# Thomas J Silhavy

### List of Publications by Citations

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202 18,214 77 130 g-index

257 20,504 9.8 7.3 L-index

ext. papers ext. citations

#	Paper	IF	Citations
202	The bacterial cell envelope. <i>Cold Spring Harbor Perspectives in Biology</i> , <b>2010</b> , 2, a000414	10.2	1674
201	Identification of a multicomponent complex required for outer membrane biogenesis in Escherichia coli. <i>Cell</i> , <b>2005</b> , 121, 235-45	56.2	565
200	Suppressor mutations that restore export of a protein with a defective signal sequence. <i>Cell</i> , <b>1981</b> , 23, 79-88	56.2	388
199	Advances in understanding bacterial outer-membrane biogenesis. <i>Nature Reviews Microbiology</i> , <b>2006</b> , 4, 57-66	22.2	353
198	Defining the roles of the periplasmic chaperones SurA, Skp, and DegP in Escherichia coli. <i>Genes and Development</i> , <b>2007</b> , 21, 2473-84	12.6	336
197	Surface sensing and adhesion of Escherichia coli controlled by the Cpx-signaling pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2002</b> , 99, 2287-92	11.5	320
196	Periplasmic stress and ECF sigma factors. <i>Annual Review of Microbiology</i> , <b>2001</b> , 55, 591-624	17.5	319
195	The ompB locus and the regulation of the major outer membrane porin proteins of Escherichia coli K12. <i>Journal of Molecular Biology</i> , <b>1981</b> , 146, 23-43	6.5	304
194	Structure and function of an essential component of the outer membrane protein assembly machine. <i>Science</i> , <b>2007</b> , 317, 961-4	33.3	302
193	Genetic analysis of the ompB locus in Escherichia coli K-12. <i>Journal of Molecular Biology</i> , <b>1981</b> , 151, 1-15	6.5	289
192	An ABC transport system that maintains lipid asymmetry in the gram-negative outer membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2009</b> , 106, 8009-14	11.5	287
191	Identification of a protein complex that assembles lipopolysaccharide in the outer membrane of Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2006</b> , 103, 11754-9	11.5	267
190	The E. coli ffh gene is necessary for viability and efficient protein export. <i>Nature</i> , <b>1992</b> , 359, 744-6	50.4	259
189	From acids to osmZ: multiple factors influence synthesis of the OmpF and OmpC porins in Escherichia coli. <i>Molecular Microbiology</i> , <b>1996</b> , 20, 911-7	4.1	258
188	Sensing external stress: watchdogs of the Escherichia coli cell envelope. <i>Current Opinion in Microbiology</i> , <b>2005</b> , 8, 122-6	7.9	257
187	EBarrel membrane protein assembly by the Bam complex. <i>Annual Review of Biochemistry</i> , <b>2011</b> , 80, 189-	 2 <u>1</u> 90,1	254
186	Lipoprotein SmpA is a component of the YaeT complex that assembles outer membrane proteins in Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2007</b> , 104, 6400-5	11.5	240

