## Gene-Wei Li

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4745190/publications.pdf

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	687363	940533
5,867	13	16
citations	h-index	g-index
0.1	0.1	0745
21	21	8745
docs citations	times ranked	citing authors
	5,867 citations  21 docs citations	5,867 13 h-index  21 21

#	Article	IF	Citations
1	Dynamic Imaging of Genomic Loci in Living Human Cells by an Optimized CRISPR/Cas System. Cell, 2013, 155, 1479-1491.	28.9	1,695
2	Quantifying Absolute Protein Synthesis Rates Reveals Principles Underlying Allocation of Cellular Resources. Cell, 2014, 157, 624-635.	28.9	1,137
3	Probing Transcription Factor Dynamics at the Single-Molecule Level in a Living Cell. Science, 2007, 316, 1191-1194.	12.6	966
4	The anti-Shine–Dalgarno sequence drives translational pausing and codon choice in bacteria. Nature, 2012, 484, 538-541.	27.8	566
5	Central dogma at the single-molecule level in living cells. Nature, 2011, 475, 308-315.	27.8	378
6	Chromosome Organization by a Nucleoid-Associated Protein in Live Bacteria. Science, 2011, 333, 1445-1449.	12.6	341
7	Effects of macromolecular crowding and DNA looping on gene regulation kinetics. Nature Physics, 2009, 5, 294-297.	16.7	202
8	A Stress Response that Monitors and Regulates mRNA Structure Is Central to Cold Shock Adaptation. Molecular Cell, 2018, 70, 274-286.e7.	9.7	157
9	The Coding and Noncoding Architecture of the Caulobacter crescentus Genome. PLoS Genetics, 2014, 10, e1004463.	3.5	136
10	Functionally uncoupled transcription–translation in Bacillus subtilis. Nature, 2020, 585, 124-128.	27.8	109
11	From coarse to fine: the absolute <i>Escherichia coli</i> proteome under diverse growth conditions. Molecular Systems Biology, 2021, 17, e9536.	7.2	82
12	Dynamic translation regulation in <i>Caulobacter</i> cell cycle control. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E6859-E6867.	7.1	57
13	Single molecule approaches to transcription factor kinetics in living cells. FEBS Letters, 2009, 583, 3979-3983.	2.8	20
14	Spurious regulatory connections dictate the expressionâ€fitness landscape of translation factors. Molecular Systems Biology, 2021, 17, e10302.	7.2	8
15	Differential translation of mRNA isoforms transcribed with distinct sigma factors. Rna, 2021, 27, 791-804.	3.5	4