

Thomas J Silhavy

List of Publications by Year in descending order

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249
papers

21,679
citations

7568

77
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11052

137
g-index

257
all docs

257
docs citations

257
times ranked

12931
citing authors

#	ARTICLE	IF	CITATIONS
1	The Bacterial Cell Envelope. Cold Spring Harbor Perspectives in Biology, 2010, 2, a000414-a000414.	5.5	2,408
2	Identification of a Multicomponent Complex Required for Outer Membrane Biogenesis in Escherichia coli. Cell, 2005, 121, 235-245.	28.9	656
3	Suppressor mutations that restore export of a protein with a defective signal sequence. Cell, 1981, 23, 79-88.	28.9	435
4	An ABC transport system that maintains lipid asymmetry in the Gram-negative outer membrane. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 8009-8014.	7.1	411
5	Defining the roles of the periplasmic chaperones SurA, Skp, and DegP in <i>Escherichia coli</i> . Genes and Development, 2007, 21, 2473-2484.	5.9	409
6	Advances in understanding bacterial outer-membrane biogenesis. Nature Reviews Microbiology, 2006, 4, 57-66.	28.6	405
7	Surface sensing and adhesion of Escherichia coli controlled by the Cpx-signaling pathway. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 2287-2292.	7.1	368
8	The ompB locus and the regulation of the major outer membrane porin proteins of Escherichia coli K12. Journal of Molecular Biology, 1981, 146, 23-43.	4.2	358
9	Periplasmic Stress and ECF Sigma Factors. Annual Review of Microbiology, 2001, 55, 591-624.	7.3	342
10	Genetic analysis of the ompB locus in Escherichia coli K-12. Journal of Molecular Biology, 1981, 151, 1-15.	4.2	341
11	Structure and Function of an Essential Component of the Outer Membrane Protein Assembly Machine. Science, 2007, 317, 961-964.	12.6	327
12	Identification of a protein complex that assembles lipopolysaccharide in the outer membrane of Escherichia coli. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 11754-11759.	7.1	322
13	Lipopolysaccharide transport and assembly at the outer membrane: the PEZ model. Nature Reviews Microbiology, 2016, 14, 337-345.	28.6	299
14	From acids to <i>osmZ</i> : multiple factors influence synthesis of the OmpF and OmpC porins in <i>Escherichia coli</i> . Molecular Microbiology, 1996, 20, 911-917.	2.5	298
15	β -Barrel Membrane Protein Assembly by the Bam Complex. Annual Review of Biochemistry, 2011, 80, 189-210.	11.1	290
16	Chemical Conditionality. Cell, 2005, 121, 307-317.	28.9	287
17	The E. coli ffh gene is necessary for viability and efficient protein export. Nature, 1992, 359, 744-746.	27.8	285
18	Sensing external stress: watchdogs of the Escherichia coli cell envelope. Current Opinion in Microbiology, 2005, 8, 122-126.	5.1	281

