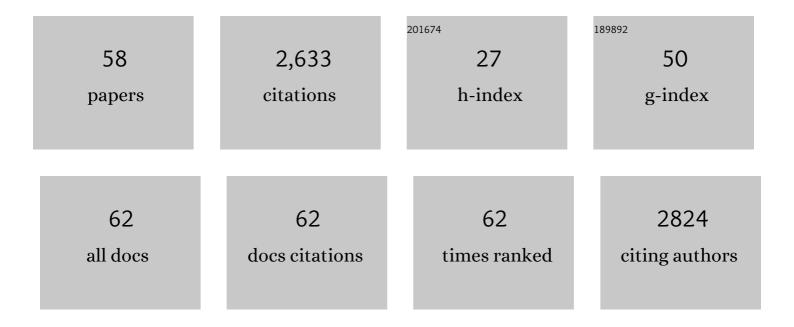
Christoph von Ballmoos

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Rapid Estimation of Membrane Protein Orientation in Liposomes. ChemBioChem, 2022, 23, .	2.6	6
2	Functional design of bacterial superoxide:quinone oxidoreductase. Biochimica Et Biophysica Acta - Bioenergetics, 2022, 1863, 148583.	1.0	2
3	Biochemical consequences of two clinically relevant ND-gene mutations in Escherichia coli respiratory complexÂl. Scientific Reports, 2021, 11, 12641.	3.3	6
4	The missing enzymatic link in syntrophic methane formation from fatty acids. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	7
5	Energy transfer between the nicotinamide nucleotide transhydrogenase and ATP synthase of Escherichia coli. Scientific Reports, 2021, 11, 21234.	3.3	4
6	CD31 (PECAM-1) Serves as the Endothelial Cell-Specific Receptor of Clostridium perfringens β-Toxin. Cell Host and Microbe, 2020, 28, 69-78.e6.	11.0	28
7	Bifunctional DNA Duplexes Permit Efficient Incorporation of pH Probes into Liposomes. ChemBioChem, 2020, 21, 2219-2224.	2.6	3
8	Kinetic coupling of the respiratory chain with ATP synthase, but not proton gradients, drives ATP production in cristae membranes. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 2412-2421.	7.1	52
9	Current problems and future avenues in proteoliposome research. Biochemical Society Transactions, 2020, 48, 1473-1492.	3.4	29
10	The proton pumping bo oxidase from Vitreoscilla. Scientific Reports, 2019, 9, 4766.	3.3	7
11	ATP synthesis at physiological nucleotide concentrations. Scientific Reports, 2019, 9, 3070.	3.3	31
12	Towards a Synthetic Mitochondrion. Chimia, 2018, 72, 291.	0.6	13
13	Scavenging of superoxide by a membrane-bound superoxide oxidase. Nature Chemical Biology, 2018, 14, 788-793.	8.0	71
14	Dissecting the proton transport pathway in electrogenic Na ⁺ /H ⁺ antiporters. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E1101-E1110.	7.1	43
15	The lateral distance between a proton pump and ATP synthase determines the ATP-synthesis rate. Scientific Reports, 2017, 7, 2926.	3.3	41
16	Splitting of the O–O bond at the heme-copper catalytic site of respiratory oxidases. Science Advances, 2017, 3, e1700279.	10.3	50
17	Activation of Proton Translocation by Respiratory Complex I. Biochemistry, 2017, 56, 5691-5697.	2.5	13
18	Delivery of membrane proteins into small and giant unilamellar vesicles by chargeâ€mediated fusion. FEBS Letters, 2016, 590, 2051-2062.	2.8	41

