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List of Publications by Year in descending order

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68 papers	5,016 citations	126907 33 h-index	98798 67 g-index
70 all docs	70 docs citations	70 times ranked	4538 citing authors

#	Article	IF	Citations
1	Pyridylpiperazine-based allosteric inhibitors of RND-type multidrug efflux pumps. Nature Communications, 2022, 13, 115.	12.8	28
2	Unidirectional mannitol synthesis ofAcinetobacter baumanniiMtlD is facilitated by the helixâ€“loopâ€“helix-mediated dimer formation. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2107994119.	7.1	2
3	Structural characterization of the EmrAB-TolC efflux complex from E. coli. Biochimica Et Biophysica Acta - Biomembranes, 2021, 1863, 183488.	2.6	17
4	Structure, Assembly, and Function of Tripartite Efflux and Type 1 Secretion Systems in Gram-Negative Bacteria. Chemical Reviews, 2021, 121, 5479-5596.	47.7	103
5	Allosteric drug transport mechanism of multidrug transporter AcrB. Nature Communications, 2021, 12, 3889.	12.8	41
6	Binding of Tetracyclines to Acinetobacter baumannii TetR Involves Two Arginines as Specificity Determinants. Frontiers in Microbiology, 2021, 12, 711158.	3.5	7
7	Structural and functional analysis of the promiscuous AcrB and AdeB efflux pumps suggests different drug binding mechanisms. Nature Communications, 2021, 12, 6919.	12.8	25
8	Characterization and Molecular Determinants for Î²-Lactam Specificity of the Multidrug Efflux Pump AcrD from Salmonella typhimurium. Antibiotics, 2021, 10, 1494.	3.7	4
9	AcrB: a mean, keen, drug efflux machine. Annals of the New York Academy of Sciences, 2020, 1459, 38-68.	3.8	99
10	Binding and Transport of Carboxylated Drugs by the Multidrug Transporter AcrB. Journal of Molecular Biology, 2020, 432, 861-877.	4.2	37
11	A novel method to determine antibiotic sensitivity in Bdellovibrio bacteriovorus reveals a DHFR-dependent natural trimethoprim resistance. Scientific Reports, 2020, 10, 5315.	3.3	12
12	Tigecycline efflux in Acinetobacter baumannii is mediated by TetA in synergy with RND-type efflux transporters. Journal of Antimicrobial Chemotherapy, 2020, 75, 1135-1139.	3.0	36
13	Antimicrobial Sensitivity Assay for Bdellovibrio bacteriovorus. Bio-protocol, 2020, 10, e3865.	0.4	0
14	The chloramphenicol/H+ antiporter CraA of Acinetobacter baumannii AYE reveals a broad substrate specificity. Journal of Antimicrobial Chemotherapy, 2019, 74, 1192-1201.	3.0	11
15	Identification of the novel class D Î²-lactamase OXA-679 involved in carbapenem resistance in Acinetobacter calcoaceticus. Journal of Antimicrobial Chemotherapy, 2019, 74, 1494-1502.	3.0	7
16	Identification and characterization of carbapenem binding sites within the RND-transporter AcrB. Biochimica Et Biophysica Acta - Biomembranes, 2019, 1861, 62-74.	2.6	18
17	A New Critical Conformational Determinant of Multidrug Efflux by an MFS Transporter. Journal of Molecular Biology, 2018, 430, 1368-1385.	4.2	27
18	High-Resolution Crystallographic Analysis of AcrB Using Designed Ankyrin Repeat Proteins (DARPins). Methods in Molecular Biology, 2018, 1700, 3-24.	0.9	3

