

David Z Rudner

List of Publications by Year in descending order

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69
papers

6,633
citations

87888

38
h-index

95266

68
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78
all docs

78
docs citations

78
times ranked

4860
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic Evidence for Signal Transduction within the <i>Bacillus subtilis</i> GerA Germinant Receptor. <i>Journal of Bacteriology</i> , 2022, 204, JB0047021.	2.2	11
2	The WalR-Walk Signaling Pathway Modulates the Activities of both CwlO and LytE through Control of the Peptidoglycan Deacetylase PdaC in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2022, 204, JB0053321.	2.2	11
3	WhyD tailors surface polymers to prevent premature bacteriolysis and direct cell elongation in <i>Streptococcus pneumoniae</i> . <i>ELife</i> , 2022, 11, .	6.0	3
4	The SpoVA membrane complex is required for dipicolinic acid import during sporulation and export during germination. <i>Genes and Development</i> , 2022, 36, 634-646.	5.9	17
5	Chromosome Segregation and Peptidoglycan Remodeling Are Coordinated at a Highly Stabilized Septal Pore to Maintain Bacterial Spore Development. <i>Developmental Cell</i> , 2021, 56, 36-51.e5.	7.0	13
6	Respiratory chain components are required for peptidoglycan recognition protein-induced thiol depletion and killing in <i>Bacillus subtilis</i> and <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2021, 11, 64.	3.3	3
7	XerD unloads bacterial SMC complexes at the replication terminus. <i>Molecular Cell</i> , 2021, 81, 756-766.e8.	9.7	27
8	FisB relies on homo-oligomerization and lipid binding to catalyze membrane fission in bacteria. <i>PLoS Biology</i> , 2021, 19, e3001314.	5.6	9
9	Dormant spores sense amino acids through the B subunits of their germination receptors. <i>Nature Communications</i> , 2021, 12, 6842.	12.8	22
10	SwsB and SafA Are Required for CwlJ-Dependent Spore Germination in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2020, 202, .	2.2	10
11	Barcoded microbial system for high-resolution object provenance. <i>Science</i> , 2020, 368, 1135-1140.	12.6	27
12	Structural coordination of polymerization and crosslinking by a SEDS- α -bBPB peptidoglycan synthase complex. <i>Nature Microbiology</i> , 2020, 5, 813-820.	13.3	91
13	A dynamic, ring-forming MucB / RseB-like protein influences spore shape in <i>Bacillus subtilis</i> . <i>PLoS Genetics</i> , 2020, 16, e1009246.	3.5	5
14	SweC and SweD are essential co-factors of the FtsEX-CwlO cell wall hydrolase complex in <i>Bacillus subtilis</i> . <i>PLoS Genetics</i> , 2019, 15, e1008296.	3.5	37
15	RNA polymerases as moving barriers to condensin loop extrusion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20489-20499.	7.1	105
16	A switch in surface polymer biogenesis triggers growth-phase-dependent and antibiotic-induced bacteriolysis. <i>ELife</i> , 2019, 8, .	6.0	47
17	Homeostatic control of cell wall hydrolysis by the WalRK two-component signaling pathway in <i>Bacillus subtilis</i> . <i>ELife</i> , 2019, 8, .	6.0	52
18	Phosphorylation-dependent activation of the cell wall synthase PBP2a in <i>Streptococcus pneumoniae</i> by MacP. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 2812-2817.	7.1	62

