List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Revised Estimates for the Number of Human and Bacteria Cells in the Body. PLoS Biology, 2016, 14, e1002533.	5.6	3,388
2	Network motifs in the transcriptional regulation network of Escherichia coli. Nature Genetics, 2002, 31, 64-68.	21.4	2,603
3	The biomass distribution on Earth. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 6506-6511.	7.1	2,102
4	Are We Really Vastly Outnumbered? Revisiting the Ratio of Bacterial to Host Cells in Humans. Cell, 2016, 164, 337-340.	28.9	1,463
5	Superfamilies of Evolved and Designed Networks. Science, 2004, 303, 1538-1542.	12.6	1,182
6	Protection of BNT162b2 Vaccine Booster against Covid-19 in Israel. New England Journal of Medicine, 2021, 385, 1393-1400.	27.0	979
7	Waning Immunity after the BNT162b2 Vaccine in Israel. New England Journal of Medicine, 2021, 385, e85.	27.0	860
8	BioNumbers—the database of key numbers in molecular and cell biology. Nucleic Acids Research, 2010, 38, D750-D753.	14.5	859
9	SARS-CoV-2 (COVID-19) by the numbers. ELife, 2020, 9, .	6.0	826
10	The Moderately Efficient Enzyme: Evolutionary and Physicochemical Trends Shaping Enzyme Parameters. Biochemistry, 2011, 50, 4402-4410.	2.5	810
11	Cell Biology by the Numbers. , 0, , .		645
12	Oscillations and variability in the p53 system. Molecular Systems Biology, 2006, 2, 2006.0033.	7.2	539
13	Variability and memory of protein levels in human cells. Nature, 2006, 444, 643-646.	27.8	526
14	Cell-to-cell spread of HIV permits ongoing replication despite antiretroviral therapy. Nature, 2011, 477, 95-98.	27.8	502
15	eQuilibratorthe biochemical thermodynamics calculator. Nucleic Acids Research, 2012, 40, D770-D775.	14.5	483
16	Network motifs in integrated cellular networks of transcription-regulation and protein-protein interaction. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 5934-5939.	7.1	479
17	What is the total number of protein molecules per cell volume? A call to rethink some published values. BioEssays, 2013, 35, 1050-1055.	2.5	477
18	Glycolytic strategy as a tradeoff between energy yield and protein cost. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 10039-10044.	7.1	446

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19	A paxillin tyrosine phosphorylation switch regulates the assembly and form of cell-matrix adhesions. Journal of Cell Science, 2007, 120, 137-148.	2.0	402
20	Design and analysis of synthetic carbon fixation pathways. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 8889-8894.	7.1	402
21	Land, irrigation water, greenhouse gas, and reactive nitrogen burdens of meat, eggs, and dairy production in the United States. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 11996-12001.	7.1	375
22	Conversion of Escherichia coli to Generate All Biomass Carbon from CO2. Cell, 2019, 179, 1255-1263.e12.	28.9	352
23	Global human-made mass exceeds all living biomass. Nature, 2020, 588, 442-444.	27.8	344
24	Visual account of protein investment in cellular functions. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 8488-8493.	7.1	304
25	Protection by a Fourth Dose of BNT162b2 against Omicron in Israel. New England Journal of Medicine, 2022, 386, 1712-1720.	27.0	303
26	Protection and Waning of Natural and Hybrid Immunity to SARS-CoV-2. New England Journal of Medicine, 2022, 386, 2201-2212.	27.0	276
27	Sugar Synthesis from CO2 in Escherichia coli. Cell, 2016, 166, 115-125.	28.9	272
28	Cross-species analysis traces adaptation of Rubisco toward optimality in a low-dimensional landscape. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3475-3480.	7.1	249
29	Pathway Thermodynamics Highlights Kinetic Obstacles in Central Metabolism. PLoS Computational Biology, 2014, 10, e1003483.	3.2	249
30	Consistent Estimation of Gibbs Energy Using Component Contributions. PLoS Computational Biology, 2013, 9, e1003098.	3.2	231
31	A survey of carbon fixation pathways through a quantitative lens. Journal of Experimental Botany, 2012, 63, 2325-2342.	4.8	212
32	Global characterization of in vivo enzyme catalytic rates and their correspondence to in vitro <i>k</i> _{cat} measurements. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 3401-3406.	7.1	212
33	Rethinking glycolysis: on the biochemical logic of metabolic pathways. Nature Chemical Biology, 2012, 8, 509-517.	8.0	211
34	Central Carbon Metabolism as a Minimal Biochemical Walk between Precursors for Biomass and Energy. Molecular Cell, 2010, 39, 809-820.	9.7	208
35	SnapShot: Key Numbers in Biology. Cell, 2010, 141, 1262-1262.e1.	28.9	206
36	The distribution of cellular turnover in the human body. Nature Medicine, 2021, 27, 45-48.	30.7	205