### (1980-2005)

185	Chemical conditionality: a genetic strategy to probe organelle assembly. <i>Cell</i> , <b>2005</b> , 121, 307-17	56.2	238
184	YfiO stabilizes the YaeT complex and is essential for outer membrane protein assembly in Escherichia coli. <i>Molecular Microbiology</i> , <b>2006</b> , 61, 151-64	4.1	234
183	CpxP, a stress-combative member of the Cpx regulon. <i>Journal of Bacteriology</i> , <b>1998</b> , 180, 831-9	3.5	220
182	Lipopolysaccharide transport and assembly at the outer membrane: the PEZ model. <i>Nature Reviews Microbiology</i> , <b>2016</b> , 14, 337-45	22.2	208
181	Transport of lipopolysaccharide across the cell envelope: the long road of discovery. <i>Nature Reviews Microbiology</i> , <b>2009</b> , 7, 677-83	22.2	205
180	Imp/OstA is required for cell envelope biogenesis in Escherichia coli. <i>Molecular Microbiology</i> , <b>2002</b> , 45, 1289-302	4.1	200
179	Genetic evidence for parallel pathways of chaperone activity in the periplasm of Escherichia coli. <i>Journal of Bacteriology</i> , <b>2001</b> , 183, 6794-800	3.5	200
178	Sequence analysis of mutations that prevent export of lambda receptor, an Escherichia coli outer membrane protein. <i>Nature</i> , <b>1980</b> , 285, 82-5	50.4	195
177	Targeting and assembly of periplasmic and outer-membrane proteins in Escherichia coli. <i>Annual Review of Genetics</i> , <b>1998</b> , 32, 59-94	14.5	194
176	The Cpx envelope stress response is controlled by amplification and feedback inhibition. <i>Journal of Bacteriology</i> , <b>1999</b> , 181, 5263-72	3.5	184
175	Identification of two inner-membrane proteins required for the transport of lipopolysaccharide to the outer membrane of Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 5537-42	11.5	183
174	Functional analysis of the protein machinery required for transport of lipopolysaccharide to the outer membrane of Escherichia coli. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 4460-9	3.5	181
173	Heat-shock proteins DnaK and GroEL facilitate export of LacZ hybrid proteins in E. coli. <i>Nature</i> , <b>1990</b> , 344, 882-4	50.4	179
172	The sec and prl genes of Escherichia coli. <i>Journal of Bioenergetics and Biomembranes</i> , <b>1990</b> , 22, 291-310	3.7	172
171	Signal detection and target gene induction by the CpxRA two-component system. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 2432-40	3.5	169
170	EnvZ controls the concentration of phosphorylated OmpR to mediate osmoregulation of the porin genes. <i>Journal of Molecular Biology</i> , <b>1991</b> , 222, 567-80	6.5	164
169	The sigmaE and Cpx regulatory pathways: overlapping but distinct envelope stress responses. <i>Current Opinion in Microbiology</i> , <b>1999</b> , 2, 159-65	7.9	159
168	A signal sequence is not sufficient to lead beta-galactosidase out of the cytoplasm. <i>Nature</i> , <b>1980</b> , 286, 356-9	50.4	157

167	Two-Component Signal Transduction Systems: Structure-Function Relationships and Mechanisms of Catalysis <b>2014</b> , 25-51		155
166	Characterization of the two-protein complex in Escherichia coli responsible for lipopolysaccharide assembly at the outer membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2010</b> , 107, 5363-8	11.5	152
165	Mutations affecting localization of an Escherichia coli outer membrane protein, the bacteriophage lambda receptor. <i>Journal of Molecular Biology</i> , <b>1980</b> , 141, 63-90	6.5	148
164	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> , <b>2008</b> , 70, 323-4	4 <del>0</del> ·1	145
163	Outer Membrane Biogenesis. <i>Annual Review of Microbiology</i> , <b>2017</b> , 71, 539-556	17.5	142
162	The Bam machine: a molecular cooper. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , <b>2012</b> , 1818, 1067-8	<b>8<u>4</u>8</b>	136
161	Quality control in the bacterial periplasm. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , <b>2004</b> , 1694, 121-34	4.9	134
160	PrlA (SecY) and PrlG (SecE) interact directly and function sequentially during protein translocation in E. coli. <i>Cell</i> , <b>1990</b> , 61, 833-42	56.2	132
159	Mutations that alter the kinase and phosphatase activities of the two-component sensor EnvZ. Journal of Bacteriology, <b>1998</b> , 180, 4538-46	3.5	126
158	A previously unidentified gene in the spc operon of Escherichia coli K12 specifies a component of the protein export machinery. <i>Cell</i> , <b>1982</b> , 31, 227-35	56.2	122
157	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> , <b>2005</b> , 102, 17775-9	11.5	117
156	Characterization of the role of the Escherichia coli periplasmic chaperone SurA using differential proteomics. <i>Proteomics</i> , <b>2009</b> , 9, 2432-43	4.8	116
155	Genetic basis for activity differences between vancomycin and glycolipid derivatives of vancomycin. <i>Science</i> , <b>2001</b> , 294, 361-4	33.3	112
154	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> , <b>2005</b> , 187, 7680-6	3.5	111
153	Structure of the malB region in Escherichia coli K12. II. Genetic map of the malE,F,G operon. <i>Molecular Genetics and Genomics</i> , <b>1979</b> , 174, 249-59		109
152	Genetic analysis of the switch that controls porin gene expression in Escherichia coli K-12. <i>Journal of Molecular Biology</i> , <b>1989</b> , 210, 281-92	6.5	104
151	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> , <b>2016</b> , 113, E15	565-74	97
150	Lipoprotein LptE is required for the assembly of LptD by the beta-barrel assembly machine in the outer membrane of Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> <b>2011</b> 108 2492-7	11.5	97

