Arnold JM Driessen

List of Publications by Year
in descending order

244 papers	17,225 citations	14655 66 h-index	20358 116 g-index
250 all docs	250 docs citations	250 times ranked	13324 citing authors

#	Article	IF	Citations
1	Genome sequencing and analysis of the versatile cell factory Aspergillus niger CBS 513.88. Nature Biotechnology, 2007, 25, 221-231.	17.5	1,047
2	Minimum Information about a Biosynthetic Gene cluster. Nature Chemical Biology, 2015, 11, 625-631.	8.0	715
3	The purified E. coli integral membrane protein SecYE is sufficient for reconstitution of SecA-dependent precursor protein translocation. Cell, 1990, 62, 649-657.	28.9	539
4	Î”Î¼H+ and ATP function at different steps of the catalytic cycle of preprotein translocase. Cell, 1991, 64, 927-939.	28.9	487
5	Genome sequencing and analysis of the filamentous fungus Penicillium chrysogenum. Nature Biotechnology, 2008, 26, 1161-1168.	17.5	427
6	Optical control of antibacterial activity. Nature Chemistry, 2013, 5, 924-928.	13.6	298
7	A presequence- and voltage-sensitive channel of the mitochondrial preprotein translocase formed by Tim23. Nature Structural Biology, 2001, 8, 1074-1082.	9.7	287
8	Multidrug resistance mediated by a bacterial homolog of the human multidrug transporter MDR1.. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 10668-10672.	7.1	282
9	Distribution and Physiology of ABC-Type Transporters Contributing to Multidrug Resistance in Bacteria. Microbiology and Molecular Biology Reviews, 2007, 71, 463-476.	6.6	270
10	CRISPR/Cas9 Based Genome Editing of <i>Penicillium chrysogenum</i>. ACS Synthetic Biology, 2016, 5, 754-764.	3.8	258
11	Photocontrol of Antibacterial Activity: Shifting from UV to Red Light Activation. Journal of the American Chemical Society, 2017, 139, 17979-17986.	13.7	224
12	<i>Escherichia coli</i> translocase: the unravelling of a molecular machine. Molecular Microbiology, 2000, 37, 226-238.	2.5	222
13	The essence of being extremophilic: the role of the unique archaeal membrane lipids. Extremophiles, 1998, 2, 163-170.	2.3	199
14	Bioenergetics and Solute Transport in Lactococci. CRC Critical Reviews in Microbiology, 1989, 16, 419-476.	4.8	188
15	F1F0 ATP synthase subunit c is a substrate of the novel YidC pathway for membrane protein biogenesis. Journal of Cell Biology, 2004, 165, 213-222.	5.2	186
16	The structural basis of protein targeting and translocation in bacteria. Nature Structural Biology, 2001, 8, 492-498.	9.7	184
17	The molecular chaperone SecB is released from the carboxy-terminus of SecA during initiation of precursor protein translocation. EMBO Journal, 1997, 16, 6105-6113.	7.8	181
18	The catalytic cycle of the Escherichia coli SecA ATPase comprises two distinct preprotein translocation events. EMBO Journal, 1997, 16, 7297-7304.	7.8	181

#	ARTICLE	IF	CITATIONS
19	In vitro pore-forming activity of the lantibiotic nisin. Role of protonmotive force and lipid composition. FEBS Journal, 1993, 212, 417-422.	0.2	161
20	Enantioselective Artificial Metalloenzymes by Creation of a Novel Active Site at the Protein Dimer Interface. Angewandte Chemie - International Edition, 2012, 51, 7472-7475.	13.8	154
21	NisT, the Transporter of the Lantibiotic Nisin, Can Transport Fully Modified, Dehydrated, and Unmodified Prenisin and Fusions of the Leader Peptide with Non-lantibiotic Peptides. Journal of Biological Chemistry, 2004, 279, 22176-22182.	3.4	152
22	SecDFyajC forms a heterotetrameric complex with YidC. Molecular Microbiology, 2002, 44, 1397-1405.	2.5	148
23	Proteomic analysis of secreted membrane vesicles of archaeal Sulfolobus species reveals the presence of endosome sorting complex components. Extremophiles, 2009, 13, 67-79.	2.3	148