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19	YfiO stabilizes the YaeT complex and is essential for outer membrane protein assembly in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2006, 61, 151-164.	2.5	278
20	Lipoprotein SmpA is a component of the YaeT complex that assembles outer membrane proteins in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 6400-6405.	7.1	267
21	CpxP, a Stress-Combative Member of the Cpx Regulon. <i>Journal of Bacteriology</i> , 1998, 180, 831-839.	2.2	265
22	Imp/OstA is required for cell envelope biogenesis in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2002, 45, 1289-1302.	2.5	232
23	Transport of lipopolysaccharide across the cell envelope: the long road of discovery. <i>Nature Reviews Microbiology</i> , 2009, 7, 677-683.	28.6	232
24	Outer Membrane Biogenesis. <i>Annual Review of Microbiology</i> , 2017, 71, 539-556.	7.3	229
25	Identification of two inner-membrane proteins required for the transport of lipopolysaccharide to the outer membrane of <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 5537-5542.	7.1	225
26	Sequence analysis of mutations that prevent export of β receptor, an <i>Escherichia coli</i> outer membrane protein. <i>Nature</i> , 1980, 285, 82-85.	27.8	224
27	Genetic Evidence for Parallel Pathways of Chaperone Activity in the Periplasm of <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2001, 183, 6794-6800.	2.2	219
28	Functional Analysis of the Protein Machinery Required for Transport of Lipopolysaccharide to the Outer Membrane of <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2008, 190, 4460-4469.	2.2	218
29	The Cpx Envelope Stress Response Is Controlled by Amplification and Feedback Inhibition. <i>Journal of Bacteriology</i> , 1999, 181, 5263-5272.	2.2	209
30	TARGETING AND ASSEMBLY OF PERIPLASMIC AND OUTER-MEMBRANE PROTEINS IN <i>ESCHERICHIA COLI</i> . <i>Annual Review of Genetics</i> , 1998, 32, 59-94.	7.6	206
31	Signal Detection and Target Gene Induction by the CpxRA Two-Component System. <i>Journal of Bacteriology</i> , 2003, 185, 2432-2440.	2.2	198
32	Heat-shock proteins DnaK and GroEL facilitate export of LacZ hybrid proteins in <i>E. coli</i> . <i>Nature</i> , 1990, 344, 882-884.	27.8	195
33	EnvZ controls the concentration of phosphorylated OmpR to mediate osmoregulation of the porin genes. <i>Journal of Molecular Biology</i> , 1991, 222, 567-580.	4.2	194
34	These <i>cndprl</i> genes of <i>Escherichia coli</i> . <i>Journal of Bioenergetics and Biomembranes</i> , 1990, 22, 291-310.	2.3	190
35	Characterization of the two-protein complex in <i>Escherichia coli</i> responsible for lipopolysaccharide assembly at the outer membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 5363-5368.	7.1	184
36	Contact-dependent growth inhibition requires the essential outer membrane protein BamA (YaeT) as the receptor and the inner membrane transport protein AcrB. <i>Molecular Microbiology</i> , 2008, 70, 323-340.	2.5	173

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37	The σ^E and Cpx regulatory pathways: Overlapping but distinct envelope stress responses. <i>Current Opinion in Microbiology</i> , 1999, 2, 159-165.	5.1	167
38	Mutations affecting localization of an Escherichia coli outer membrane protein, the bacteriophage λ receptor. <i>Journal of Molecular Biology</i> , 1980, 141, 63-90.	4.2	166
39	A signal sequence is not sufficient to lead β -galactosidase out of the cytoplasm. <i>Nature</i> , 1980, 286, 356-359.	27.8	165
40	Two-Component Signal Transduction Systems: Structure-Function Relationships and Mechanisms of Catalysis. <i>Journal of Molecular Biology</i> , 1990, 215, 25-51.		164
41	Structure of the malB region in Escherichia coli K12. <i>Molecular Genetics and Genomics</i> , 1979, 174, 249-259.	2.4	157
42	Envelope stress responses: balancing damage repair and toxicity. <i>Nature Reviews Microbiology</i> , 2019, 17, 417-428.	28.6	153
43	The Bam machine: A molecular cooper. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 1067-1084.	2.6	145
44	PrIA (SecY) and PrIC (SecE) interact directly and function sequentially during protein translocation in E. coli. <i>Cell</i> , 1990, 61, 833-842.	28.9	143
45	Quality control in the bacterial periplasm. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2004, 1694, 121-134.	4.1	143
46	A previously unidentified gene in the spc operon of Escherichia coli K12 specifies a component of the protein export machinery. <i>Cell</i> , 1982, 31, 227-235.	28.9	142
47	The extracytoplasmic adaptor protein CpxP is degraded with substrate by DegP. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 17775-17779.	7.1	142
48	Disruption of lipid homeostasis in the Gram-negative cell envelope activates a novel cell death pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E1565-74.	7.1	142
49	Mutations That Alter the Kinase and Phosphatase Activities of the Two-Component Sensor EnvZ. <i>Journal of Bacteriology</i> , 1998, 180, 4538-4546.	2.2	141
50	A small-molecule inhibitor of BamA impervious to efflux and the outer membrane permeability barrier. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 21748-21757.	7.1	136
51	Characterization of the role of the <i>Escherichia coli</i> periplasmic chaperone SurA using differential proteomics. <i>Proteomics</i> , 2009, 9, 2432-2443.	2.2	128
52	Genetic Basis for Activity Differences Between Vancomycin and Glycolipid Derivatives of Vancomycin. <i>Science</i> , 2001, 294, 361-364.	12.6	127
53	Periplasmic Peptidyl Prolyl cis-trans Isomerases Are Not Essential for Viability, but SurA Is Required for Pilus Biogenesis in Escherichia coli. <i>Journal of Bacteriology</i> , 2005, 187, 7680-7686.	2.2	126
54	Genetic analysis of the switch that controls porin gene expression in Escherichia coli K-12. <i>Journal of Molecular Biology</i> , 1989, 210, 281-292.	4.2	123

