

Flemming Cornelius

List of Publications by Year in descending order

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92
papers

3,798
citations

136740

32
h-index

133063

59
g-index

93
all docs

93
docs citations

93
times ranked

2680
citing authors

#	ARTICLE	IF	CITATIONS
1	Crystal structure of the sodium-potassium pump at 2.4-Å resolution. <i>Nature</i> , 2009, 459, 446-450.	13.7	557
2	Crystal structure of the sodium-potassium pump (Na ⁺ ,K ⁺ -ATPase) with bound potassium and ouabain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 13742-13747.	3.3	298
3	Crystal structure of a Na ⁺ -bound Na ⁺ ,K ⁺ -ATPase preceding the E1P state. <i>Nature</i> , 2013, 502, 201-206.	13.7	271
4	Modulation of Na ⁺ ,K ⁺ -ATPase and Na ⁺ -ATPase Activity by Phospholipids and Cholesterol. I. Steady-State Kinetics. <i>Biochemistry</i> , 2001, 40, 8842-8851.	1.2	166
5	Reversible Oxidative Modification. <i>Circulation Research</i> , 2009, 105, 185-193.	2.0	147
6	General and specific lipid-protein interactions in Na ⁺ ,K ⁺ -ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 1729-1743.	1.4	120
7	Functional reconstitution of the sodium pump. Kinetics of exchange reactions performed by reconstituted Na ⁺ /K ⁺ -ATPase. <i>BBA - Biomembranes</i> , 1991, 1071, 19-66.	7.9	112
8	Identification of a Phospholemman-like Protein from Shark Rectal Glands. <i>Journal of Biological Chemistry</i> , 2000, 275, 35969-35977.	1.6	108
9	Reconstitution of (Na ⁺ + K ⁺)-ATPase into phospholipid vesicles with full recovery of its specific activity. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1984, 772, 357-373.	1.4	104
10	First Crystal Structures of Na ⁺ ,K ⁺ -ATPase: New Light on the Oldest Ion Pump. <i>Structure</i> , 2011, 19, 1732-1738.	1.6	102
11	FXFD Proteins Reverse Inhibition of the Na ⁺ -K ⁺ Pump Mediated by Glutathionylation of Its Î²1 Subunit. <i>Journal of Biological Chemistry</i> , 2011, 286, 18562-18572.	1.6	79
12	Modulation of Na ⁺ ,K ⁺ -ATPase by Phospholipids and Cholesterol. II. Steady-State and Presteady-State Kinetics. <i>Biochemistry</i> , 2003, 42, 8541-8549.	1.2	69
13	Na ⁺ -Na ⁺ exchange mediated by (Na ⁺ + K ⁺)-ATPase reconstituted into liposomes. Evaluation of pump stoichiometry and response to ATP and ADP. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1985, 818, 211-221.	1.4	62
14	A Structural View on the Functional Importance of the Sugar Moiety and Steroid Hydroxyls of Cardiotonic Steroids in Binding to Na ⁺ ,K ⁺ -ATPase*. <i>Journal of Biological Chemistry</i> , 2013, 288, 6602-6616.	1.6	61
15	Regulation of Na ⁺ ,K ⁺ -ATPase by PLMS, the Phospholemman-like Protein from Shark. <i>Journal of Biological Chemistry</i> , 2003, 278, 37427-37438.	1.6	59
16	New crystal structures of PII-type ATPases: excitement continues. <i>Current Opinion in Structural Biology</i> , 2013, 23, 507-514.	2.6	58
17	Rate Limitation of the Na ⁺ ,K ⁺ -ATPase Pump Cycle. <i>Biophysical Journal</i> , 2001, 81, 2069-2081.	0.2	57
18	Cholesterol modulation of molecular activity of reconstituted shark Na ⁺ ,K ⁺ -ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1995, 1235, 205-212.	1.4	50