149	Crl stimulates RpoS activity during stationary phase. <i>Molecular Microbiology</i> , <b>1998</b> , 29, 1225-36	4.1	96
148	The essential tension: opposed reactions in bacterial two-component regulatory systems. <i>Trends in Microbiology</i> , <b>1993</b> , 1, 306-10	12.4	95
147	Transmembrane domain of surface-exposed outer membrane lipoprotein RcsF is threaded through the lumen of Ebarrel proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, E4350-8	11.5	90
146	The free and bound forms of Lpp occupy distinct subcellular locations in Escherichia coli. <i>Molecular Microbiology</i> , <b>2011</b> , 79, 1168-81	4.1	90
145	Effects of antibiotics and a proto-oncogene homolog on destruction of protein translocator SecY. <i>Science</i> , <b>2009</b> , 325, 753-6	33.3	89
144	Mapping an interface of SecY (PrlA) and SecE (PrlG) by using synthetic phenotypes and in vivo cross-linking. <i>Journal of Bacteriology</i> , <b>1999</b> , 181, 3438-44	3.5	89
143	Sirtuins are evolutionarily conserved viral restriction factors. MBio, 2014, 5,	7.8	86
142	Nonconsecutive disulfide bond formation in an essential integral outer membrane protein.  Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12245-50	11.5	86
141	Escherichia coli starvation diets: essential nutrients weigh in distinctly. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 7549-53	3.5	86
140	Conformation-specific labeling of BamA and suppressor analysis suggest a cyclic mechanism for Ebarrel assembly in Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 5151-6	11.5	85
139	Tethering of CpxP to the inner membrane prevents spheroplast induction of the cpx envelope stress response. <i>Molecular Microbiology</i> , <b>2000</b> , 37, 1186-97	4.1	82
138	Mutational activation of the Cpx signal transduction pathway of Escherichia coli suppresses the toxicity conferred by certain envelope-associated stresses. <i>Molecular Microbiology</i> , <b>1995</b> , 18, 491-505	4.1	82
137	Sequence information within the lamB genes in required for proper routing of the bacteriophage lambda receptor protein to the outer membrane of Escherichia coli K-12. <i>Journal of Molecular Biology</i> , <b>1982</b> , 156, 93-112	6.5	82
136	prlF and yhaV encode a new toxin-antitoxin system in Escherichia coli. <i>Journal of Molecular Biology</i> , <b>2007</b> , 372, 894-905	6.5	81
135	Outer membrane lipoprotein biogenesis: Lol is not the end. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , <b>2015</b> , 370,	5.8	80
134	Olaf Schneewind, 1961 2019: Scientist, Mentor, Friend. <i>Journal of Bacteriology</i> , <b>2019</b> , 201,	3.5	78
133	Envelope Stress Responses: An Interconnected Safety Net. <i>Trends in Biochemical Sciences</i> , <b>2017</b> , 42, 232	2-243	78
132	2019 Jack Kenney Award for Outstanding Service. <i>Journal of Bacteriology</i> , <b>2019</b> , 202,	3.5	78