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19	Bacterial efflux transporters in the limelight. Research in Microbiology, 2018, 169, 349-350.	2.1	0
20	Multidrug efflux pumps: structure, function and regulation. Nature Reviews Microbiology, 2018, 16, 523-539.	28.6	580
21	Crystal structure and mechanistic basis of a functional homolog of the antigen transporter TAP. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E438-E447.	7.1	67
22	Mechanisms of envelope permeability and antibiotic influx and efflux in Gram-negative bacteria. Nature Microbiology, 2017, 2, 17001.	13.3	238
23	Switch Loop Flexibility Affects Substrate Transport of the AcrB Efflux Pump. Journal of Molecular Biology, 2017, 429, 3863-3874.	4.2	33
24	Dynamics of Intact MexAB-OprM Efflux Pump: Focusing on the MexA-OprM Interface. Scientific Reports, 2017, 7, 16521.	3.3	30
25	Cytochrome c Oxidase Biogenesis and Metallochaperone Interactions: Steps in the Assembly Pathway of a Bacterial Complex. PLoS ONE, 2017, 12, e0170037.	2.5	23
26	Editorial: Bad Bugs in the XXIst Century: Resistance Mediated by Multi-Drug Efflux Pumps in Gram-Negative Bacteria. Frontiers in Microbiology, 2016, 7, 833.	3.5	26
27	BGA66 and BGA71 facilitate complement resistance of Borrelia bavariensis by inhibiting assembly of the membrane attack complex. Molecular Microbiology, 2016, 99, 407-424.	2.5	63
28	Transport of lipophilic carboxylates is mediated by transmembrane helix 2 in multidrug transporter AcrB. Nature Communications, 2016, 7, 13819.	12.8	51
29	Tripartite assembly of RND multidrug efflux pumps. Nature Communications, 2016, 7, 10731.	12.8	166
30	Molecular basis for inhibition of AcrB multidrug efflux pump by novel and powerful pyranopyridine derivatives. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 3509-3514.	7.1	172
31	Molecular basis of polyspecificity of the Small Multidrug Resistance Efflux Pump AbeS from Acinetobacter baumannii. Journal of Molecular Biology, 2016, 428, 644-657.	4.2	24
32	Approved Drugs Containing Thiols as Inhibitors of Metallo-Î²-lactamases: Strategy To Combat Multidrug-Resistant Bacteria. Journal of Medicinal Chemistry, 2015, 58, 3626-3630.	6.4	127
33	Structure, mechanism and cooperation of bacterial multidrug transporters. Current Opinion in Structural Biology, 2015, 33, 76-91.	5.7	129
34	The assembly and disassembly of the AcrAB-TolC three-component multidrug efflux pump. Biological Chemistry, 2015, 396, 1083-1089.	2.5	23
35	Coupling of remote alternating-access transport mechanisms for protons and substrates in the multidrug efflux pump AcrB. ELife, 2014, 3, .	6.0	137
36	Switch-Loop Flexibility Affects Transport of Large Drugs by the Promiscuous AcrB Multidrug Efflux Transporter. Antimicrobial Agents and Chemotherapy, 2014, 58, 4767-4772.	3.2	52

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37	Detecting Substrates Bound to the Secondary Multidrug Efflux Pump EmrE by DNP-Enhanced Solid-State NMR. Journal of the American Chemical Society, 2013, 135, 15754-15762.	13.7	51
38	The Outer Membrane TolC-like Channel HgdD Is Part of Tripartite Resistance-Nodulation-Cell Division (RND) Efflux Systems Conferring Multiple-drug Resistance in the Cyanobacterium Anabaena sp. PCC7120. Journal of Biological Chemistry, 2013, 288, 31192-31205.	3.4	22
39	RND Efflux Pumps: Structural Information Translated into Function and Inhibition Mechanisms. Current Topics in Medicinal Chemistry, 2013, 13, 3079-3100.	2.1	122
40	Transport of drugs by the multidrug transporter AcrB involves an access and a deep binding pocket that are separated by a switch-loop. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 5687-5692.	7.1	285
41	Effect of the F610A Mutation on Substrate Extrusion in the AcrB Transporter: Explanation and Rationale by Molecular Dynamics Simulations. Journal of the American Chemical Society, 2011, 133, 10704-10707.	13.7	79
42	A natural prodrug activation mechanism in nonribosomal peptide synthesis. Nature Chemical Biology, 2011, 7, 888-890.	8.0	118
43	Analysis of AcrB and AcrB/DARPin ligand complexes by LILBID MS. Biochimica Et Biophysica Acta - Biomembranes, 2011, 1808, 2189-2196.	2.6	20
44	Drug Resistance: A Periplasmic MÃ©nage Ã Trois. Chemistry and Biology, 2011, 18, 405-407.	6.0	4
45	The use of novel organic gels and hydrogels in protein crystallization. Journal of Applied Crystallography, 2010, 43, 58-63.	4.5	18
46	Trinity revealed: Stoichiometric complex assembly of a bacterial multidrug efflux pump. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 6893-6894.	7.1	18
47	Structural and functional aspects of the multidrug efflux pump AcrB. Biological Chemistry, 2009, 390, 693-699.	2.5	50
48	Drug transport mechanism of the AcrB efflux pump. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2009, 1794, 782-793.	2.3	256
49	Crucial Role of Asp408 in the Proton Translocation Pathway of Multidrug Transporter AcrB: Evidence from Site-Directed Mutagenesis and Carbodiimide Labeling. Biochemistry, 2009, 48, 5801-5812.	2.5	74
50	Engineered disulfide bonds support the functional rotation mechanism of multidrug efflux pump AcrB. Nature Structural and Molecular Biology, 2008, 15, 199-205.	8.2	142
51	Site-Directed Mutagenesis Reveals Putative Substrate Binding Residues in the Escherichia coli RND Efflux Pump AcrB. Journal of Bacteriology, 2008, 190, 8225-8229.	2.2	126
52	Molecular Analysis of BcrR, a Membrane-bound Bacitracin Sensor and DNA-binding Protein from Enterococcus faecalis. Journal of Biological Chemistry, 2008, 283, 8591-8600.	3.4	23
53	The AcrB Efflux Pump: Conformational Cycling and Peristalsis Lead to Multidrug Resistance. Current Drug Targets, 2008, 9, 729-749.	2.1	116
54	Functional characterization of a NapA Na+/H+antiporter fromThermus thermophilus. FEBS Letters, 2007, 581, 572-578.	2.8	17