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19	Lipid-mediated Protein-protein Interactions Modulate Respiration-driven ATP Synthesis. Scientific Reports, 2016, 6, 24113.	3.3	38
20	Regulatory role of the respiratory supercomplex factors in <i>Saccharomyces cerevisiae</i> . Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E4476-85.	7.1	45
21	Isolation of yeast complex IV in native lipid nanodiscs. Biochimica Et Biophysica Acta - Biomembranes, 2016, 1858, 2984-2992.	2.6	45
22	Rapid Electron Transfer within the III-IV Supercomplex in Corynebacterium glutamicum. Scientific Reports, 2016, 6, 34098.	3.3	20
23	Mimicking respiratory phosphorylation using purified enzymes. Biochimica Et Biophysica Acta - Bioenergetics, 2016, 1857, 321-331.	1.0	40
24	Mutation of a single residue in the <i>ba</i> ₃ oxidase specifically impairs protonation of the pump site. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 3397-3402.	7.1	23
25	Effect of lipid bilayer properties on the photocycle of green proteorhodopsin. Biochimica Et Biophysica Acta - Bioenergetics, 2015, 1847, 698-708.	1.0	17
26	Isolation, N-glycosylations and Function of a Hyaluronidase-Like Enzyme from the Venom of the Spider Cupiennius salei. PLoS ONE, 2015, 10, e0143963.	2.5	23
27	Crystal structure of the sodium–proton antiporter NhaA dimer and new mechanistic insights. Journal of General Physiology, 2014, 144, 529-544.	1.9	79
28	SNARE-fusion mediated insertion of membrane proteins into native and artificial membranes. Nature Communications, 2014, 5, 4303.	12.8	26
29	A two-domain elevator mechanism for sodium/proton antiport. Nature, 2013, 501, 573-577.	27.8	221
30	Single Mutations That Redirect Internal Proton Transfer in the <i>ba</i> ₃ Oxidase from <i>Thermus thermophilus</i> . Biochemistry, 2013, 52, 7022-7030.	2.5	6
31	An alternative role of FoF1-ATP synthase in Escherichia coli: synthesis of thiamine triphosphate. Scientific Reports, 2013, 3, 1071.	3.3	19
32	Selective and ATP-driven transport of ions across supported membranes into nanoporous carriers using gramicidin A and ATP synthase. Physical Chemistry Chemical Physics, 2013, 15, 2733.	2.8	9
33	The Membrane Modulates Internal Proton Transfer in Cytochrome <i>c</i> Oxidase. Biochemistry, 2012, 51, 1092-1100.	2.5	19
34	Timing of Electron and Proton Transfer in the <i>ba</i> ₃ Cytochrome <i>c</i> Oxidase from <i>Thermus thermophilus</i> . Biochemistry, 2012, 51, 4507-4517.	2.5	15
35	Reconstitution of respiratory oxidases in membrane nanodiscs for investigation of protonâ€coupled electron transfer. FEBS Letters, 2012, 586, 640-645.	2.8	21
36	Proton transfer in ba3 cytochrome c oxidase from Thermus thermophilus. Biochimica Et Biophysica Acta - Bioenergetics, 2012, 1817, 650-657.	1.0	52

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37	Kinetic design of the respiratory oxidases. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 11057-11062.	7.1	36
38	Functional Role of Thr-312 and Thr-315 in the Proton-Transfer Pathway in <i>ba</i> ₃ Cytochrome <i>c</i> Oxidase from <i>Thermus thermophilus</i> . Biochemistry, 2010, 49, 7033-7039.	2.5	18
39	Functional asymmetry of the F ₀ motor in bacterial ATP synthases. Molecular Microbiology, 2009, 72, 479-490.	2.5	27
40	Essentials for ATP Synthesis by F ₁ F ₀ ATP Synthases. Annual Review of Biochemistry, 2009, 78, 649-672.	11.1	326
41	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
42	Arginineâ€induced conformational change in the <i>c</i> â€ring/ <i>a</i> â€subunit interface of ATP synthase. FEBS Journal, 2008, 275, 2137-2150.	4.7	25
43	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
44	Unique Rotary ATP Synthase and Its Biological Diversity. Annual Review of Biophysics, 2008, 37, 43-64.	10.0	167
45	l̃"Γ̈́ and l̃"pH are equivalent driving forces for proton transport through isolated FO complexes of ATP synthases. Biochimica Et Biophysica Acta - Bioenergetics, 2008, 1777, 1301-1310.	1.0	48
46	Two Distinct Proton Binding Sites in the ATP Synthase Family. Biochemistry, 2007, 46, 11800-11809.	2.5	54
47	ATP Synthesis by Decarboxylation Phosphorylation. Results and Problems in Cell Differentiation, 2007, 45, 153-184.	0.7	25
48	Alternative proton binding mode in ATP synthases. Journal of Bioenergetics and Biomembranes, 2007, 39, 441-445.	2.3	15
49	Catalytic and mechanical cycles in Fâ€ATP synthases. EMBO Reports, 2006, 7, 276-282.	4.5	119
50	The ion channel of F-ATP synthase is the target of toxic organotin compounds. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 11239-11244.	7.1	88
51	A continuous fluorescent method for measuring Na+ transport. Analytical Biochemistry, 2004, 335, 334-337.	2.4	9
52	Torque Generation by the Fo motor of the Sodium ATPase. Biophysical Journal, 2004, 87, 2148-2163.	0.5	49
53	Electrical Power Fuels Rotary ATP Synthase. Structure, 2003, 11, 1469-1473.	3.3	45
54	Complete DNA sequence of the atp operon of the sodium-dependent F1Fo ATP synthase from Ilyobacter tartaricus and identification of the encoded subunits. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2003, 1625, 221-226.	2.4	7

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55	Evidence for Structural Integrity in the Undecameric c-Rings Isolated from Sodium ATP Synthases. Journal of Molecular Biology, 2003, 325, 389-397.	4.2	80
56	Purification and Biochemical Characterization of the F 1 F o -ATP Synthase from Thermoalkaliphilic Bacillus sp. Strain TA2.A1. Journal of Bacteriology, 2003, 185, 4442-4449.	2.2	59
57	Membrane Topography of the Coupling Ion Binding Site in Na+-translocating F1F0 ATP Synthase. Journal of Biological Chemistry, 2002, 277, 3504-3510.	3.4	36
58	Membrane embedded location of Na+ or H+ binding sites on the rotor ring of F1 F0 ATP synthases. FEBS Journal, 2002, 269, 5581-5589.	0.2	19