Juquan Jiang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	N-type fast inactivation of a eukaryotic voltage-gated sodium channel. Nature Communications, 2022, 13, 2713.	5.8	8
2	Structural basis of ligand binding modes of human EAAT2. Nature Communications, 2022, 13, .	5.8	12
3	A novel three-TMH Na+/H+ antiporter and the functional role of its oligomerization. Journal of Molecular Biology, 2021, 433, 166730.	2.0	11
4	A Novel MFS-MDR Transporter, MdrP, Employs D223 as a Key Determinant in the Na+ Translocation Coupled to Norfloxacin Efflux. Frontiers in Microbiology, 2020, 11, 955.	1.5	2
5	Polar or Charged Residues Located in Four Highly Conserved Motifs Play a Vital Role in the Function or pH Response of a UPF0118 Family Na+(Li+)/H+ Antiporter. Frontiers in Microbiology, 2020, 11, 841.	1.5	8
6	Implications for Cation Selectivity and Evolution by a Novel Cation Diffusion Facilitator Family Member From the Moderate Halophile Planococcus dechangensis. Frontiers in Microbiology, 2019, 10, 607.	1.5	10
7	Characterization of a Functionally Unknown Arginine–Aspartate–Aspartate Family Protein From Halobacillus andaensis and Functional Analysis of Its Conserved Arginine/Aspartate Residues. Frontiers in Microbiology, 2018, 9, 807.	1.5	15
8	An Uncharacterized Major Facilitator Superfamily Transporter From Planococcus maritimus Exhibits Dual Functions as a Na+(Li+, K+)/H+ Antiporter and a Multidrug Efflux Pump. Frontiers in Microbiology, 2018, 9, 1601.	1.5	21
9	A novel NhaD-type Na ⁺ /H ⁺ antiporter from the moderate halophile and alkaliphile <i>Halomonas alkaliphila</i> . Canadian Journal of Microbiology, 2017, 63, 596-607.	0.8	20
10	A UPF0118 family protein with uncharacterized function from the moderate halophile Halobacillus andaensis represents a novel class of Na+(Li+)/H+ antiporter. Scientific Reports, 2017, 7, 45936.	1.6	20
11	Characterization of a novel two-component Na+(Li+, K+)/H+ antiporter from Halomonas zhaodongensis. Scientific Reports, 2017, 7, 4221.	1.6	20
12	Novosphingobium oryzae sp. nov., a potential plant-promoting endophytic bacterium isolated from rice roots. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 302-307.	0.8	32
13	Pseudomonas songnenensis sp. nov., isolated from saline and alkaline soils in Songnen Plain, China. Antonie Van Leeuwenhoek, 2015, 107, 711-721.	0.7	16
14	Pseudomonas zhaodongensis sp. nov., isolated from saline and alkaline soils. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 1022-1030.	0.8	22
15	Planococcus dechangensis sp. nov., a moderately halophilic bacterium isolated from saline and alkaline soils in Dechang Township, Zhaodong City, China. Antonie Van Leeuwenhoek, 2015, 107, 1075-1083.	0.7	17
16	Halobacillus andaensis sp. nov., a moderately halophilic bacterium isolated from saline and alkaline soil. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 1908-1914.	0.8	23
17	Kocuria dechangensis sp. nov., an actinobacterium isolated from saline and alkaline soils. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 3024-3030.	0.8	15
18	Cloning and identification of Group 1 mrp operon encoding a novel monovalent cation/proton antiporter system from the moderate halophile Halomonas zhaodongensis. Extremophiles, 2014, 18, 963-972.	0.9	18

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19	Cloning and identification of a novel NhaD-type Na+/H+ antiporter from metagenomic DNA of the halophilic bacteria in soil samples around Daban Salt Lake. Extremophiles, 2014, 18, 89-98.	0.9	16
20	Halomonas songnenensis sp. nov., a moderately halophilic bacterium isolated from saline and alkaline soils. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1662-1669.	0.8	36
21	Halomonas zhaodongensis sp. nov., a slightly halophilic bacterium isolated from saline–alkaline soils in Zhaodong, China. Antonie Van Leeuwenhoek, 2013, 104, 685-694.	0.7	18
22	Identification of important charged residues for alkali cation exchange or pH regulation of NhaH, a Na+/H+ antiporter of Halobacillus dabanensis. Biochimica Et Biophysica Acta - Biomembranes, 2013, 1828, 997-1003.	1.4	14
23	Putative paired small multidrug resistance family proteins <i>PsmrAB</i> , the homolog of YvdSR, actually function as a novel two-component Na ⁺ /H ⁺ antiporter. FEMS Microbiology Letters, 2013, 338, 31-38.	0.7	19
24	Metabolism of [¹³ C ₅]hydroxyproline in vitro and in vivo: implications for primary hyperoxaluria. American Journal of Physiology - Renal Physiology, 2012, 302, G637-G643.	1.6	25
25	Oxalate and Sucralose Absorption in Idiopathic Calcium Oxalate Stone Formers. Urology, 2011, 78, 475.e9-475.e13.	0.5	16
26	Oral Antibiotic Treatment of Helicobacter pylori Leads to Persistently Reduced Intestinal Colonization Rates with Oxalobacter formigenes. Journal of Endourology, 2011, 25, 1781-1785.	1.1	55
27	Impact of Dietary Calcium and Oxalate, and Oxalobacter Formigenes Colonization on Urinary Oxalate Excretion. Journal of Urology, 2011, 186, 135-139.	0.2	92
28	A primary sodium pump gene of the moderate halophile Halobacillus dabanensis exhibits secondary antiporter properties. Biochemical and Biophysical Research Communications, 2006, 346, 612-617.	1.0	14
29	A Na+/H+antiporter gene of the moderately halophilic bacteriumHalobacillus dabanensisD-8T: cloning and molecular characterization. FEMS Microbiology Letters, 2006, 255, 89-95.	0.7	36
30	Thepha2gene cluster involved in Na+resistance and adaption to alkaline pH inSinorhizobium frediiRT19 encodes a monovalent cation/proton antiporter. FEMS Microbiology Letters, 2006, 262, 172-177.	0.7	20
31	Hydroxyproline ingestion and urinary oxalate and glycolate excretion. Kidney International, 2006, 70, 1929-1934.	2.6	134
32	Isolation of salt-sensitive mutants from Sinorhizobium meliloti and characterization of genes involved in salt tolerance. Letters in Applied Microbiology, 2004, 39, 278-283.	1.0	51
33	Salt-tolerance genes involved in cation efflux and osmoregulation ofSinorhizobium frediiRT19 detected by isolation and characterization of Tn5 mutants. FEMS Microbiology Letters, 2004, 239, 139-146.	0.7	32