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55	Sirtuins Are Evolutionarily Conserved Viral Restriction Factors. <i>MBio</i> , 2014, 5, .	4.1	122
56	Lipoprotein LptE is required for the assembly of LptD by the β^2 -barrel assembly machine in the outer membrane of <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2492-2497.	7.1	116
57	Outer membrane lipoprotein biogenesis: Lol is not the end. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20150030.	4.0	116
58	Crl stimulates RpoS activity during stationary phase. <i>Molecular Microbiology</i> , 1998, 29, 1225-1236.	2.5	114
59	Envelope Stress Responses: An Interconnected Safety Net. <i>Trends in Biochemical Sciences</i> , 2017, 42, 232-242.	7.5	112
60	The essential tension: opposed reactions in bacterial two-component regulatory systems. <i>Trends in Microbiology</i> , 1993, 1, 306-310.	7.7	111
61	Physical properties of the bacterial outer membrane. <i>Nature Reviews Microbiology</i> , 2022, 20, 236-248.	28.6	111
62	The free and bound forms of Lpp occupy distinct subcellular locations in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2011, 79, 1168-1181.	2.5	109
63	Transmembrane domain of surface-exposed outer membrane lipoprotein RcsF is threaded through the lumen of β^2 -barrel proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E4350-8.	7.1	109
64	<i>Escherichia coli</i> Starvation Diets: Essential Nutrients Weigh in Distinctly. <i>Journal of Bacteriology</i> , 2005, 187, 7549-7553.	2.2	107
65	Effects of Antibiotics and a Proto-Oncogene Homolog on Destruction of Protein Translocator SecY. <i>Science</i> , 2009, 325, 753-756.	12.6	105
66	The Cpx Stress Response Confers Resistance to Some, but Not All, Bactericidal Antibiotics. <i>Journal of Bacteriology</i> , 2013, 195, 1869-1874.	2.2	103
67	Redefining the essential trafficking pathway for outer membrane lipoproteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 4769-4774.	7.1	101
68	Mutational activation of the Cpx signal transduction pathway of <i>Escherichia coli</i> suppresses the toxicity conferred by certain envelope-associated stresses. <i>Molecular Microbiology</i> , 1995, 18, 491-505.	2.5	98
69	Nonconsecutive disulfide bond formation in an essential integral outer membrane protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 12245-12250.	7.1	96
70	Mapping an Interface of SecY (PrIA) and SecE (PrIG) by Using Synthetic Phenotypes and In Vivo Cross-Linking. <i>Journal of Bacteriology</i> , 1999, 181, 3438-3444.	2.2	96
71	Sequence information within the lamB gene is required for proper routing of the bacteriophage λ receptor protein to the outer membrane of <i>Escherichia coli</i> K-12. <i>Journal of Molecular Biology</i> , 1982, 156, 93-112.	4.2	94
72	Conformation-specific labeling of BamA and suppressor analysis suggest a cyclic mechanism for β^2 -barrel assembly in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 5151-5156.	7.1	94

