exchange timers in yeast. Nature Biotechnology, 2021, 39, 1434-1443.	17.5	15
4	Gene Transcription as a Limiting Factor in Protein Production and Cell Growth. G3: Genes, Genomes, Genetics, 2020, 10, 3229-3242.	1.8	12
5	Dynamics of Spaetzle morphogen shuttling in the Drosophila embryo shapes gastrulation patterning. Development (Cambridge), 2019, 146, .	2.5	16
6	Resolving noiseâ€“control conflict by gene duplication. PLoS Biology, 2019, 17, e3000289.	5.6	60
7	A repressor-decay timer for robust temporal patterning in embryonic Drosophila neuroblast lineages. ELife, 2018, 7, .	6.0	31
8	Buffering Global Variability of Morphogen Gradients. Developmental Cell, 2017, 40, 429-438.	7.0	36
9	Hybrid vigor: The best of both parents, or a genomic clash?. Current Opinion in Systems Biology, 2017, 6, 22-27.	2.6	27
10	Principles of cellular resource allocation revealed by condition-dependent proteome profiling. ELife, 2017, 6, .	6.0	174
11	Expression homeostasis during DNA replication. Science, 2016, 351, 1087-1090.	12.6	101
12	A WntD-Dependent Integral Feedback Loop Attenuates Variability in Drosophila Toll Signaling. Developmental Cell, 2016, 36, 401-414.	7.0	36
13	Coordination of Gene Expression and Growth-Rate in Natural Populations of Budding Yeast. PLoS ONE, 2014, 9, e88801.	2.5	17
14	Systematic identification of cell size regulators in budding yeast. Molecular Systems Biology, 2014, 10, 761.	7.2	67
15	Loss of growth homeostasis by genetic decoupling of cell division from biomass growth: implication for size control mechanisms. Molecular Systems Biology, 2014, 10, 769.	7.2	11
16	Scaling of dorsalâ€ventral patterning in the Xenopus laevis embryo. BioEssays, 2014, 36, 151-156.	2.5	24
17	Disentangling signaling gradients generated by equivalent sources. Journal of Biological Physics, 2012, 38, 267-278.	1.5	20
18	The Competitive Advantage of a Dual-Transporter System. Science, 2011, 334, 1408-1412.	12.6	74

#	Article	IF	Citations
19	Scaling of morphogen gradients. Current Opinion in Genetics and Development, 2011, 21, 704-710.	3.3	74
20	Robust selection of sensory organ precursors by the Notchâ€“Delta pathway. Current Opinion in Cell Biology, 2011, 23, 663-667.	5.4	38
21	Coordination of gene expression with growth rate: A feedback or a feedâ€forward strategy?. FEBS Letters, 2009, 583, 3974-3978.	2.8	39
22	â€˜Big frog, small frogâ€™â€“ maintaining proportions in embryonic development. FEBS Journal, 2009, 276, 1196-1207.	4.7	31
23	Noise Propagation and Signaling Sensitivity in Biological Networks: A Role for Positive Feedback. PLoS Computational Biology, 2008, 4, e8.	3.2	180
24	Pre-Steady-State Decoding of the Bicoid Morphogen Gradient. PLoS Biology, 2007, 5, e46.	5.6	183
25	Variability and Robustness in Biomolecular Systems. Molecular Cell, 2007, 28, 755-760.	9.7	106
26	Strategy of Transcription Regulation in the Budding Yeast. PLoS ONE, 2007, 2, e250.	2.5	67
27	Comparative biology: beyond sequence analysis. Current Opinion in Biotechnology, 2007, 18, 371-377.	6.6	45
28	Comparative Gene Expression Analysis by a Differential Clustering Approach: Application to the Candida albicans Transcription Program. PLoS Genetics, 2005, 1, e39.	3.5	124