

Henrik Strahl

List of Publications by Year in descending order

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33
papers

3,106
citations

257450

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361022

35
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44
docs citations

44
times ranked

4338
citing authors

#	ARTICLE	IF	CITATIONS
1	Regulation of para-cresol production in <i>Clostridioides difficile</i> . <i>Current Opinion in Microbiology</i> , 2022, 65, 131-137.	5.1	1
2	Low membrane fluidity triggers lipid phase separation and protein segregation in living bacteria. <i>EMBO Journal</i> , 2022, 41, e109800.	7.8	52
3	A hyperpromiscuous antitoxin protein domain for the neutralization of diverse toxin domains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	22
4	Disruption of the Cytoplasmic Membrane Structure and Barrier Function Underlies the Potent Antiseptic Activity of Octenidine in Gram-Positive Bacteria. <i>Applied and Environmental Microbiology</i> , 2022, 88, e0018022.	3.1	9
5	A membrane-depolarizing toxin substrate of the <i>Staphylococcus aureus</i> type VII secretion system mediates intraspecies competition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 20836-20847.	7.1	57
6	A widespread toxin-antitoxin system exploiting growth control via alarmone signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 10500-10510.	7.1	81
7	The Epipeptide YydF Intrinsically Triggers the Cell Envelope Stress Response of <i>Bacillus subtilis</i> and Causes Severe Membrane Perturbations. <i>Frontiers in Microbiology</i> , 2020, 11, 151.	3.5	29
8	Metabolism of multiple glycosaminoglycans by <i>Bacteroides thetaiotaomicron</i> is orchestrated by a versatile core genetic locus. <i>Nature Communications</i> , 2020, 11, 646.	12.8	58
9	Extreme slow growth as alternative strategy to survive deep starvation in bacteria. <i>Nature Communications</i> , 2019, 10, 890.	12.8	153
10	A family of Type VI secretion system effector proteins that form ion-selective pores. <i>Nature Communications</i> , 2019, 10, 5484.	12.8	57
11	ABC ATPases Involved in Protein Synthesis, Ribosome Assembly and Antibiotic Resistance: Structural and Functional Diversification across the Tree of Life. <i>Journal of Molecular Biology</i> , 2019, 431, 3568-3590.	4.2	90
12	Membrane Curvature and the Tol-Pal Complex Determine Polar Localization of the Chemoreceptor Tar in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2018, 200, .	2.2	12
13	Segregation of mitochondrial DNA heteroplasmy through a developmental genetic bottleneck in human embryos. <i>Nature Cell Biology</i> , 2018, 20, 144-151.	10.3	182
14	Mode of Action and Heterologous Expression of the Natural Product Antibiotic Vancoresmycin. <i>ACS Chemical Biology</i> , 2018, 13, 207-214.	3.4	50
15	The type VI secretion system deploys antifungal effectors against microbial competitors. <i>Nature Microbiology</i> , 2018, 3, 920-931.	13.3	199
16	The Gram-positive model organism <i>Bacillus subtilis</i> does not form microscopically detectable cardiolipin-specific lipid domains. <i>Microbiology (United Kingdom)</i> , 2018, 164, 475-482.	1.8	15
17	Assessing Membrane Fluidity and Visualizing Fluid Membrane Domains in Bacteria Using Fluorescent Membrane Dyes. <i>Bio-protocol</i> , 2018, 8, e3063.	0.4	31
18	Growth rate control of flagellar assembly in <i>Escherichia coli</i> strain RP437. <i>Scientific Reports</i> , 2017, 7, 41189.	3.3	45

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19	Antimicrobial peptide cWFW kills by combining lipid phase separation with autolysis. <i>Scientific Reports</i> , 2017, 7, 44332.	3.3	98
20	Bacterial Membranes: Structure, Domains, and Function. <i>Annual Review of Microbiology</i> , 2017, 71, 519-538.	7.3	178
21	Measurement of Cell Membrane Fluidity by Laurdan GP: Fluorescence Spectroscopy and Microscopy. <i>Methods in Molecular Biology</i> , 2017, 1520, 159-174.	0.9	47
22	Analysis of Antimicrobial-Triggered Membrane Depolarization Using Voltage Sensitive Dyes. <i>Frontiers in Cell and Developmental Biology</i> , 2016, 4, 29.	3.7	207
23	Daptomycin inhibits cell envelope synthesis by interfering with fluid membrane microdomains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E7077-E7086.	7.1	326
24	Against the mainstream: the membrane-associated type I toxin BsrG from <i>Bacillus subtilis</i> interferes with cell envelope biosynthesis without increasing membrane permeability. <i>Molecular Microbiology</i> , 2015, 98, 651-666.	2.5	54
25	Membrane Recognition and Dynamics of the RNA Degradosome. <i>PLoS Genetics</i> , 2015, 11, e1004961.	3.5	93
26	The actin homologue MreB organizes the bacterial cell membrane. <i>Nature Communications</i> , 2014, 5, 3442.	12.8	223
27	Time-Delayed In Vivo Assembly of Subunit a into Preformed <i>Escherichia coli</i> FoF1 ATP Synthase. <i>Journal of Bacteriology</i> , 2013, 195, 4074-4084.	2.2	13
28	Structural and genetic analyses reveal the protein SepF as a new membrane anchor for the Z ring. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E4601-10.	7.1	116
29	Finding the corners in a cell. <i>Current Opinion in Microbiology</i> , 2012, 15, 731-736.	5.1	31
30	Archaeal transcriptional regulation of the prokaryotic KdpFABC complex mediating K ⁺ uptake in <i>H. salinarum</i> . <i>Extremophiles</i> , 2011, 15, 643-652.	2.3	11
31	Membrane potential is important for bacterial cell division. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 12281-12286.	7.1	426
32	The extremely halophilic archaeon <i>Halobacterium salinarum</i> R1 responds to potassium limitation by expression of the K ⁺ -transporting KdpFABC P-type ATPase and by a decrease in intracellular K ⁺ . <i>Extremophiles</i> , 2008, 12, 741-752.	2.3	39
33	Localization of general and regulatory proteolysis in <i>Bacillus subtilis</i> cells. <i>Molecular Microbiology</i> , 2008, 70, 682-694.	2.5	48