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19	Tonic contraction and the control of relaxation in a chemically skinned molluscan smooth muscle.. Journal of General Physiology, 1982, 79, 821-834.	0.9	45
20	Susceptibility of $\hat{1}^{21}$ Na ⁺ -K ⁺ Pump Subunit to Glutathionylation and Oxidative Inhibition Depends on Conformational State of Pump. Journal of Biological Chemistry, 2012, 287, 12353-12364.	1.6	43
21	Functional regulation of reconstituted Na, K-ATPase by protein kinase A phosphorylation. FEBS Letters, 1996, 380, 277-280.	1.3	40
22	E2P Phosphoforms of Na,K-ATPase. I. Comparison of Phosphointermediates Formed from ATP and Pi by Their Reactivity toward Hydroxylamine and Vanadate. Biochemistry, 1998, 37, 13634-13642.	1.2	40
23	Cholesterol-Dependent Interaction of Polyunsaturated Phospholipids with Na,K-ATPase. Biochemistry, 2008, 47, 1652-1658.	1.2	40
24	Fluorescent styryl dyes as probes for Na,K-ATPase reaction mechanism: significance of the charge of the hydrophilic moiety of RH dyes. Biochemistry, 1995, 34, 16806-16814.	1.2	39
25	Mechanism of the Rate-Determining Step of the Na ⁺ ,K ⁺ -ATPase Pump Cycle. Biochemistry, 2002, 41, 9496-9507.	1.2	36
26	Functional Modulation of the Sodium Pump: The Regulatory Proteins. Physiology, 2003, 18, 119-124.	1.6	36
27	Intrinsic reaction-cycle time scale of Na ⁺ ,K ⁺ -ATPase manifests itself in the lipid-protein interactions of nonequilibrium membranes. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18442-18446.	3.3	36
28	Spatial distribution and activity of Na ⁺ /K ⁺ -ATPase in lipid bilayer membranes with phase boundaries. Biochimica Et Biophysica Acta - Biomembranes, 2016, 1858, 1390-1399.	1.4	36
29	E2P Phosphoforms of Na,K-ATPase. II. Interaction of Substrate and Cation-Binding Sites in Pi Phosphorylation of Na,K-ATPase. Biochemistry, 1998, 37, 16686-16696.	1.2	35
30	Metal Fluoride Complexes of Na,K-ATPase. Journal of Biological Chemistry, 2011, 286, 29882-29892.	1.6	35
31	Binding of cardiotonic steroids to Na ⁺ ,K ⁺ -ATPase in the E2P state. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	35
32	Protein Kinase C Phosphorylation of Purified Na,K-ATPase: C-Terminal Phosphorylation Sites at the $\hat{1}^{\pm}$ - and $\hat{1}^3$ -Subunits Close to the Inner Face of the Plasma Membrane. Biophysical Journal, 2002, 82, 1907-1919.	0.2	34
33	The regulation of tension in a chemically skinned molluscan smooth muscle: effect of Mg ²⁺ on the Ca ²⁺ -activated tension generation.. Journal of General Physiology, 1980, 75, 709-725.	0.9	32
34	Modulation of Na,K-ATPase by associated small transmembrane regulatory proteins and by lipids. Journal of Bioenergetics and Biomembranes, 2001, 33, 415-423.	1.0	29
35	[15] Incorporation of C12E8-solubilized Na ⁺ ,K ⁺ -ATPase into liposomes: determination of sidedness and orientation. Methods in Enzymology, 1988, 156, 156-167.	0.4	28
36	Uncoupled Na ⁺ -efflux on reconstituted shark Na,K-ATPase is electrogenic. Biochemical and Biophysical Research Communications, 1989, 160, 801-807.	1.0	28