24	Crystal Structures of the ATPase Subunit of the Glucose ABC Transporter from Sulfolobus solfataricus: Nucleotide-free and Nucleotide-bound Conformations. Journal of Molecular Biology, 2003, 330, 343-358.	4.2	145
25	Identification of Diverse Archaeal Proteins with Class III Signal Peptides Cleaved by Distinct Archaeal Prepilin Peptidases. Journal of Bacteriology, 2007, 189, 772-778.	2.2	139
26	Dissection and Modulation of the Four Distinct Activities of Nisin by Mutagenesis of Rings A and B and by C-Terminal Truncation. Applied and Environmental Microbiology, 2007, 73, 5809-5816.	3.1	139
27	Substrate Specificity of the SecB Chaperone. Journal of Biological Chemistry, 1999, 274, 34219-34225.	3.4	137
28	UVâ€inducible cellular aggregation of the hyperthermophilic archaeon <i>Sulfolobus solfataricus</i> is mediated by pili formation. Molecular Microbiology, 2008, 70, 938-952.	2.5	137
29	Biosynthesis of archaeal membrane ether lipids. Frontiers in Microbiology, 2014, 5, 641.	3.5	137
30	A conserved function of YidC in the biogenesis of respiratory chain complexes. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 5801-5806.	7.1	133
31	Direct Observation of Chaperone-Induced Changes in a Protein Folding Pathway. Science, 2007, 318, 1458-1461.	12.6	133
32	Archaeal Homolog of Bacterial Type IV Prepilin Signal Peptidases with Broad Substrate Specificity. Journal of Bacteriology, 2003, 185, 3918-3925.	2.2	129
33	The uncoupling efficiency and affinity of flavonoids for vesicles. Biochemical Pharmacology, 2000, 60, 1593-1600.	4.4	123
34	Sugar transport in Sulfolobus solfataricus is mediated by two families of binding protein-dependent ABC transporters. Molecular Microbiology, 2001, 39, 1494-1503.	2.5	121
35	Bioenergetics and cytoplasmic membrane stability of the extremely acidophilic, thermophilic archaeon Picrophilus oshimae. Extremophiles, 1998, 2, 67-74.	2.3	120
36	Preprotein transfer to theEscherichia colitranslocase requires the coâ€operative binding of SecB and the signal sequence to SecA. Molecular Microbiology, 1998, 29, 1179-1190.	2.5	119

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37	Structure of the transcriptional regulator LmrR and its mechanism of multidrug recognition. EMBO Journal, 2009, 28, 156-166.	7.8	118
38	Reshaping of the conformational search of a protein by the chaperone trigger factor. Nature, 2013, 500, 98-101.	27.8	118
39	The Lactococcal lmrP Gene Encodes a Proton Motive Force- dependent Drug Transporter. Journal of Biological Chemistry, 1995, 270, 26092-26098.	3.4	116
40	Production of Recombinant and Tagged Proteins in the Hyperthermophilic Archaeon Sulfolobus solfataricus. Applied and Environmental Microbiology, 2006, 72, 102-111.	3.1	116
41	Non-bilayer Lipids Stimulate the Activity of the Reconstituted Bacterial Protein Translocase. Journal of Biological Chemistry, 2000, 275, 2472-2478.	3.4	114
42	Supramolecular Assembly of Artificial Metalloenzymes Based on the Dimeric Protein LmrR as Promiscuous Scaffold. Journal of the American Chemical Society, 2015, 137, 9796-9799.	13.7	114
43	UV-inducible DNA exchange in hyperthermophilic archaea mediated by type IV pili. Molecular Microbiology, 2011, 82, 807-817.	2.5	113
44	Isoprenoid biosynthesis in Archaea â€“ Biochemical and evolutionary implications. Research in Microbiology, 2011, 162, 39-52.	2.1	109
45	Translocation of proteins across the cell envelope of Gram-positive bacteria. FEMS Microbiology Reviews, 2001, 25, 437-454.	8.6	106
46	Structural Organization of Essential Iron-Sulfur Clusters in the Evolutionarily Highly Conserved ATP-binding Cassette Protein ABCE1. Journal of Biological Chemistry, 2007, 282, 14598-14607.	3.4	99
47	The role of transport processes in survival of lactic acid bacteria. Energy transduction and multidrug resistance. Antonie Van Leeuwenhoek, 1997, 71, 117-128.	1.7	98