131	Current Issues in Scientific Publishing. <i>Journal of Bacteriology</i> , <b>2019</b> , 202,	3.5	78
130	Accumulation of the enterobacterial common antigen lipid II biosynthetic intermediate stimulates degP transcription in Escherichia coli. <i>Journal of Bacteriology</i> , <b>1998</b> , 180, 5875-84	3.5	76
129	Complex spatial distribution and dynamics of an abundant Escherichia coli outer membrane protein, LamB. <i>Molecular Microbiology</i> , <b>2004</b> , 53, 1771-83	4.1	75
128	Information within the mature LamB protein necessary for localization to the outer membrane of E coli K12. <i>Cell</i> , <b>1983</b> , 32, 1325-35	56.2	75
127	Kinetic analysis of the assembly of the outer membrane protein LamB in Escherichia coli mutants each lacking a secretion or targeting factor in a different cellular compartment. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 446-54	3.5	73
126	The Cpx stress response confers resistance to some, but not all, bactericidal antibiotics. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 1869-74	3.5	71
125	Envelope stress responses: balancing damage repair and toxicity. <i>Nature Reviews Microbiology</i> , <b>2019</b> , 17, 417-428	22.2	68
124	Activation of the Escherichia coli Ebarrel 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> , <b>2012</b> , 109, 3487-91	11.5	68
123	The art and design of genetic screens: Escherichia coli. <i>Nature Reviews Genetics</i> , <b>2003</b> , 4, 419-31	30.1	68
122	Mutations that affect separate functions of OmpR the phosphorylated regulator of porin transcription in Escherichia coli. <i>Journal of Molecular Biology</i> , <b>1993</b> , 231, 261-73	6.5	68
121	Starvation for different nutrients in Escherichia coli results in differential modulation of RpoS levels and stability. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 434-42	3.5	66
120	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> , <b>2019</b> , 116, 217	48-277	5 <b>9</b> 4
119	Porin Regulon of Escherichia coli105-127		64
118	BamE modulates the Escherichia coli beta-barrel assembly machine component BamA. <i>Journal of Bacteriology</i> , <b>2012</b> , 194, 1002-8	3.5	63
117	Characterization of a stalled complex on the Ebarrel assembly machine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2016</b> , 113, 8717-22	11.5	62
116	Making a beta-barrel: assembly of outer membrane proteins in Gram-negative bacteria. <i>Current Opinion in Microbiology</i> , <b>2012</b> , 15, 189-93	7.9	61
115	OmpR mutants specifically defective for transcriptional activation. <i>Journal of Molecular Biology</i> , <b>1994</b> , 243, 579-94	6.5	60
114	Redefining the essential trafficking pathway for outer membrane lipoproteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2017</b> , 114, 4769-4774	11.5	59

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113	A suppressor of cell death caused by the loss of sigmal downregulates extracytoplasmic stress responses and outer membrane vesicle production in Escherichia coli. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 1523-30	3.5	59
112	Secretion of LamB-LacZ by the signal recognition particle pathway of Escherichia coli. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 5697-705	3.5	59
111	Constitutive activation of the Escherichia coli Pho regulon upregulates rpoS translation in an Hfq-dependent fashion. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 5984-92	3.5	58
110	A lipoprotein/Ebarrel complex monitors lipopolysaccharide integrity transducing information across the outer membrane. <i>ELife</i> , <b>2016</b> , 5,	8.9	58
109	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> , <b>1983</b> , 166, 273-82	6.5	57
108	Probing the barrier function of the outer membrane with chemical conditionality. <i>ACS Chemical Biology</i> , <b>2006</b> , 1, 385-95	4.9	55
107	Identification of base pairs important for OmpR-DNA interaction. <i>Molecular Microbiology</i> , <b>1995</b> , 17, 565-	-743 <u>1</u>	53
106	LptE binds to and alters the physical state of LPS to catalyze its assembly at the cell surface.  Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9467-72	11.5	52
105	Genetic fusions as experimental tools. <i>Methods in Enzymology</i> , <b>1991</b> , 204, 213-48	1.7	51
104	Engineering Escherichia coli to secrete heterologous gene products. <i>Methods in Enzymology</i> , <b>1990</b> , 185, 166-87	1.7	49
103	The LysR homolog LrhA promotes RpoS degradation by modulating activity of the response regulator sprE. <i>Journal of Bacteriology</i> , <b>1999</b> , 181, 563-71	3.5	48
102	Dissecting the Escherichia coli periplasmic chaperone network using differential proteomics. <i>Proteomics</i> , <b>2012</b> , 12, 1391-401	4.8	46
101	RpoS proteolysis is regulated by a mechanism that does not require the SprE (RssB) response regulator phosphorylation site. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 7403-10	3.5	45
100	Regulation of Capsule Synthesis: Modification of the Two-Component Paradigm by an Accessory Unstable Regulator253-262		45
99	The response regulator SprE (RssB) is required for maintaining poly(A) polymerase I-degradosome association during stationary phase. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 3713-21	3.5	44
98	LrhA regulates rpoS translation in response to the Rcs phosphorelay system in Escherichia coli. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 3175-81	3.5	44
97	Crl facilitates RNA polymerase holoenzyme formation. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 7966-70	3.5	44
96	Decline in ribosomal fidelity contributes to the accumulation and stabilization of the master stress response regulator sigmaS upon carbon starvation. <i>Genes and Development</i> , <b>2007</b> , 21, 862-74	12.6	44

