

Alessio Accardi

List of Publications by Year in descending order

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44
papers

3,367
citations

218381

26
h-index

301761

39
g-index

51
all docs

51
docs citations

51
times ranked

2424
citing authors

#	ARTICLE	IF	CITATIONS
1	TMEM16 scramblases thin the membrane to enable lipid scrambling. <i>Nature Communications</i> , 2022, 13, 2604.	5.8	22
2	Not so transport incompetent after all: Revisiting a CLC-7 mutant sheds new mechanistic light on lysosomal physiology. <i>Journal of General Physiology</i> , 2021, 153, .	0.9	1
3	A quantitative flux assay for the study of reconstituted Cl ⁻ channels and transporters. <i>Methods in Enzymology</i> , 2021, 652, 243-272.	0.4	2
4	Membrane lipids are both the substrates and a mechanistically responsive environment of TMEM16 scramblase proteins. <i>Journal of Computational Chemistry</i> , 2020, 41, 538-551.	1.5	15
5	What biologists want from their chloride reporters – a conversation between chemists and biologists. <i>Journal of Cell Science</i> , 2020, 133, .	1.2	27
6	Reconstitution of Proteoliposomes for Phospholipid Scrambling and Nonselective Channel Assays. <i>Methods in Molecular Biology</i> , 2020, 2127, 207-225.	0.4	10
7	Divergent Cl ⁻ and H ⁺ pathways underlie transport coupling and gating in CLC exchangers and channels. <i>ELife</i> , 2020, 9, .	2.8	17
8	Structural Basis of Lipid Scrambling and Ion Conduction by TMEM16 Scramblases. <i>FASEB Journal</i> , 2020, 34, 1-1.	0.2	0
9	Dynamic modulation of the lipid translocation groove generates a conductive ion channel in Ca ²⁺ -bound nhTMEM16. <i>Nature Communications</i> , 2019, 10, 4972.	5.8	23
10	The structural basis of lipid scrambling and inactivation in the endoplasmic reticulum scramblase TMEM16K. <i>Nature Communications</i> , 2019, 10, 3956.	5.8	101
11	Structural basis of Ca ²⁺ -dependent activation and lipid transport by a TMEM16 scramblase. <i>ELife</i> , 2019, 8, .	2.8	87
12	Known structures and unknown mechanisms of TMEM16 scramblases and channels. <i>Journal of General Physiology</i> , 2018, 150, 933-947.	0.9	92
13	Gating mechanism of the extracellular entry to the lipid pathway in a TMEM16 scramblase. <i>Nature Communications</i> , 2018, 9, 3251.	5.8	70
14	Out-of-the-groove transport of lipids by TMEM16 and GPCR scramblases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E7033-E7042.	3.3	49
15	Probing the conformation of a conserved glutamic acid within the Cl ⁻ pathway of a CLC H ⁺ /Cl ⁻ exchanger. <i>Journal of General Physiology</i> , 2017, 149, 523-529.	0.9	18
16	The nhTMEM16 Scramblase Is Also a Nonselective Ion Channel. <i>Biophysical Journal</i> , 2016, 111, 1919-1924.	0.2	70
17	Structure and gating of CLC channels and exchangers. <i>Journal of Physiology</i> , 2015, 593, 4129-4138.	1.3	65
18	A Proteoliposome-Based Efflux Assay to Determine Single-molecule Properties of Cl ⁻ Channels and Transporters. <i>Journal of Visualized Experiments</i> , 2015, . .	0.2	4