48	Energetics and Mechanism of Drug Transport Mediated by the Lactococcal Multidrug Transporter LmrP. Journal of Biological Chemistry, 1996, 271, 24123-24128.	3.4	91
49	PrlA4 prevents the rejection of signal sequence defective preproteins by stabilizing the SecA-SecY interaction during the initiation of translocation. EMBO Journal, 1998, 17, 3631-3639.	7.8	91
50	LmrCD is a major multidrug resistance transporter in Lactococcus lactis. Molecular Microbiology, 2006, 61, 771-781.	2.5	91
51	Nark is a nitrite-extrusion system involved in anaerobic nitrate respiration by Escherichia coli. Molecular Microbiology, 1994, 12, 579-586.	2.5	87
52	The Lateral Gate of SecYEG Opens during Protein Translocation. Journal of Biological Chemistry, 2009, 284, 15805-15814.	3.4	87
53	Ciprofloxacinâ€“Photoswitch Conjugates: A Facile Strategy for Photopharmacology. Bioconjugate Chemistry, 2015, 26, 2592-2597.	3.6	86
54	SecA Is Not Required for Signal Recognition Particle-mediated Targeting and Initial Membrane Insertion of a Nascent Inner Membrane Protein. Journal of Biological Chemistry, 1999, 274, 29883-29888.	3.4	85

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55	Formation of the Productive ATP-Mg 2+ -bound Dimer of GlcV, an ABC-ATPase from Sulfolobus solfataricus. Journal of Molecular Biology, 2003, 334, 255-267.	4.2	84
56	Appendage-Mediated Surface Adherence of <i>Sulfolobus solfataricus</i>. Journal of Bacteriology, 2010, 192, 104-110.	2.2	84
57	How hyperthermophiles adapt to change their lives: DNA exchange in extreme conditions. Extremophiles, 2013, 17, 545-563.	2.3	84
58	The <i>Escherichia coli</i> multidrug transporter MdfA catalyzes both electrogenic and electroneutral transport reactions. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 1667-1672.	7.1	80
59	Characterization of a Bacillus subtilis SecA mutant protein deficient in translocation ATPase and release from the membrane. Molecular Microbiology, 1993, 8, 31-42.	2.5	77
60	Insights into ABC Transport in Archaea. Journal of Bioenergetics and Biomembranes, 2004, 36, 5-15.	2.3	76
61	Covalently Dimerized SecA Is Functional in Protein Translocation. Journal of Biological Chemistry, 2005, 280, 35255-35260.	3.4	75
62	Subunit a of Cytochrome o Oxidase Requires Both YidC and SecYEG for Membrane Insertion. Journal of Biological Chemistry, 2006, 281, 12248-12252.	3.4	75
63	Expanding and understanding the genetic toolbox of the hyperthermophilic genus <i>Sulfolobus</i>. Biochemical Society Transactions, 2009, 37, 97-101.	3.4	75
64	Production of Dehydroamino Acid-Containing Peptides by Lactococcus lactis. Applied and Environmental Microbiology, 2007, 73, 1792-1796.	3.1	74
65	Flagellar Motility and Structure in the Hyperthermoacidophilic Archaeon Sulfolobus solfataricus. Journal of Bacteriology, 2007, 189, 4305-4309.	2.2	73
66	A Branched Biosynthetic Pathway Is Involved in Production of Roquefortine and Related Compounds in Penicillium chrysogenum. PLoS ONE, 2013, 8, e65328.	2.5	73
67	Orthogonal Control of Antibacterial Activity with Light. ACS Chemical Biology, 2014, 9, 1969-1974.	3.4	73
68	Regulation of archaella expression by the FHA and von Willebrand domainâ€containing proteins ArnA and ArnB in <i>Sulfolobus acidocaldarius</i>. Molecular Microbiology, 2012, 86, 24-36.	2.5	72
69	SecB, a molecular chaperone with two faces. Trends in Microbiology, 2001, 9, 193-196.	7.7	64
70	Probing the SecYEG translocation pore size with preproteins conjugated with sizable rigid spherical molecules. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7775-7780.	7.1	64
71	Identification of a system required for the functional surface localization of sugar binding proteins with class III signal peptides in Sulfolobus solfataricus. Molecular Microbiology, 2007, 64, 795-806.	2.5	63
