

Morgan Beeby

List of Publications by Year in descending order

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Version: 2024-02-01

63
papers

3,373
citations

218677

26
h-index

155660

55
g-index

80
all docs

80
docs citations

80
times ranked

3745
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein Structures Forming the Shell of Primitive Bacterial Organelles. <i>Science</i> , 2005, 309, 936-938.	12.6	420
2	Crystal Structure of 3-Amino-5-hydroxybenzoic Acid (AHBA) Synthase. <i>Biochemistry</i> , 1999, 38, 9840-9849.	2.5	291
3	Structural diversity of bacterial flagellar motors. <i>EMBO Journal</i> , 2011, 30, 2972-2981.	7.8	281
4	The flagellum in bacterial pathogens: For motility and a whole lot more. <i>Seminars in Cell and Developmental Biology</i> , 2015, 46, 91-103.	5.0	275
5	Architecture of the major component of the type III secretion system export apparatus. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 99-104.	8.2	200
6	The Genomics of Disulfide Bonding and Protein Stabilization in Thermophiles. <i>PLoS Biology</i> , 2005, 3, e309.	5.6	180
7	Diverse high-torque bacterial flagellar motors assemble wider stator rings using a conserved protein scaffold. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E1917-26.	7.1	170
8	Architecture and assembly of the γ -positive cell wall. <i>Molecular Microbiology</i> , 2013, 88, 664-672.	2.5	116
9	Communication across the bacterial cell envelope depends on the size of the periplasm. <i>PLoS Biology</i> , 2017, 15, e2004303.	5.6	108
10	Escherichia coli Peptidoglycan Structure and Mechanics as Predicted by Atomic-Scale Simulations. <i>PLoS Computational Biology</i> , 2014, 10, e1003475.	3.2	92
11	Nanoscale-length control of the flagellar driveshaft requires hitting the tethered outer membrane. <i>Science</i> , 2017, 356, 197-200.	12.6	86
12	β -proteobacteria eject their polar flagella under nutrient depletion, retaining flagellar motor relic structures. <i>PLoS Biology</i> , 2019, 17, e3000165.	5.6	72
13	Evolution of higher torque in Campylobacter-type bacterial flagellar motors. <i>Scientific Reports</i> , 2018, 8, 97.	3.3	70
14	Growth and Localization of Polyhydroxybutyrate Granules in <i>Ralstonia eutropha</i> . <i>Journal of Bacteriology</i> , 2012, 194, 1092-1099.	2.2	65
15	Propulsive nanomachines: the convergent evolution of archaella, flagella and cilia. <i>FEMS Microbiology Reviews</i> , 2020, 44, 253-304.	8.6	60
16	Visualization and interpretation of protein networks in <i>Mycobacterium tuberculosis</i> based on hierarchical clustering of genome-wide functional linkage maps. <i>Nucleic Acids Research</i> , 2003, 31, 7099-7109.	14.5	55
17	Short FtsZ filaments can drive asymmetric cell envelope constriction at the onset of bacterial cytokinesis. <i>EMBO Journal</i> , 2017, 36, 1577-1589.	7.8	55
18	Coarse-grained simulations of bacterial cell wall growth reveal that local coordination alone can be sufficient to maintain rod shape. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E3689-98.	7.1	50

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19	Expanding the archaeellum regulatory network - the eukaryotic protein kinases ArnC and ArnD influence motility of <i>Sulfolobus acidocaldarius</i> . <i>MicrobiologyOpen</i> , 2017, 6, e00414.	3.0	45
20	Motility in the epsilon-proteobacteria. <i>Current Opinion in Microbiology</i> , 2015, 28, 115-121.	5.1	40
21	<i>Campylobacter jejuni</i> motility integrates specialized cell shape, flagellar filament, and motor, to coordinate action of its opposed flagella. <i>PLoS Pathogens</i> , 2020, 16, e1008620.	4.7	40
22	Inter-membrane association of the Sec and BAM translocons for bacterial outer-membrane biogenesis. <i>ELife</i> , 2020, 9, .	6.0	39
23	Activated chemoreceptor arrays remain intact and hexagonally packed. <i>Molecular Microbiology</i> , 2011, 82, 748-757.	2.5	38
24	Adenita: interactive 3D modelling and visualization of DNA nanostructures. <i>Nucleic Acids Research</i> , 2020, 48, 8269-8275.	14.5	33
25	The structure of the periplasmic FlaF-FlaF complex and its essential role for archaeellar swimming motility. <i>Nature Microbiology</i> , 2020, 5, 216-225.	13.3	32
26	Exploiting genomic patterns to discover new supramolecular protein assemblies. <i>Protein Science</i> , 2009, 18, 69-79.	7.6	31
27	Insights into the evolution of bacterial flagellar motors from high-throughput <i>in situ</i> electron cryotomography and subtomogram averaging. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 585-594.	2.3	30
28	<i>ScpA</i> , a kinase involved in starvation-induced archaeellum expression. <i>Molecular Microbiology</i> , 2017, 103, 181-194.	2.5	29
29	Ultrastructure and complex polar architecture of the human pathogen <i>Campylobacter jejuni</i> . <i>MicrobiologyOpen</i> , 2014, 3, 702-710.	3.0	25
30	Bacterial flagellar motor PL-ring disassembly subcomplexes are widespread and ancient. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 8941-8947.	7.1	23
31	Diversification of <i>Campylobacter jejuni</i> Flagellar C-Ring Composition Impacts Its Structure and Function in Motility, Flagellar Assembly, and Cellular Processes. <i>MBio</i> , 2020, 11, .	4.1	23
32	<i>In situ</i> structure of the <i>Caulobacter crescentus</i> flagellar motor and visualization of binding of a CheY homolog. <i>Molecular Microbiology</i> , 2020, 114, 443-453.	2.5	22
33	The Brownian and Flow-Driven Rotational Dynamics of a Multicomponent DNA Origami-Based Rotor. <i>Small</i> , 2020, 16, e2001855.	10.0	20
34	An ATP-dependent partner switch links flagellar C-ring assembly with gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 20826-20835.	7.1	17
35	Adaptation of the periplasm to maintain spatial constraints essential for cell envelope processes and cell viability. <i>ELife</i> , 2022, 11, .	6.0	17
36	<i>In situ</i> imaging of bacterial outer membrane projections and associated protein complexes using electron cryo-tomography. <i>ELife</i> , 2021, 10, .	6.0	16