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19	Lipids link ion channels and cancer. <i>Science</i> , 2015, 349, 789-790.	6.0	31
20	TMEM16 Proteins: Unknown Structure and Confusing Functions. <i>Journal of Molecular Biology</i> , 2015, 427, 94-105.	2.0	115
21	Unveiling the Secret Lives of Glutamate Transporters: VGLUTs Engage in Multiple Transport Modes. <i>Neuron</i> , 2014, 84, 1110-1112.	3.8	1
22	Nucleotide binding triggers a conformational change of the CBS module of the magnesium transporter CNNM2 from a twisted towards a flat structure. <i>Biochemical Journal</i> , 2014, 464, 23-34.	1.7	41
23	Conformational changes required for H ⁺ /Cl ⁻ exchange mediated by a CLC transporter. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 456-463.	3.6	76
24	Ca ²⁺ -dependent phospholipid scrambling by a reconstituted TMEM16 ion channel. <i>Nature Communications</i> , 2013, 4, 2367.	5.8	202
25	Purified TMEM16A is sufficient to form Ca ²⁺ -activated Cl ⁻ channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 19354-19359.	3.3	97
26	Synergistic substrate binding determines the stoichiometry of transport of a prokaryotic H ⁺ /Cl ⁻ exchanger. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 525-531.	3.6	71
27	3PT118 Functional in vitro reconstitution of Calcium-activated Chloride channel, TMEM16A/anoctamin1(The 50th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuru</i> , 2012, 52, S160.	0.0	0
28	Proton Transport and Conformational Changes in H ⁺ /CL ⁻ Exchangers. <i>Biophysical Journal</i> , 2011, 100, 245a.	0.2	0
29	Proton block of the CLC-5 Cl ⁻ /H ⁺ exchanger. <i>Journal of General Physiology</i> , 2010, 135, 653-659.	0.9	28
30	CLC channels and transporters: Proteins with borderline personalities. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2010, 1798, 1457-1464.	1.4	87
31	Proton Block of Human CLC-5 Cl ⁻ /H ⁺ Exchanger. <i>Biophysical Journal</i> , 2010, 98, 320a.	0.2	0
32	Basis of substrate binding and conservation of selectivity in the CLC family of channels and transporters. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 1294-1301.	3.6	106
33	Ion permeation through a Cl ⁻ -selective channel designed from a CLC Cl ⁻ /H ⁺ exchanger. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 11194-11199.	3.3	97
34	To ATP or Not To ATP: This Is the Question. <i>Journal of General Physiology</i> , 2008, 131, 105-108.	0.9	4
35	Uncoupling and Turnover in a Cl ⁻ /H ⁺ Exchange Transporter. <i>Journal of General Physiology</i> , 2007, 129, 317-329.	0.9	131
36	Synergism Between Halide Binding and Proton Transport in a CLC-type Exchanger. <i>Journal of Molecular Biology</i> , 2006, 362, 691-699.	2.0	103

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37	Structure and Function of CLC Chloride Channels and Transporters. <i>Advances in Molecular and Cell Biology</i> , 2006, , 59-82.	0.1	0
38	Separate Ion Pathways in a Cl ⁻ /H ⁺ Exchanger. <i>Journal of General Physiology</i> , 2005, 126, 563-570.	0.9	203
39	Ionic Currents Mediated by a Prokaryotic Homologue of CLC Cl ⁻ Channels. <i>Journal of General Physiology</i> , 2004, 123, 109-119.	0.9	138
40	Secondary active transport mediated by a prokaryotic homologue of CLC Cl ⁻ channels. <i>Nature</i> , 2004, 427, 803-807.	13.7	602
41	Conservation of Chloride Channel Structure Revealed by an Inhibitor Binding Site in CLC-1. <i>Neuron</i> , 2003, 38, 47-59.	3.8	161
42	Conformational Changes in the Pore of CLC-0. <i>Journal of General Physiology</i> , 2003, 122, 277-294.	0.9	82
43	A biological role for prokaryotic CLC chloride channels. <i>Nature</i> , 2002, 419, 715-718.	13.7	204
44	Fast and Slow Gating Relaxations in the Muscle Chloride Channel Clc-1. <i>Journal of General Physiology</i> , 2000, 116, 433-444.	0.9	101