72	Quaternary Structure of SecA in Solution and Bound to SecYEG Probed at the Single Molecule Level. Structure, 2011, 19, 430-439.	3.3	63

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73	SecBâ€"A chaperone dedicated to protein translocation. Molecular BioSystems, 2010, 6, 620-627.	2.9	62
74	Identification of a Polyketide Synthase Involved in Sorbicillin Biosynthesis by Penicillium chrysogenum. Applied and Environmental Microbiology, 2016, 82, 3971-3978.	3.1	62
75	Diversity of transport mechanisms: common structural principles. Trends in Biochemical Sciences, 2000, 25, 397-401.	7.5	61
76	Protein secretion in the Archaea: multiple paths towards a unique cell surface. Nature Reviews Microbiology, 2006, 4, 537-547.	28.6	61
77	Cell Surface Structures of Archaea. Journal of Bacteriology, 2008, 190, 6039-6047.	2.2	61
78	LmrR Is a Transcriptional Repressor of Expression of the Multidrug ABC Transporter LmrCD in <i>Lactococcus lactis</i>. Journal of Bacteriology, 2008, 190, 759-763.	2.2	61
79	SecA, a remarkable nanomachine. Cellular and Molecular Life Sciences, 2011, 68, 2053-2066.	5.4	61
80	Bacterial protein translocation: kinetic and thermodynamic role of ATP and the protonmotive force. Trends in Biochemical Sciences, 1992, 17, 219-223.	7.5	60
81	Kinetic Analysis of the Translocation of Fluorescent Precursor Proteins into Escherichia coli Membrane Vesicles. Journal of Biological Chemistry, 2002, 277, 46059-46065.	3.4	60
82	A single copy of SecYEG is sufficient for preprotein translocation. EMBO Journal, 2011, 30, 4387-4397.	7.8	60
83	Growing Membranes <i>In Vitro</i> by Continuous Phospholipid Biosynthesis from Free Fatty Acids. ACS Synthetic Biology, 2018, 7, 153-165.	3.8	60
84	Conditions for gene disruption by homologous recombination of exogenous DNA into theSulfolobus solfataricusgenome. Archaea, 2008, 2, 145-149.	2.3	59
85	Compartmentalization and transport in beta-lactam antibiotic biosynthesis by filamentous fungi. Antonie Van Leeuwenhoek, 1999, 75, 41-78.	1.7	58
86	The polymerization mechanism of the bacterial cell division protein FtsZ. FEBS Letters, 2001, 506, 6-10.	2.8	57
87	YidC â€" an evolutionary conserved device for the assembly of energy-transducing membrane protein complexes. Current Opinion in Microbiology, 2005, 8, 182-187.	5.1	57
88	Substrate Recognition and Specificity of the NisB Protein, the Lantibiotic Dehydratase Involved in Nisin Biosynthesis. Journal of Biological Chemistry, 2011, 286, 30552-30560.	3.4	57
89	SecA Supports a Constant Rate of Preprotein Translocation*. Journal of Biological Chemistry, 2006, 281, 15709-15713.	3.4	56
90	Mechanisms of YidC-mediated Insertion and Assembly of Multimeric Membrane Protein Complexes. Journal of Biological Chemistry, 2008, 283, 31269-31273.	3.4	56

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91	Lightâ€Induced Control of Protein Translocation by the SecYEG Complex. Angewandte Chemie - International Edition, 2010, 49, 7234-7238.	13.8	56
92	How proteins cross the bacterial cytoplasmic membrane. Journal of Membrane Biology, 1994, 142, 145-59.	2.1	55
93	â€œHot standardsâ€•for the thermoacidophilic archaeon Sulfolobus solfataricus. Extremophiles, 2010, 14, 119-142.	2.3	55
94	Mapping the Sites of Interaction between SecY and SecE by Cysteine Scanning Mutagenesis. Journal of Biological Chemistry, 2001, 276, 32559-32566.	3.4	54
95	YidC and SecY Mediate Membrane Insertion of a Type I Transmembrane Domain. Journal of Biological Chemistry, 2002, 277, 35880-35886.	3.4	54
96	Reconstitution of Purified Bacterial Preprotein Translocase in Liposomes. Methods in Enzymology, 2003, 372, 86-98.	1.0	54