95	The Phospholipase PldA Regulates Outer Membrane Homeostasis via Lipid Signaling. MBio, 2018, 9,	7.8	42
94	Characterization and in vivo cloning of prlC, a suppressor of signal sequence mutations in Escherichia coli K12. <i>Genetics</i> , <b>1987</b> , 116, 513-21	4	42
93	Control of Cellular Development in Sporulating Bacteria by the Phosphorelay Two-Component Signal Transduction System <b>2014</b> , 129-144		41
92	The CpxQ sRNA Negatively Regulates Skp To Prevent Mistargeting of Barrel Outer Membrane Proteins into the Cytoplasmic Membrane. <i>MBio</i> , <b>2016</b> , 7, e00312-16	7.8	40
91	RpoS proteolysis is controlled directly by ATP levels in Escherichia coli. <i>Genes and Development</i> , <b>2012</b> , 26, 548-53	12.6	38
90	Dual Sensors and Dual Response Regulators Interact to Control Nitrate- and Nitrite-Responsive Gene Expression in Escherichia coli233-252		38
89	Role for Skp in LptD assembly in Escherichia coli. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 3734-42	3.5	36
88	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> , <b>2018</b> , 115, 2359-2364	11.5	35
87	Continuous control in bacterial regulatory circuits. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 7618-25	3.5	34
86	RpoS-dependent transcriptional control of sprE: regulatory feedback loop. <i>Journal of Bacteriology</i> , <b>2001</b> , 183, 5974-81	3.5	34
85	Classifying EBarrel Assembly Substrates by Manipulating Essential Bam Complex Members. Journal of Bacteriology, <b>2016</b> , 198, 1984-92	3.5	34
84	P pilus assembly motif necessary for activation of the CpxRA pathway by PapE in Escherichia coli. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 4326-37	3.5	32
83	Making a membrane on the other side of the wall. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , <b>2017</b> , 1862, 1386-1393	5	31
82	The genetics of protein secretion in E. coli. <i>Trends in Genetics</i> , <b>1990</b> , 6, 329-34	8.5	31
81	Inhibitor of intramembrane protease RseP blocks the Iresponse causing lethal accumulation of unfolded outer membrane proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, E6614-E6621	11.5	31
80	Predicting functionally informative mutations in Escherichia coli BamA using evolutionary covariance analysis. <i>Genetics</i> , <b>2013</b> , 195, 443-55	4	30
79	Ti Plasmid and Chromosomally Encoded Two-Component Systems Important in Plant Cell Transformation by Agrobacterium Species367-385		30
78	Transcriptional occlusion caused by overlapping promoters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, 1557-61	11.5	29