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55	Structural Asymmetry of AcrB Trimer Suggests a Peristaltic Pump Mechanism. Science, 2006, 313, 1295-1298.	12.6	512
56	Basolateral aromatic amino acid transporter TAT1 (Slc16a10) functions as an efflux pathway. Journal of Cellular Physiology, 2006, 206, 771-779.	4.1	81
57	Identification of a domain in the Î±-subunit of the oxaloacetate decarboxylase Na+ pump that accomplishes complex formation with the Î³-subunit. FEBS Journal, 2005, 272, 846-855.	4.7	16
58	Oxaloacetate decarboxylase of Archaeoglobus fulgidus: cloning of genes and expression in Escherichia coli. Archives of Microbiology, 2004, 182, 414-420.	2.2	7
59	Crystallographic analysis of AcrB. FEBS Letters, 2004, 564, 333-339.	2.8	36
60	An antibody library for stabilizing and crystallizing membrane proteins - selecting binders to the citrate carrier CitS. FEBS Letters, 2004, 564, 340-348.	2.8	43
61	Characterization of the citrate/acetate antiporter CitW of Klebsiella pneumoniae. Archives of Microbiology, 2002, 177, 500-506.	2.2	19
62	Purification, crystallization and preliminary diffraction studies of AcrB, an inner-membrane multi-drug efflux protein. Acta Crystallographica Section D: Biological Crystallography, 2002, 58, 1865-1867.	2.5	40
63	Role of conserved residues within helices IV and VIII of the oxaloacetate decarboxylase Î²â€Šsubunit in the energy coupling mechanism of the Na+pump. FEBS Journal, 2002, 269, 2997-3004.	0.2	17
64	The Na + -dependent citrate carrier of Klebsiella pneumoniae : high-level expression and site-directed mutagenesis of asparagine-185 and glutamate-194. Archives of Microbiology, 2000, 174, 67-73.	2.2	15
65	Identification of an Na ⁺ -Dependent Malonate Transporter of Malonomonas rubra and Its Dependence on Two Separate Genes. Journal of Bacteriology, 1998, 180, 2689-2693.	2.2	14
66	The Escherichia coli Citrate Carrier CitT: a Member of a Novel Eubacterial Transporter Family Related to the 2-Oxoglutarate/Malate Translocator from Spinach Chloroplasts. Journal of Bacteriology, 1998, 180, 4160-4165.	2.2	110
67	Purification of two active fusion proteins of the Na+-dependent citrate carrier ofKlebsiella pneumoniae. FEBS Letters, 1994, 347, 37-41.	2.8	33
68	Localization of Peptidases in Lactococci. Applied and Environmental Microbiology, 1992, 58, 285-290.	3.1	96