97	ydaG and ydbA of Lactococcus lactis Encode a Heterodimeric ATP-binding Cassette-type Multidrug Transporter. Journal of Biological Chemistry, 2004, 279, 34449-34455.	3.4	54
98	Pushing, pulling and trapping - Modes of motor protein supported protein translocation. FEBS Letters, 2007, 581, 2820-2828.	2.8	54
99	Sec-dependent preprotein translocation in bacteria. Archives of Microbiology, 1996, 165, 1-8.	2.2	52
100	Elucidating the Native Architecture of the YidC: Ribosome Complex. Journal of Molecular Biology, 2013, 425, 4112-4124.	4.2	52
101	Non-hydrolysable GTP-gamma-S stabilizes the FtsZ polymer in a GDP-bound state. Molecular Microbiology, 2000, 35, 1211-1219.	2.5	51
102	Signal peptides of secreted proteins of the archaeon Sulfolobus solfataricus : a genomic survey. Archives of Microbiology, 2002, 177, 209-216.	2.2	51
103	Stepwise evolution of the Sec machinery in Proteobacteria. Trends in Microbiology, 2006, 14, 105-108.	7.7	51
104	Cellobiose Uptake in the Hyperthermophilic Archaeon Pyrococcus furiosus Is Mediated by an Inducible, High-Affinity ABC Transporter. Journal of Bacteriology, 2001, 183, 4979-4984.	2.2	50
105	Interaction of SecB with soluble SecA1. FEBS Letters, 1997, 416, 35-38.	2.8	49
106	SecYEG Proteoliposomes Catalyze the Î”Ï•-Dependent Membrane Insertion of FtsQ. Journal of Biological Chemistry, 2004, 279, 1659-1664.	3.4	49
107	The Oligomeric Distribution of SecYEG is Altered by SecA and Translocation Ligands. Journal of Molecular Biology, 2005, 354, 258-271.	4.2	49
108	Novel Key Metabolites Reveal Further Branching of the Roquefortine/Meleagrin Biosynthetic Pathway. Journal of Biological Chemistry, 2013, 288, 37289-37295.	3.4	49

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109	Measurements of the proton motive force generated by cytochrome c oxidase from Bacillus subtilis in proteoliposomes and membrane vesicles. FEBS Journal, 1986, 156, 431-440.	0.2	48
110	Bioenergetics and solute uptake under extreme conditions. Extremophiles, 2001, 5, 285-294.	2.3	48
111	Î´-(l-Î±-Aminoadipyl)-l-cysteinyl-d-valine synthetase, that mediates the first committed step in penicillin biosynthesis, is a cytosolic enzyme. Fungal Genetics and Biology, 2002, 37, 49-55.	2.1	48
112	Co- and post-translational translocation through the protein-conducting channel: analogous mechanisms at work?. Nature Structural and Molecular Biology, 2006, 13, 957-964.	8.2	48
113	Sec-Mediated Transport of Posttranslationally Dehydrated Peptides in <i>Lactococcus lactis</i>. Applied and Environmental Microbiology, 2006, 72, 7626-7633.	3.1	47
114	Nonlinear Biosynthetic Gene Cluster Dose Effect on Penicillin Production by <i>Penicillium chrysogenum</i>. Applied and Environmental Microbiology, 2010, 76, 7109-7115.	3.1	46
115	Co-operation between different targeting pathways during integration of a membrane protein. Journal of Cell Biology, 2012, 199, 303-315.	5.2	46
116	Comparative study of the extracellular proteome of Sulfolobus species reveals limited secretion. Extremophiles, 2010, 14, 87-98.	2.3	45
117	Active-Site Residues in the Type IV Prepilin Peptidase Homologue PibD from the Archaeon Sulfolobus solfataricus. Journal of Bacteriology, 2006, 188, 1437-1443.	2.2	44
118	Ssâ€LrpB, a transcriptional regulator from <i>Sulfolobus solfataricus</i>, regulates a gene cluster with a pyruvate ferredoxin oxidoreductaseâ€encoding operon and permease genes. Molecular Microbiology, 2009, 71, 972-988.	2.5	44
119	Differential effect of YidC depletion on the membrane proteome of <i>Escherichia coli</i> under aerobic and anaerobic growth conditions. Proteomics, 2010, 10, 3235-3247.	2.2	44
120	Competitive Binding of the SecA ATPase and Ribosomes to the SecYEG Translocon. Journal of Biological Chemistry, 2012, 287, 7885-7895.	3.4	44