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73	Tethering of CpxP to the inner membrane prevents spheroplast induction of the Cpx envelope stress response. <i>Molecular Microbiology</i> , 2000, 37, 1186-1197.	2.5	91
74	Accumulation of the Enterobacterial Common Antigen Lipid II Biosynthetic Intermediate Stimulates σ^{pP} Transcription in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 1998, 180, 5875-5884.	2.2	90
75	A lipoprotein/ β -barrel complex monitors lipopolysaccharide integrity transducing information across the outer membrane. <i>ELife</i> , 2016, 5, .	6.0	88
76	prfF and yhaV Encode a New Toxin-Antitoxin System in <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 2007, 372, 894-905.	4.2	87
77	Starvation for Different Nutrients in <i>Escherichia coli</i> Results in Differential Modulation of RpoS Levels and Stability. <i>Journal of Bacteriology</i> , 2005, 187, 434-442.	2.2	85
78	The art and design of genetic screens: <i>Escherichia coli</i> . <i>Nature Reviews Genetics</i> , 2003, 4, 419-431.	16.3	84
79	Kinetic Analysis of the Assembly of the Outer Membrane Protein LamB in <i>Escherichia coli</i> Mutants Each Lacking a Secretion or Targeting Factor in a Different Cellular Compartment. <i>Journal of Bacteriology</i> , 2007, 189, 446-454.	2.2	83
80	Complex spatial distribution and dynamics of an abundant <i>Escherichia coli</i> outer membrane protein, LamB. <i>Molecular Microbiology</i> , 2004, 53, 1771-1783.	2.5	82
81	Information within the mature LamB protein necessary for localization to the outer membrane of <i>E. coli</i> K12. <i>Cell</i> , 1983, 32, 1325-1335.	28.9	80
82	Mutations that Affect Separate Functions of OmpR the Phosphorylated Regulator of Porin Transcription in <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 1993, 231, 261-273.	4.2	80
83	Characterization of a stalled complex on the β -barrel assembly machine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 8717-8722.	7.1	77
84	Activation of the <i>Escherichia coli</i> β -barrel assembly machine (Bam) is required for essential components to interact properly with substrate. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 3487-3491.	7.1	76
85	LptE binds to and alters the physical state of LPS to catalyze its assembly at the cell surface. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9467-9472.	7.1	74
86	Probing the Barrier Function of the Outer Membrane with Chemical Conditionality. <i>ACS Chemical Biology</i> , 2006, 1, 385-395.	3.4	72
87	BamE Modulates the <i>Escherichia coli</i> Beta-Barrel Assembly Machine Component BamA. <i>Journal of Bacteriology</i> , 2012, 194, 1002-1008.	2.2	72
88	Porin Regulon of <i>Escherichia coli</i> . , 0, , 105-127.		70
89	Isolation and characterization of mutations altering expression of the major outer membrane porin proteins using the local anaesthetic procaine. <i>Journal of Molecular Biology</i> , 1983, 166, 273-282.	4.2	69
90	A Suppressor of Cell Death Caused by the Loss of σ^E Downregulates Extracytoplasmic Stress Responses and Outer Membrane Vesicle Production in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2007, 189, 1523-1530.	2.2	68

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91	Making a beta-barrel: assembly of outer membrane proteins in Gram-negative bacteria. <i>Current Opinion in Microbiology</i> , 2012, 15, 189-193.	5.1	67
92	OmpR mutants specifically defective for transcriptional activation. <i>Journal of Molecular Biology</i> , 1994, 243, 579-594.	4.2	65
93	The <i>Escherichia coli</i> Phospholipase PldA Regulates Outer Membrane Homeostasis via Lipid Signaling. <i>MBio</i> , 2018, 9, .	4.1	65
94	The LysR Homolog LrhA Promotes RpoS Degradation by Modulating Activity of the Response Regulator SprE. <i>Journal of Bacteriology</i> , 1999, 181, 563-571.	2.2	65
95	[9] Genetic fusions as experimental tools. <i>Methods in Enzymology</i> , 1991, 204, 213-248.	1.0	64
96	Secretion of LamB-LacZ by the Signal Recognition Particle Pathway of <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2003, 185, 5697-5705.	2.2	64
97	[15] Engineering <i>Escherichia coli</i> to secrete heterologous gene products. <i>Methods in Enzymology</i> , 1990, 185, 166-187.	1.0	63
98	Constitutive Activation of the <i>Escherichia coli</i> Pho Regulon Upregulates rpoS Translation in an Hfq-Dependent Fashion. <i>Journal of Bacteriology</i> , 2003, 185, 5984-5992.	2.2	60
99	Dissecting the <i>Escherichia coli</i> periplasmic chaperone network using differential proteomics. <i>Proteomics</i> , 2012, 12, 1391-1401.	2.2	58
100	Identification of base pairs important for OmpR-DNA interaction. <i>Molecular Microbiology</i> , 1995, 17, 565-573.	2.5	57
101	RpoS Proteolysis Is Regulated by a Mechanism That Does Not Require the SprE (RssB) Response Regulator Phosphorylation Site. <i>Journal of Bacteriology</i> , 2004, 186, 7403-7410.	2.2	56
102	Making a membrane on the other side of the wall. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2017, 1862, 1386-1393.	2.4	55
103	Cyclic Enterobacterial Common Antigen Maintains the Outer Membrane Permeability Barrier of <i>Escherichia coli</i> in a Manner Controlled by YhdP. <i>MBio</i> , 2018, 9, .	4.1	54
104	Phase separation in the outer membrane of <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	53
105	LrhA Regulates rpoS Translation in Response to the Rcs Phosphorelay System in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2006, 188, 3175-3181.	2.2	52
106	Decline in ribosomal fidelity contributes to the accumulation and stabilization of the master stress response regulator σ^S upon carbon starvation. <i>Genes and Development</i> , 2007, 21, 862-874.	5.9	52
107	RpoS proteolysis is controlled directly by ATP levels in <i>Escherichia coli</i> . <i>Genes and Development</i> , 2012, 26, 548-553.	5.9	52
108	The CpxQ sRNA Negatively Regulates Skp To Prevent Mistargeting of β -Barrel Outer Membrane Proteins into the Cytoplasmic Membrane. <i>MBio</i> , 2016, 7, e00312-16.	4.1	52