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37	The total number and mass of SARS-CoV-2 virions. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	187
38	Protection against Covid-19 by BNT162b2 Booster across Age Groups. New England Journal of Medicine, 2021, 385, 2421-2430.	27.0	185
39	Promoters maintain their relative activity levels under different growth conditions. Molecular Systems Biology, 2013, 9, 701.	7.2	181
40	A central role for Necl4 (SynCAM4) in Schwann cell–axon interaction and myelination. Nature Neuroscience, 2007, 10, 861-869.	14.8	178
41	Input–output robustness in simple bacterial signaling systems. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 19931-19935.	7.1	170
42	Massively Parallel Interrogation of the Effects of Gene Expression Levels on Fitness. Cell, 2016, 166, 1282-1294.e18.	28.9	168
43	HSP90 affects the expression of genetic variation and developmental stability in quantitative traits. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 2963-2968.	7.1	167
44	pH determines the energetic efficiency of the cyanobacterial CO ₂ concentrating mechanism. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E5354-62.	7.1	166
45	Spanning high-dimensional expression space using ribosome-binding site combinatorics. Nucleic Acids Research, 2013, 41, e98-e98.	14.5	165
46	The global mass and average rate of rubisco. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 4738-4743.	7.1	154
47	Design and analysis of metabolic pathways supporting formatotrophic growth for electricity-dependent cultivation of microbes. Biochimica Et Biophysica Acta - Bioenergetics, 2013, 1827, 1039-1047.	1.0	150
48	Prediction of Microbial Growth Rate versus Biomass Yield by a Metabolic Network with Kinetic Parameters. PLoS Computational Biology, 2012, 8, e1002575.	3.2	148
49	The opportunity cost of animal based diets exceeds all food losses. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 3804-3809.	7.1	144
50	The Protein Cost of Metabolic Fluxes: Prediction from Enzymatic Rate Laws and Cost Minimization. PLoS Computational Biology, 2016, 12, e1005167.	3.2	144
51	Revisiting Trade-offs between Rubisco Kinetic Parameters. Biochemistry, 2019, 58, 3365-3376.	2.5	142
52	Estimating disease severity of Omicron and Delta SARS-CoV-2 infections. Nature Reviews Immunology, 2022, 22, 267-269.	22.7	138
53	Thermodynamic constraints shape the structure of carbon fixation pathways. Biochimica Et Biophysica Acta - Bioenergetics, 2012, 1817, 1646-1659.	1.0	126
54	Dynamic proteomics in individual human cells uncovers widespread cell-cycle dependence of nuclear proteins. Nature Methods, 2006, 3, 525-531.	19.0	125

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55	An integrated open framework for thermodynamics of reactions that combines accuracy and coverage. Bioinformatics, 2012, 28, 2037-2044.	4.1	108
56	A note on the kinetics of enzyme action: A decomposition that highlights thermodynamic effects. FEBS Letters, 2013, 587, 2772-2777.	2.8	108
57	A feeling for the numbers in biology. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 21465-21471.	7.1	100
58	Quantifying Translational Coupling in <i>E. coli</i> Synthetic Operons Using RBS Modulation and Fluorescent Reporters. ACS Synthetic Biology, 2013, 2, 327-336.	3.8	100
59	Coarse-graining and self-dissimilarity of complex networks. Physical Review E, 2005, 71, 016127.	2.1	92
60	The Moderately Efficient Enzyme: Futile Encounters and Enzyme Floppiness. Biochemistry, 2015, 54, 4969-4977.	2.5	89
61	Energetic cost of building a virus. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4324-E4333.	7.1	89
62	A Bird's-Eye View of Enzyme Evolution: Chemical, Physicochemical, and Physiological Considerations. Chemical Reviews, 2018, 118, 8786-8797.	47.7	88
63	Noise in gene expression is coupled to growth rate. Genome Research, 2015, 25, 1893-1902.	5.5	83
64	Highly active rubiscos discovered by systematic interrogation of natural sequence diversity. EMBO Journal, 2020, 39, e104081.	7.8	72
65	Functional reconstitution of a bacterial CO2 concentrating mechanism in Escherichia coli. ELife, 2020, 9, .	6.0	72
66	Design principles of autocatalytic cycles constrain enzyme kinetics and force low substrate saturation at flux branch points. ELife, 2017, 6, .	6.0	70
67	eQuilibrator 3.0: a database solution for thermodynamic constant estimation. Nucleic Acids Research, 2022, 50, D603-D609.	14.5	70
68	Photovoltaic-driven microbial protein production can use land and sunlight more efficiently than conventional crops. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	68
69	Steady-State Metabolite Concentrations Reflect a Balance between Maximizing Enzyme Efficiency and Minimizing Total Metabolite Load. PLoS ONE, 2013, 8, e75370.	2.5	67
70	The Biomass Composition of the Oceans: A Blueprint of Our Blue Planet. Cell, 2019, 179, 1451-1454.	28.9	67
71	Omicron infection enhances Delta antibody immunity in vaccinated persons. Nature, 2022, 607, 356-359.	27.8	66
72	Hydrophobicity and Charge Shape Cellular Metabolite Concentrations. PLoS Computational Biology, 2011, 7, e1002166.	3.2	65