121	Increased Penicillin Production in Penicillium chrysogenum Production Strains via Balanced Overexpression of Isopenicillin N Acyltransferase. Applied and Environmental Microbiology, 2012, 78, 7107-7113.	3.1	44
122	Identification of the Magnesium-binding Domain of the High-affinity ATP-binding Site of the Bacillus subtilis and Escherichia coli SecA Protein. Journal of Biological Chemistry, 1995, 270, 18975-18982.	3.4	43
123	Regulation of expression of the arabinose and glucose transporter genes in the thermophilic archaeon Sulfolobus solfataricus. Extremophiles, 2006, 10, 383-391.	2.3	43
124	The <i>Bifidobacterium longum</i> NCIMB 702259 ^T <i>ctr</i> Gene Codes for a Novel Cholate Transporter. Applied and Environmental Microbiology, 2006, 72, 923-926.	3.1	43
125	Expression of the transporter encoded by the cefT gene of Acremonium chrysogenum increases cephalosporin production in Penicillium chrysogenum. Fungal Genetics and Biology, 2008, 45, 1415-1421.	2.1	43
126	The Sulfolobicin Genes of Sulfolobus acidocaldariusEncode Novel Antimicrobial Proteins. Journal of Bacteriology, 2011, 193, 4380-4387.	2.2	43

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127	SecY-SecY and SecY-SecG contacts revealed by site-specific crosslinking. FEBS Letters, 2002, 527, 159-165.	2.8	42
128	A Non-Canonical NRPS Is Involved in the Synthesis of Fungisporin and Related Hydrophobic Cyclic Tetrapeptides in Penicillium chrysogenum. PLoS ONE, 2014, 9, e98212.	2.5	42
129	Lipids Activate SecA for High Affinity Binding to the SecYEG Complex. Journal of Biological Chemistry, 2016, 291, 22534-22543.	3.4	42
130	Sec-Mediated Secretion of Bacteriocin Enterocin P by Lactococcus lactis. Applied and Environmental Microbiology, 2005, 71, 1959-1963.	3.1	40
131	Identification of Two Interaction Sites in SecY that Are Important for the Functional Interaction with SecA. Journal of Molecular Biology, 2006, 361, 839-849.	4.2	40
132	Functional incorporation of beef-heart cytochrome c oxidase into membranes of Streptococcus cremoris. FEBS Journal, 1986, 154, 617-624.	0.2	39
133	Immediate GTP hydrolysis upon FtsZ polymerization. Molecular Microbiology, 2002, 43, 1517-1521.	2.5	39
134	Direct Demonstration of ATP-dependent Release of SecA from a Translocating Preprotein by Surface Plasmon Resonance. Journal of Biological Chemistry, 2003, 278, 29581-29586.	3.4	39
135	Bacillus subtilis SpoIIIJ and YqjG Function in Membrane Protein Biogenesis. Journal of Bacteriology, 2009, 191, 6749-6757.	2.2	39
136	Distinct Contributions of the Nisin Biosynthesis Enzymes NisB and NisC and Transporter NisT to Prenisin Production by Lactococcus lactis. Applied and Environmental Microbiology, 2008, 74, 5541-5548.	3.1	38
137	Glutamate transport in Rhodobacter sphaeroides is mediated by a novel binding protein-dependent secondary transport system. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 12786-12790.	7.1	37
138	Enterocin P Causes Potassium Ion Efflux from Enterococcus faecium T136 Cells. Antimicrobial Agents and Chemotherapy, 2001, 45, 901-904.	3.2	37
139	The Large First Periplasmic Loop of SecD and SecF Plays an Important Role in SecDF Functioning. Journal of Bacteriology, 2005, 187, 5857-5860.	2.2	37
140	The ABC-Type Multidrug Resistance Transporter LmrCD Is Responsible for an Extrusion-Based Mechanism of Bile Acid Resistance in <i>Lactococcus lactis</i>. Journal of Bacteriology, 2008, 190, 7357-7366.	2.2	37
141	Subunit a of the F1F0 ATP Synthase Requires YidC and SecYEG for Membrane Insertion. Journal of Molecular Biology, 2009, 390, 893-901.	4.2	37
142	Conserved Negative Charges in the Transmembrane Segments of Subunit K of the NADH:Ubiquinone Oxidoreductase Determine Its Dependence on YidC for Membrane Insertion. Journal of Biological Chemistry, 2010, 285, 3575-3581.	3.4	37