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109	Inhibitor of intramembrane protease RseP blocks the σ^E response causing lethal accumulation of unfolded outer membrane proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E6614-E6621.	7.1	51
110	Characterization and <i>in Vivo</i> Cloning of <i>prlC</i> , a Suppressor of Signal Sequence Mutations in <i>Escherichia coli</i> K12. <i>Genetics</i> , 1987, 116, 513-521.	2.9	50
111	Regulation of Capsule Synthesis: Modification of the Two-Component Paradigm by an Accessory Unstable Regulator. , 0, , 253-262.		49
112	YejM Modulates Activity of the YciM/FtsH Protease Complex To Prevent Lethal Accumulation of Lipopolysaccharide. <i>MBio</i> , 2020, 11, .	4.1	48
113	Substrate binding to BamD triggers a conformational change in BamA to control membrane insertion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 2359-2364.	7.1	47
114	The Response Regulator SprE (RssB) Is Required for Maintaining Poly(A) Polymerase I-Degradosome Association during Stationary Phase. <i>Journal of Bacteriology</i> , 2010, 192, 3713-3721.	2.2	46
115	Crl Facilitates RNA Polymerase Holoenzyme Formation. <i>Journal of Bacteriology</i> , 2006, 188, 7966-7970.	2.2	45
116	Control of Cellular Development in Sporulating Bacteria by the Phosphorelay Two-Component Signal Transduction System. , 0, , 129-144.		45
117	Predicting Functionally Informative Mutations in <i>Escherichia coli</i> BamA Using Evolutionary Covariance Analysis. <i>Genetics</i> , 2013, 195, 443-455.	2.9	42
118	Transcriptional occlusion caused by overlapping promoters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 1557-1561.	7.1	41
119	RpoS-Dependent Transcriptional Control of sprE : Regulatory Feedback Loop. <i>Journal of Bacteriology</i> , 2001, 183, 5974-5981.	2.2	40
120	Role for Skp in LptD Assembly in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2013, 195, 3734-3742.	2.2	40
121	Novel RpoS-Dependent Mechanisms Strengthen the Envelope Permeability Barrier during Stationary Phase. <i>Journal of Bacteriology</i> , 2017, 199, .	2.2	40
122	Border Control: Regulating LPS Biogenesis. <i>Trends in Microbiology</i> , 2021, 29, 334-345.	7.7	40
123	Dual Sensors and Dual Response Regulators Interact to Control Nitrate- and Nitrite-Responsive Gene Expression in <i>Escherichia coli</i> . , 0, , 233-252.		40
124	Continuous Control in Bacterial Regulatory Circuits. <i>Journal of Bacteriology</i> , 2004, 186, 7618-7625.	2.2	39
125	Classifying β -Barrel Assembly Substrates by Manipulating Essential Bam Complex Members. <i>Journal of Bacteriology</i> , 2016, 198, 1984-1992.	2.2	38
126	Distinctive Roles for Periplasmic Proteases in the Maintenance of Essential Outer Membrane Protein Assembly. <i>Journal of Bacteriology</i> , 2017, 199, .	2.2	37