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37	Sequential substitution of K ⁺ bound to Na ⁺ ,K ⁺ -ATPase visualized by X-ray crystallography. <i>Nature Communications</i> , 2015, 6, 8004.	5.8	27
38	K ⁺ -Dependence of electrogenic transport by the Na ⁺ ,K ⁺ -ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1998, 1368, 184-200.	1.4	26
39	Rate Determination in Phosphorylation of Shark Rectal Na,K-ATPase by ATP: Temperature Sensitivity and Effects of ADP. <i>Biophysical Journal</i> , 1999, 77, 934-942.	0.2	26
40	Interaction of N-terminal peptide analogues of the Na ⁺ ,K ⁺ -ATPase with membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2018, 1860, 1282-1291.	1.4	26
41	The sided action of Na ⁺ on reconstituted shark Na ⁺ /K ⁺ -ATPase engaged in Na ⁺ -Na ⁺ exchange accompanied by ATP hydrolysis. II. Transmembrane allosteric effects on Na ⁺ affinity. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1988, 944, 223-232.	1.4	25
42	Mechanism of Mg ²⁺ Binding in the Na ⁺ ,K ⁺ -ATPase. <i>Biophysical Journal</i> , 2009, 96, 3753-3761.	0.2	25
43	Cholesterol depletion inhibits Na ⁺ ,K ⁺ -ATPase activity in a near-native membrane environment. <i>Journal of Biological Chemistry</i> , 2019, 294, 5956-5969.	1.6	25
44	Electrogenic pump current of sarcoplasmic reticulum Ca ²⁺ -ATPase reconstituted at high lipid/protein ratio. <i>FEBS Letters</i> , 1991, 284, 46-50.	1.3	23
45	Phosphorylation/dephosphorylation of reconstituted shark Na ⁺ ,K ⁺ -ATPase: one phosphorylation site per 1±1 ² protomer. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1995, 1235, 197-204.	1.4	23
46	Investigation of the enzymatic activity of the Na ⁺ ,K ⁺ -ATPase via isothermal titration microcalorimetry. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2010, 1797, 1540-1545.	0.5	23
47	Electrostatic Stabilization Plays a Central Role in Autoinhibitory Regulation of the Na ⁺ ,K ⁺ -ATPase. <i>Biophysical Journal</i> , 2017, 112, 288-299.	0.2	22
48	Functional Significance of the Shark Na,K-ATPase N-Terminal Domain. Is the Structurally Variable N-Terminus Involved in Tissue-Specific Regulation by FXYP Proteins?. <i>Biochemistry</i> , 2005, 44, 13051-13062.	1.2	21
49	Tension-length behaviour of a molluscan smooth muscle related to filament organisation. <i>Acta Physiologica Scandinavica</i> , 1978, 102, 167-180.	2.3	20
50	Interaction between Cardiotonic Steroids and Na,K-ATPase. Effects of pH and Ouabain-Induced Changes in Enzyme Conformation. <i>Biochemistry</i> , 2009, 48, 10056-10065.	1.2	20
51	The sided action of Na ⁺ and of K ⁺ on reconstituted shark (Na ⁺ + K ⁺)-ATPase engaged in Na ⁺ -Na ⁺ exchange accompanied by ATP hydrolysis. I. The ATP activation curve. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1987, 904, 353-364.	1.4	19
52	Interaction of FXYP10 (PLMS) with Na,K-ATPase from Shark Rectal Glands. <i>Journal of Biological Chemistry</i> , 2005, 280, 27776-27782.	1.6	19
53	Identification of Electric-Field-Dependent Steps in the Na ⁺ ,K ⁺ -Pump Cycle. <i>Biophysical Journal</i> , 2014, 107, 1352-1363.	0.2	18
54	Direct Activation of Gastric H,K-ATPase by N-Terminal Protein Kinase C Phosphorylation. Comparison of the Acute Regulation Mechanisms of H,K-ATPase and Na,K-ATPase. <i>Biophysical Journal</i> , 2003, 84, 1690-1700.	0.2	17

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55	Interaction of ATP with the Phosphoenzyme of the Na ⁺ ,K ⁺ -ATPase. <i>Biochemistry</i> , 2010, 49, 1248-1258.	1.2	16
56	Variable stoichiometry in reconstituted shark Na,K-ATPase engaged in uncoupled efflux. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1990, 1026, 147-152.	1.4	14
57	ATP Binding Equilibria of the Na ⁺ ,K ⁺ -ATPase. <i>Biochemistry</i> , 2008, 47, 13103-13114.	1.2	14
58	Modulation of FXYP Interaction with Na,K-ATPase by Anionic Phospholipids and Protein Kinase Phosphorylation. <i>Biochemistry</i> , 2007, 46, 2371-2379.	1.2	13
59	Inhibition of K ⁺ Transport through Na ⁺ , K ⁺ -ATPase by Capsazepine: Role of Membrane Span 10 of the β -Subunit in the Modulation of Ion Gating. <i>PLoS ONE</i> , 2014, 9, e96909.	1.1	13
60	Exploring the raft-hypothesis by probing planar bilayer patches of free-standing giant vesicles at nanoscale resolution, with and without Na,K-ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2016, 1858, 3041-3049.	1.4	13
61	The voltage-sensitive dye RH421 detects a Na ⁺ ,K ⁺ -ATPase conformational change at the membrane surface. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017, 1859, 813-823.	1.4	13
62	Dual Mechanisms of Allosteric Acceleration of the Na ⁺ ,K ⁺ -ATPase by ATP. <i>Biophysical Journal</i> , 2010, 98, 2290-2298.	0.2	12
63	Kinetics of K ⁺ Occlusion by the Phosphoenzyme of the Na ⁺ ,K ⁺ -ATPase. <i>Biophysical Journal</i> , 2011, 100, 70-79.	0.2	12
64	Membrane accessibility of glutathione. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 2430-2436.	1.4	12
65	Themes in Ion Pump Regulation. <i>Annals of the New York Academy of Sciences</i> , 2003, 986, 579-586.	1.8	11
66	Rb ⁺ occlusion stabilized by vanadate in gastric H ⁺ /K ⁺ -ATPase at 25°C. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011, 1808, 316-322.	1.4	11
67	Glutathionylation-Dependence of Na ⁺ -K ⁺ -Pump Currents Can Mimic Reduced Subsarcolemmal Na ⁺ Diffusion. <i>Biophysical Journal</i> , 2016, 110, 1099-1109.	0.2	11
68	The effect of cytoplasmic K ⁺ on the activity of the Na ⁺ /K ⁺ -ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1991, 1067, 227-234.	1.4	10
69	Hydrophobic ion interaction on Na ⁺ activation and dephosphorylation of reconstituted Na ⁺ ,K ⁺ -ATPase. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1995, 1235, 183-196.	1.4	10
70	Penetration of phospholipid membranes by poly-L-lysine depends on cholesterol and phospholipid composition. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2020, 1862, 183128.	1.4	10
71	Cryoelectron microscopy of Na ⁺ ,K ⁺ -ATPase in the two E2P states with and without cardiotonic steroids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2123226119.	3.3	10
72	Inorganic phosphate in ehrlich ascites tumor cells and its distribution across the cell membrane. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1978, 511, 213-223.	1.4	9