#### (2019-2013)

77	Dominant negative lptE mutation that supports a role for LptE as a plug in the LptD barrel. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 1327-34	3.5	28	
76	Involvement of a tryptophan residue in the binding site of Escherichia coli galactose-binding protein. <i>Biochemistry</i> , <b>1974</b> , 13, 993-9	3.2	28	
75	Cyclic Enterobacterial Common Antigen Maintains the Outer Membrane Permeability Barrier of Escherichia coli in a Manner Controlled by YhdP. <i>MBio</i> , <b>2018</b> , 9,	7.8	28	
74	Accumulation of phosphatidic acid increases vancomycin resistance in Escherichia coli. <i>Journal of Bacteriology</i> , <b>2014</b> , 196, 3214-20	3.5	27	
73	Distinctive Roles for Periplasmic Proteases in the Maintenance of Essential Outer Membrane Protein Assembly. <i>Journal of Bacteriology</i> , <b>2017</b> , 199,	3.5	27	
7 <sup>2</sup>	YejM Modulates Activity of the YciM/FtsH Protease Complex To Prevent Lethal Accumulation of Lipopolysaccharide. <i>MBio</i> , <b>2020</b> , 11,	7.8	27	
71	The Activity of Escherichia coli Chaperone SurA Is Regulated by Conformational Changes Involving a Parvulin Domain. <i>Journal of Bacteriology</i> , <b>2016</b> , 198, 921-9	3.5	25	
70	Transposition of lambda placMu is mediated by the A protein altered at its carboxy-terminal end. <i>Gene</i> , <b>1988</b> , 71, 177-86	3.8	24	
69	Novel RpoS-Dependent Mechanisms Strengthen the Envelope Permeability Barrier during Stationary Phase. <i>Journal of Bacteriology</i> , <b>2017</b> , 199,	3.5	23	
68	PrlC, a suppressor of signal sequence mutations in Escherichia coli, can direct the insertion of the signal sequence into the membrane. <i>Journal of Molecular Biology</i> , <b>1989</b> , 205, 665-76	6.5	23	
67	Sirtuin Lipoamidase Activity Is Conserved in Bacteria as a Regulator of Metabolic Enzyme Complexes. <i>MBio</i> , <b>2017</b> , 8,	7.8	22	
66	Structural and Functional Conservation in Response Regulators53-64		22	
65	The activity and specificity of the outer membrane protein chaperone SurA are modulated by a proline isomerase domain. <i>MBio</i> , <b>2013</b> , 4,	7.8	21	
64	HisAsp Phosphorelay: Two Components or More?. <i>Cell</i> , <b>1996</b> , 85, 13-14	56.2	21	
63	Two-Component Signal Transduction and Its Role in the Expression of Bacterial Virulence Factors303-3	317	21	
62	A Signal Transduction Network in Bacillus subtilis Includes the DegS/DegU and ComP/ComA Two-Component Systems447-471		21	
61	Control of Nitrogen Assimilation by the NRI-NRII Two-Component System of Enteric Bacteria65-88		21	
60	The Synthetic Phenotype of IDouble Mutants Results from a Lethal Jamming of the Bam Complex by the Lipoprotein RcsF. <i>MBio</i> , <b>2019</b> , 10,	7.8	21	

59	Genetic Approaches for Signaling Pathways and Proteins <b>2014</b> , 7-23		20
58	Absence of the outer membrane phospholipase A suppresses the temperature-sensitive phenotype of Escherichia coli degP mutants and induces the Cpx and sigma(E) extracytoplasmic stress responses. <i>Journal of Bacteriology</i> , <b>2001</b> , 183, 5230-8	3.5	20
57	Conferral of transposable properties to a chromosomal gene in Escherichia coli. <i>Journal of Molecular Biology</i> , <b>1980</b> , 141, 235-48	5.5	20
56	Flagellar Switch181-199		20
55	Regulation of Salmonella Virulence by Two-Component Regulatory Systems319-332		20
54	Bordetella pertussis BvgAS Virulence Control System <b>2014</b> , 333-349		19
53	Assembly of Outer Membrane Barrel Proteins: the Bam Complex. <i>EcoSal Plus</i> , <b>2011</b> , 4,	7.7	19
52	Gene fusions. <i>Journal of Bacteriology</i> , <b>2000</b> , 182, 5935-8	3.5	19
51	Genetic analysis of protein export in Escherichia coli. <i>Methods in Enzymology</i> , <b>1983</b> , 97, 3-11	1.7	19
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4	Regulation of Glycopeptide Resistance Genes of Enterococcal Transposon Tn1546 by the VanR-VanS Two-Component Regulatory System387-391		2
3	The sacrificial adaptor protein Skp functions to remove stalled substrates from the Ebarrel assembly machine <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119,	11.5	2
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