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73	Lessons on enzyme kinetics from quantitative proteomics. Current Opinion in Biotechnology, 2017, 46, 81-89.	6.6	64
74	Generation of a fluorescently labeled endogenous protein library in living human cells. Nature Protocols, 2007, 2, 1515-1527.	12.0	62
75	Achieving Diversity in the Face of Constraints: Lessons from Metabolism. Science, 2012, 336, 1663-1667.	12.6	61
76	Pyruvate Formate-Lyase Enables Efficient Growth of <i>Escherichia coli</i> on Acetate and Formate. Biochemistry, 2016, 55, 2423-2426.	2.5	57
77	Protein Dynamics in Individual Human Cells: Experiment and Theory. PLoS ONE, 2009, 4, e4901.	2.5	54
78	Reconstructing a puzzle: existence of cyanophages containing both photosystemâ€l and photosystemâ€l gene suites inferred from oceanic metagenomic datasets. Environmental Microbiology, 2011, 13, 24-32.	3.8	46
79	The quantified cell. Molecular Biology of the Cell, 2014, 25, 3497-3500.	2.1	44
80	The genetic basis for the adaptation of E. coli to sugar synthesis from CO2. Nature Communications, 2017, 8, 1705.	12.8	39
81	Engineering carbon fixation in E. coli : from heterologous RuBisCO expression to the Calvin–Benson–Bassham cycle. Current Opinion in Biotechnology, 2017, 47, 83-91.	6.6	38
82	The relationship between evolutionary and physiological variation in hemoglobin. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16998-17003.	7.1	37
83	Environmentally Optimal, Nutritionally Aware Beef Replacement Plant-Based Diets. Environmental Science & Technology, 2016, 50, 8164-8168.	10.0	28
84	What governs the reaction center excitation wavelength of photosystems I and II?. Photosynthesis Research, 2009, 101, 59-67.	2.9	23
85	A model for â€~sustainable' US beef production. Nature Ecology and Evolution, 2018, 2, 81-85.	7.8	23
86	Efficiency in Evolutionary Trade-Offs. Science, 2012, 336, 1114-1115.	12.6	22
87	A proof for loop-law constraints in stoichiometric metabolic networks. BMC Systems Biology, 2012, 6, 140.	3.0	21
88	Noise Genetics: Inferring Protein Function by Correlating Phenotype with Protein Levels and Localization in Individual Human Cells. PLoS Genetics, 2014, 10, e1004176.	3.5	20
89	Towards a quantitative view of the global ubiquity of biofilms. Nature Reviews Microbiology, 2019, 17, 199-200.	28.6	20
90	A Minimalistic Resource Allocation Model to Explain Ubiquitous Increase in Protein Expression with Growth Rate. PLoS ONE, 2016, 11, e0153344.	2.5	18

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91	Dynamic Proteomics of Human Protein Level and Localization across the Cell Cycle. PLoS ONE, 2012, 7, e48722.	2.5	17
92	An In Vivo Metabolic Approach for Deciphering the Product Specificity of Glycerate Kinase Proves that Both E. coli's Glycerate Kinases Generate 2-Phosphoglycerate. PLoS ONE, 2015, 10, e0122957.	2.5	15
93	Point mutations in topoisomerase I alter the mutation spectrum in E. coli and impact the emergence of drug resistance genotypes. Nucleic Acids Research, 2020, 48, 761-769.	14.5	13
94	Robust Control of PEP Formation Rate in the Carbon Fixation Pathway of C4 Plants by a Bi-functional Enzyme. BMC Systems Biology, 2011, 5, 171.	3.0	10
95	Protection following BNT162b2 booster in adolescents substantially exceeds that of a fresh 2-dose vaccine. Nature Communications, 2022, 13, 1971.	12.8	10
96	RECURRING HARMONIC WALKS AND NETWORK MOTIFS IN WESTERN MUSIC. International Journal of Modeling, Simulation, and Scientific Computing, 2006, 09, 121-132.	1.4	5
97	Dynamic proteomics in mammalian cells: capabilities and challenges. Molecular BioSystems, 2007, 3, 542.	2.9	1
98	Reply to Metson et al.: The importance of phosphorus perturbations. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4908-E4908.	7.1	0
99	Reply to Tichenor: Proposed update to beef greenhouse gas footprint is numerically questionable and well within current uncertainty bounds. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, F822-F823	7.1	Ο