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127	Accumulation of Phosphatidic Acid Increases Vancomycin Resistance in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2014, 196, 3214-3220.	2.2	36
128	The inner membrane protein YhdP modulates the rate of anterograde phospholipid flow in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 26907-26914.	7.1	36
129	Dominant Negative <i>lptE</i> Mutation That Supports a Role for LptE as a Plug in the LptD Barrel. <i>Journal of Bacteriology</i> , 2013, 195, 1327-1334.	2.2	35
130	The Synthetic Phenotype of $\hat{\Gamma}^*$ <i>bamB</i> / <i>l</i> $\hat{\Gamma}^*$ <i>bamE</i> / <i>l</i> Double Mutants Results from a Lethal Jamming of the Bam Complex by the Lipoprotein RcsF. <i>MBio</i> , 2019, 10, .	4.1	35
131	The Activity and Specificity of the Outer Membrane Protein Chaperone SurA Are Modulated by a Proline Isomerase Domain. <i>MBio</i> , 2013, 4, .	4.1	34
132	The genetics of protein secretion in <i>E. coli</i> . <i>Trends in Genetics</i> , 1990, 6, 329-334.	6.7	33
133	P Pilus Assembly Motif Necessary for Activation of the CpxRA Pathway by PapE in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2004, 186, 4326-4337.	2.2	33
134	Involvement of a tryptophan residue in the binding site of <i>Escherichia coli</i> galactose-binding protein. <i>Biochemistry</i> , 1974, 13, 993-999.	2.5	32
135	Ti Plasmid and Chromosomally Encoded Two-Component Systems Important in Plant Cell Transformation by <i>Agrobacterium</i> Species. , 0, , 367-385.		32
136	The Activity of <i>Escherichia coli</i> Chaperone SurA Is Regulated by Conformational Changes Involving a Parvulin Domain. <i>Journal of Bacteriology</i> , 2016, 198, 921-929.	2.2	29
137	Outer Membrane Protein Insertion by the $\hat{\Gamma}^2$ -barrel Assembly Machine. <i>EcoSal Plus</i> , 2019, 8, .	5.4	29
138	Sirtuin Lipoamidase Activity Is Conserved in Bacteria as a Regulator of Metabolic Enzyme Complexes. <i>MBio</i> , 2017, 8, .	4.1	28
139	PrlC, a suppressor of signal sequence mutations in <i>Escherichia coli</i> , can direct the insertion of the signal sequence into the membrane. <i>Journal of Molecular Biology</i> , 1989, 205, 665-676.	4.2	27
140	A Signal Transduction Network in <i>Bacillus subtilis</i> Includes the DegS/DegU and Comp/ComA Two-Component Systems. , 0, , 447-471.		27
141	Assembly of Outer Membrane $\hat{\Gamma}^2$ -Barrel Proteins: the Bam Complex. <i>EcoSal Plus</i> , 2011, 4, .	5.4	26
142	Regulation of <i>Salmonella</i> Virulence by Two-Component Regulatory Systems. , 0, , 319-332.		26
143	[1] Genetic analysis of protein export in <i>Escherichia coli</i> . <i>Methods in Enzymology</i> , 1983, 97, 3-11.	1.0	25
144	Transposition of $\hat{\Gamma}^*$ placMu is mediated by the A protein altered at its carboxy-terminal end. <i>Gene</i> , 1988, 71, 177-186.	2.2	25

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145	Genetic Approaches for Signaling Pathways and Proteins. , 2014, , 7-23.		25
146	Two-Component Signal Transduction and Its Role in the Expression of Bacterial Virulence Factors. , 0, , 303-317.		25
147	Structural and Functional Conservation in Response Regulators. , 0, , 53-64.		25
148	Control of Nitrogen Assimilation by the NRI-NRII Two-Component System of Enteric Bacteria. , 0, , 65-88.		25
149	Signal Transduction and Cross Regulation in the Escherichia coli Phosphate Regulon by PhoR, CreC, and Acetyl Phosphate. , 0, , 201-221.		24
150	Conferral of transposable properties to a chromosomal gene in Escherichia coli. Journal of Molecular Biology, 1980, 141, 235-248.	4.2	23
151	Metabolite turns master regulator. Nature, 2013, 500, 283-284.	27.8	23
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