

# Matthew J Culyba

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1594058/publications.pdf>

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times ranked

356  
citing authors

#	ARTICLE	IF	CITATIONS
1	Targets for Combating the Evolution of Acquired Antibiotic Resistance. <i>Biochemistry</i> , 2015, 54, 3573-3582.	1.2	101
2	Systematically Altering Bacterial SOS Activity under Stress Reveals Therapeutic Strategies for Potentiating Antibiotics. <i>MSphere</i> , 2016, 1, .	1.3	74
3	Inhibitors of LexA Autoproteolysis and the Bacterial SOS Response Discovered by an Academic-Industry Partnership. <i>ACS Infectious Diseases</i> , 2018, 4, 349-359.	1.8	45
4	Non-equilibrium repressor binding kinetics link DNA damage dose to transcriptional timing within the SOS gene network. <i>PLoS Genetics</i> , 2018, 14, e1007405.	1.5	37
5	Bacterial evolution during human infection: Adapt and live or adapt and die. <i>PLoS Pathogens</i> , 2021, 17, e1009872.	2.1	33
6	Advancement of the 5-Amino-1-(Carbamoylmethyl)-1H-1,2,3-Triazole-4-Carboxamide Scaffold to Disarm the Bacterial SOS Response. <i>Frontiers in Microbiology</i> , 2018, 9, 2961.	1.5	22
7	Convergent Evolution of Antibiotic Tolerance in Patients with Persistent Methicillin-Resistant <i>Staphylococcus aureus</i> Bacteremia. <i>Infection and Immunity</i> , 2022, 90, e0000122.	1.0	8
8	Ordering up gene expression by slowing down transcription factor binding kinetics. <i>Current Genetics</i> , 2019, 65, 401-406.	0.8	7
9	The Parameter-Fitness Landscape of <i>lexA</i> Autoregulation in <i>Escherichia coli</i> . <i>MSphere</i> , 2020, 5, .	1.3	7
10	A Small-Molecule Inducible Synthetic Circuit for Control of the SOS Gene Network without DNA Damage. <i>ACS Synthetic Biology</i> , 2017, 6, 2067-2076.	1.9	4
11	DNA cytosine methylation at the <i>lexA</i> promoter of <i>Escherichia coli</i> is stationary phase specific. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	0.8	4
12	Effect of mismatch repair on the mutational footprint of the bacterial SOS mutator activity. <i>DNA Repair</i> , 2021, 103, 103130.	1.3	3