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73	Capturing suboptical dynamic structures in lipid bilayer patches formed from free-standing giant unilamellar vesicles. <i>Nature Protocols</i> , 2017, 12, 1563-1575.	5.5	9
74	A voltage-activated cation transport pathway associated with the sodium pump. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1991, 1070, 497-500.	1.4	5
75	Cis-allosteric effects of cytoplasmic Na ⁺ /K ⁺ discrimination at varying pH. Low-affinity multisite inhibition of cytoplasmic K ⁺ in reconstituted Na ⁺ /K ⁺ -ATPase engaged in uncoupled Na ⁺ -efflux. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1992, 1108, 190-200.	1.4	5
76	Kinetic Investigations of the Mechanism of the Rate-Determining Step of the Na ⁺ ,K ⁺ -ATPase Pump Cycle. <i>Annals of the New York Academy of Sciences</i> , 2003, 986, 159-162.	1.8	5
77	Liposomes in Reconstitution of Ion-Pumps. Electrogenic Properties of the Na ⁺ ,K ⁺ -Atpase and the Sarcoplasmic Ca ²⁺ -Atpase. <i>Journal of Liposome Research</i> , 1995, 5, 399-412.	1.5	4
78	Fluorescent Styryl Dyes as Probes for Na,K-ATPase Reaction.. <i>Annals of the New York Academy of Sciences</i> , 1997, 834, 394-396.	1.8	4
79	Protein Kinase C Phosphorylation Directed at Novel C-Terminal Sites in Na,K-ATPase. <i>Annals of the New York Academy of Sciences</i> , 2003, 986, 541-542.	1.8	4
80	Distinct pH dependencies of Na ⁺ /K ⁺ selectivity at the two faces of Na,K-ATPase. <i>Journal of Biological Chemistry</i> , 2018, 293, 2195-2205.	1.6	4
81	Reconstitution of transmembrane protein Na ⁺ ,K ⁺ -ATPase in giant unilamellar vesicles of lipid mixtures involving PSM, DOPC, DPPC and cholesterol at physiological buffer and temperature conditions. <i>Protocol Exchange</i> , 0, , .	0.3	4
82	Interaction between Substrate Site and Cation Binding Sites in PiPhosphorylation of Na,K-ATPase. <i>Annals of the New York Academy of Sciences</i> , 1997, 834, 390-393.	1.8	2
83	Order-disorder transitions of cytoplasmic N-termini in the mechanisms of P-type ATPases. <i>Faraday Discussions</i> , 2021, 232, 172-187.	1.6	2
84	Diversity of the E2P Phosphoforms of Na, K-ATPase. <i>Annals of the New York Academy of Sciences</i> , 1997, 834, 386-389.	1.8	1
85	PKA and PKC Phosphorylation of Gastric H,K-ATPase. <i>Annals of the New York Academy of Sciences</i> , 2003, 986, 548-549.	1.8	1
86	X-Ray Crystallographic Study of Na,K-ATPase in Complex with Cardiotonic Steroids. <i>Biophysical Journal</i> , 2015, 108, 197a.	0.2	1
87	The sodium PUMP. <i>Biomembranes: A Multi-Volume Treatise</i> , 1996, 5, 133-184.	0.1	0
88	Species-specific peculiarities of functional reactions of the sodium pump to phosphorylation by protein kinase A. <i>Journal of Evolutionary Biochemistry and Physiology</i> , 2000, 36, 11-16.	0.2	0
89	Kinetics by X-Ray Crystallography: Sequential Substitution of K ⁺ Bound To Na ⁺ , K ⁺ -ATPase. <i>Biophysical Journal</i> , 2016, 110, 629a.	0.2	0
90	To Image the Orientation and Spatial Distribution of Reconstituted Na ⁺ ,K ⁺ -ATPase in Model Lipid Membranes. , 2019, , 29-46.		0

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91	Active Biomimetic Membranes. Biological and Medical Physics Series, 2011, , 113-135.	0.3	0
92	Displacement of Native FXYP Protein From Na ⁺ /K ⁺ -ATPase With Novel FXYP Peptide Derivatives: Effects on Doxorubicin Cytotoxicity. Frontiers in Oncology, 2022, 12, 859216.	1.3	0