143	Molecular analysis of the UVâ€inducible pili operon from <i>Sulfolobus acidocaldarius</i>. MicrobiologyOpen, 2013, 2, 928-937.	3.0	37
144	Biosynthetic concepts for the production of Î²â€lactam antibiotics in <i>Penicillium chrysogenum</i>. Biotechnology Journal, 2012, 7, 225-236.	3.5	36

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145	Identification of CDP-Archaeol Synthase, a Missing Link of Ether Lipid Biosynthesis in Archaea. Chemistry and Biology, 2014, 21, 1392-1401.	6.0	36
146	Controlling the activity of quorum sensing autoinducers with light. Chemical Science, 2015, 6, 3593-3598.	7.4	36
147	Getting out: protein traffic in prokaryotes. Molecular Microbiology, 2004, 52, 3-11.	2.5	35
148	Monitoring the Activity of Single Translocons. Journal of Molecular Biology, 2013, 425, 4145-4153.	4.2	35
149	New promoters for strain engineering of Penicillium chrysogenum. Fungal Genetics and Biology, 2016, 89, 62-71.	2.1	35
150	Modular Synthetic Biology Toolkit for Filamentous Fungi. ACS Synthetic Biology, 2021, 10, 2850-2861.	3.8	35
151	The positive inside rule is not determined by the polarity of the Î”Ïˆ. Molecular Microbiology, 1998, 29, 1125-1126.	2.5	34
152	SecDFyajC is not required for the maintenance of the proton motive force. FEBS Letters, 2001, 508, 103-106.	2.8	33
153	YidC-mediated Membrane Insertion of Assembly Mutants of Subunit c of the F1F0 ATPase. Journal of Biological Chemistry, 2006, 281, 29762-29768.	3.4	33
154	Energy-transducing properties of primary proton pumps reconstituted into archaeal bipolar lipid vesicles. FEBS Journal, 1993, 214, 917-925.	0.2	32
155	The YidC/Oxa1/Alb3 protein family: common principles and distinct features. Biological Chemistry, 2012, 393, 1279-1290.	2.5	32
156	Bacterial Sec-translocase Unfolds and Translocates a Class of Folded Protein Domains. Journal of Molecular Biology, 2007, 372, 422-433.	4.2	31
157	The <i>Sulfolobus solfataricus</i> AAA protein Sso0909, a homologue of the eukaryotic ESCRT Vps4 ATPase. Biochemical Society Transactions, 2008, 36, 94-98.	3.4	30
158	The aminoâ€terminal tail of Hxt11 confers membrane stability to the Hxt2 sugar transporter and improves xylose fermentation in the presence of acetic acid. Biotechnology and Bioengineering, 2017, 114, 1937-1945.	3.3	30
159	Binding of SecA to the SecYEG Complex Accelerates the Rate of Nucleotide Exchange on SecA. Journal of Biological Chemistry, 2004, 279, 13769-13777.	3.4	29
160	YidC Is Involved in the Biogenesis of Anaerobic Respiratory Complexes in the Inner Membrane of Escherichia coli. Journal of Biological Chemistry, 2008, 283, 26921-26927.	3.4	29
161	The bindosome is a structural component of the Sulfolobus solfataricus cell envelope. Extremophiles, 2011, 15, 235-244.	2.3	29
162	Improved xylose uptake in<i>Saccharomyces cerevisiae</i>due to directed evolution of galactose permease Gal2 for sugar co-consumption. Journal of Applied Microbiology, 2015, 119, 99-111.	3.1	29

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163	Role of the Cytosolic Loop C2 and the C Terminus of YidC in Ribosome Binding and Insertion Activity. Journal of Biological Chemistry, 2015, 290, 17250-17261.	3.4	29
164	Substitution of a conserved aspartate allows cation-induced polymerization of FtsZ. FEBS Letters, 2001, 494, 34-37.	2.8	28
165	Facilitated Drug Influx by an Energy-uncoupled Secondary Multidrug Transporter. Journal of Biological Chemistry, 2004, 279, 103-108.	3.4	28
166	Cloning and molecular characterization of the secY genes from Bacillus licheniformis and Staphylococcus carnosus: comparative analysis of nine members of the SecY family. Molecular Genetics and Genomics, 1992, 235, 147-152.	2.4	27
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