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19	Structure of the peptidoglycan polymerase RodA resolved by evolutionary coupling analysis. <i>Nature</i> , 2018, 556, 118-121.	27.8	110
20	Evidence that regulation of intramembrane proteolysis is mediated by substrate gating during sporulation in <i>Bacillus subtilis</i> . <i>PLoS Genetics</i> , 2018, 14, e1007753.	3.5	11
21	Structural characterization of the sporulation protein GerM from <i>Bacillus subtilis</i> . <i>Journal of Structural Biology</i> , 2018, 204, 481-490.	2.8	8
22	InÂVivo Evidence for ATPase-Dependent DNA Translocation by the <i>Bacillus subtilis</i> SMC Condensin Complex. <i>Molecular Cell</i> , 2018, 71, 841-847.e5.	9.7	66
23	<i>Bacillus subtilis</i> SMC complexes juxtapose chromosome arms as they travel from origin to terminus. <i>Science</i> , 2017, 355, 524-527.	12.6	267
24	Construction and Analysis of Two Genome-Scale Deletion Libraries for <i>Bacillus subtilis</i> . <i>Cell Systems</i> , 2017, 4, 291-305.e7.	6.2	457
25	The <i>Bacillus subtilis</i> germinant receptor GerA triggers premature germination in response to morphological defects during sporulation. <i>Molecular Microbiology</i> , 2017, 105, 689-704.	2.5	23
26	CozE is a member of the MreCD complex that directs cell elongation in <i>Streptococcus pneumoniae</i> . <i>Nature Microbiology</i> , 2017, 2, 16237.	13.3	70
27	The nucleoid occlusion factor Noc controls DNA replication initiation in <i>Staphylococcus aureus</i> . <i>PLoS Genetics</i> , 2017, 13, e1006908.	3.5	43
28	A two-step transport pathway allows the mother cell to nurture the developing spore in <i>Bacillus subtilis</i> . <i>PLoS Genetics</i> , 2017, 13, e1007015.	3.5	32
29	A ring-shaped conduit connects the mother cell and forespore during sporulation in <i>Bacillus subtilis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 11585-11590.	7.1	24
30	SEDS proteins are a widespread family of bacterial cell wall polymerases. <i>Nature</i> , 2016, 537, 634-638.	27.8	448
31	GerM is required to assemble the basal platform of the SpoIIAâ€“SpoIIQ transenvelope complex during sporulation in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2016, 102, 260-273.	2.5	27
32	Saltâ€“sensitivity of σ^H and Spo0A prevents sporulation of <i>Bacillus subtilis</i> at high osmolarity avoiding death during cellular differentiation. <i>Molecular Microbiology</i> , 2016, 100, 108-124.	2.5	25
33	High-Throughput Genetic Screens Identify a Large and Diverse Collection of New Sporulation Genes in <i>Bacillus subtilis</i> . <i>PLoS Biology</i> , 2016, 14, e1002341.	5.6	87
34	An experimentally supported model of the <i>Bacillus subtilis</i> global transcriptional regulatory network. <i>Molecular Systems Biology</i> , 2015, 11, 839.	7.2	186
35	Condensin promotes the juxtaposition of DNA flanking its loading site in <i>Bacillus subtilis</i> . <i>Genes and Development</i> , 2015, 29, 1661-1675.	5.9	215
36	MurJ and a novel lipid II flippase are required for cell wall biogenesis in <i>Bacillus subtilis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 6437-6442.	7.1	166

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37	Condensation and localization of the partitioning protein ParB on the bacterial chromosome. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 8809-8814.	7.1	96
38	Spatial organization of bacterial chromosomes. Current Opinion in Microbiology, 2014, 22, 66-72.	5.1	51
39	<i>Bacillus subtilis</i> chromosome organization oscillates between two distinct patterns. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 12877-12882.	7.1	116
40	ParB spreading requires DNA bridging. Genes and Development, 2014, 28, 1228-1238.	5.9	177
41	The SMC Condensin Complex Is Required for Origin Segregation in <i>Bacillus subtilis</i> . Current Biology, 2014, 24, 287-292.	3.9	109
42	<i>FtsEX</i> is required for <i>CwlO</i> peptidoglycan hydrolase activity during cell wall elongation in <i>Bacillus subtilis</i> . Molecular Microbiology, 2013, 89, 1069-1083.	2.5	145
43	Organization and segregation of bacterial chromosomes. Nature Reviews Genetics, 2013, 14, 191-203.	16.3	252
44	CtpB Assembles a Gated Protease Tunnel Regulating Cell-Cell Signaling during Spore Formation in <i>Bacillus subtilis</i> . Cell, 2013, 155, 647-658.	28.9	31
45	FisB mediates membrane fission during sporulation in <i>Bacillus subtilis</i> . Genes and Development, 2013, 27, 322-334.	5.9	47
46	Peptidoglycan hydrolysis is required for assembly and activity of the transenvelope secretion complex during sporulation in <i>Bacillus subtilis</i> . Molecular Microbiology, 2013, 89, 1039-1052.	2.5	28
47	RefZ Facilitates the Switch from Medial to Polar Division during Spore Formation in <i>Bacillus subtilis</i> . Journal of Bacteriology, 2012, 194, 4608-4618.	2.2	23
48	Coupled, Circumferential Motions of the Cell Wall Synthesis Machinery and MreB Filaments in <i>B. subtilis</i> . Science, 2011, 333, 222-225.	12.6	505
49	Nucleoid occlusion prevents cell division during replication fork arrest in <i>Bacillus subtilis</i> . Molecular Microbiology, 2010, 78, 866-882.	2.5	47
50	A highly coordinated cell wall degradation machine governs spore morphogenesis in <i>Bacillus subtilis</i> . Genes and Development, 2010, 24, 411-422.	5.9	91
51	Protein Subcellular Localization in Bacteria. Cold Spring Harbor Perspectives in Biology, 2010, 2, a000307-a000307.	5.5	163
52	Novel Secretion Apparatus Maintains Spore Integrity and Developmental Gene Expression in <i>Bacillus subtilis</i> . PLoS Genetics, 2009, 5, e1000566.	3.5	93
53	SirA enforces diploidy by inhibiting the replication initiator DnaA during spore formation in <i>Bacillus subtilis</i> . Molecular Microbiology, 2009, 73, 963-974.	2.5	72
54	Recruitment of SMC by ParB-parS Organizes the Origin Region and Promotes Efficient Chromosome Segregation. Cell, 2009, 137, 697-707.	28.9	275