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37	Molecular Models for the Core Components of the Flagellar Type-III Secretion Complex. PLoS ONE, 2016, 11, e0164047.	2.5	14
38	Electron Cryotomography of Bacterial Cells. Journal of Visualized Experiments, 2010, , .	0.3	13
39	Bacterial Flagellins: Does Size Matter?. Trends in Microbiology, 2018, 26, 575-581.	7.7	13
40	Structure of the cytoplasmic domain of SctV (SsaV) from the Salmonella SPI-2 injectisome and implications for a pH sensing mechanism. Journal of Structural Biology, 2021, 213, 107729.	2.8	13
41	Lpp positions peptidoglycan at the AcrA-TolC interface in the AcrAB-TolC multidrug efflux pump. Biophysical Journal, 2021, 120, 3973-3982.	0.5	13
42	Analysis of Cell-Cell Bridges in <i>Haloferax volcanii</i> Using Electron Cryo-Tomography Reveal a Continuous Cytoplasm and S-Layer. Frontiers in Microbiology, 2020, 11, 612239.	3.5	13
43	Proteins in a Small World. Science, 2006, 314, 1882-1883.	12.6	12
44	Simulations suggest a constrictive force is required for Gram-negative bacterial cell division. Nature Communications, 2019, 10, 1259.	12.8	12
45	Giant flagellins form thick flagellar filaments in two species of marine β -proteobacteria. PLoS ONE, 2018, 13, e0206544.	2.5	10
46	<i>Trichinella spiralis</i> secretes abundant unencapsulated small RNAs with potential effects on host gene expression. International Journal for Parasitology, 2020, 50, 697-705.	3.1	10
47	CryoEM of bacterial secretion systems: A primer for microbiologists. Molecular Microbiology, 2021, 115, 366-382.	2.5	10
48	Novel transient cytoplasmic rings stabilize assembling bacterial flagellar motors. EMBO Journal, 2022, 41, e109523.	7.8	10
49	Advances in Mycobacterium tuberculosis Structural Genomics: Investigating Potential Chinks in the Armor of a Deadly Pathogen. Infectious Disorders - Drug Targets, 2009, 9, 475-492.	0.8	8
50	The "Jack-of-all-Trades" Flagellum From Salmonella and E. coli Was Horizontally Acquired From an Ancestral β -Proteobacterium. Frontiers in Microbiology, 2021, 12, 643180.	3.5	6
51	Loss of the Bacterial Flagellar Motor Switch Complex upon Cell Lysis. MBio, 2021, 12, e0029821.	4.1	6
52	Evolution of a family of molecular Rube Goldberg contraptions. PLoS Biology, 2019, 17, e3000405.	5.6	5
53	High-Throughput Electron Cryo-tomography of Protein Complexes and Their Assembly. Methods in Molecular Biology, 2018, 1764, 29-44.	0.9	4
54	Toward Organism-scale Structural Biology: S-layer Reined in by Bacterial LPS. Trends in Biochemical Sciences, 2020, 45, 549-551.	7.5	3

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55	Evolution of Archaellum Rotation Involved Invention of a Stator Complex by Duplicating and Modifying a Core Component. <i>Frontiers in Microbiology</i> , 2021, 12, 773386.	3.5	3
56	How Did the Archaellum Get Its Rotation?. <i>Frontiers in Microbiology</i> , 2021, 12, 803720.	3.5	3
57	Electron Cryo-Tomography. <i>Biological and Medical Physics Series</i> , 2018, , 61-94.	0.4	2
58	Title is missing!. , 2020, 16, e1008620.		0
59	Title is missing!. , 2020, 16, e1008620.		0
60	Title is missing!. , 2020, 16, e1008620.		0
61	Title is missing!. , 2020, 16, e1008620.		0
62	Title is missing!. , 2020, 16, e1008620.		0
63	Title is missing!. , 2020, 16, e1008620.		0