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55	SpoIIQ Anchors Membrane Proteins on Both Sides of the Sporulation Septum in <i>Bacillus subtilis</i> . <i>Journal of Biological Chemistry</i> , 2008, 283, 4975-4982.	3.4	34
56	SpoIIIE strips proteins off the DNA during chromosome translocation. <i>Genes and Development</i> , 2008, 22, 1786-1795.	5.9	63
57	SpoIVB and CtpB Are Both Forespore Signals in the Activation of the Sporulation Transcription Factor σ^K in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2007, 189, 6021-6027.	2.2	37
58	The ATPase SpoIIIE Transports DNA across Fused Septal Membranes during Sporulation in <i>Bacillus subtilis</i> . <i>Cell</i> , 2007, 131, 1301-1312.	28.9	112
59	Perturbations to engulfment trigger a degradative response that prevents cell-cell signalling during sporulation in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2007, 64, 500-511.	2.5	21
60	A Branched Pathway Governing the Activation of a Developmental Transcription Factor by Regulated Intramembrane Proteolysis. <i>Molecular Cell</i> , 2006, 23, 25-35.	9.7	63
61	Subcellular localization of a sporulation membrane protein is achieved through a network of interactions along and across the septum. <i>Molecular Microbiology</i> , 2005, 55, 1767-1781.	2.5	109
62	Defining a Centromere-like Element in <i>Bacillus subtilis</i> by Identifying the Binding Sites for the Chromosome-Anchoring Protein RacA. <i>Molecular Cell</i> , 2005, 17, 773-782.	9.7	93
63	The Program of Gene Transcription for a Single Differentiating Cell Type during Sporulation in <i>Bacillus subtilis</i> . <i>PLoS Biology</i> , 2004, 2, e328.	5.6	308
64	RacA, a Bacterial Protein That Anchors Chromosomes to the Cell Poles. <i>Science</i> , 2003, 299, 532-536.	12.6	287
65	A Second PDZ-Containing Serine Protease Contributes to Activation of the Sporulation Transcription Factor σ^K in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2003, 185, 6051-6056.	2.2	33
66	A sporulation membrane protein tethers the pro- σ^K processing enzyme to its inhibitor and dictates its subcellular localization. <i>Genes and Development</i> , 2002, 16, 1007-1018.	5.9	115
67	Evidence that subcellular localization of a bacterial membrane protein is achieved by diffusion and capture. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 8701-8706.	7.1	122
68	Morphological Coupling in Development. <i>Developmental Cell</i> , 2001, 1, 733-742.	7.0	89
69	Intercompartmental Signal Transduction during Sporulation in <i>Bacillus subtilis</i> . , 0, , 1-12.		0