Roderick MacKinnon

List of Publications by Year in Descending Order

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128 33,507 109 73 h-index g-index citations papers 128 37,024 25.4 7.59 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
109	Correlation between structure and function in phosphatidylinositol lipid-dependent Kir2.2 gating Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e211404611	9 ^{11.5}	O
108	Molecular structure of an open human K channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	7
107	Analysis of the mechanosensor channel functionality of TACAN. ELife, 2021, 10,	8.9	7
106	Cryo-EM analysis of PIP regulation in mammalian GIRK channels. ELife, 2020, 9,	8.9	21
105	Structural Basis of Human KCNQ1 Modulation and Gating. <i>Cell</i> , 2020 , 180, 340-347.e9	56.2	85
104	Force-induced conformational changes in PIEZO1. <i>Nature</i> , 2019 , 573, 230-234	50.4	106
103	Regulation of Eag1 gating by its intracellular domains. <i>ELife</i> , 2019 , 8,	8.9	9
102	The mechanosensitive ion channel TRAAK is localized to the mammalian node of Ranvier. <i>ELife</i> , 2019 , 8,	8.9	33
101	Molecular structures of the human Slo1 K channel in complex with 4. ELife, 2019, 8,	8.9	35
100	Cryo-EM structure of the KvAP channel reveals a non-domain-swapped voltage sensor topology. <i>ELife</i> , 2019 , 8,	8.9	10
99	Voltage Sensor Movements during Hyperpolarization in the HCN Channel. <i>Cell</i> , 2019 , 179, 1582-1589.e7	′56.2	41
98	Activation mechanism of a human SK-calmodulin channel complex elucidated by cryo-EM structures. <i>Science</i> , 2018 , 360, 508-513	33.3	87
97	Piezo1 forms a slowly-inactivating mechanosensory channel in mouse embryonic stem cells. <i>ELife</i> , 2018 , 7,	8.9	38
96	Piezo's membrane footprint and its contribution to mechanosensitivity. <i>ELife</i> , 2018 , 7,	8.9	67
95	Molecular basis of signaling specificity between GIRK channels and GPCRs. ELife, 2018, 7,	8.9	20
94	Author response: Piezol membrane footprint and its contribution to mechanosensitivity 2018,		2
93	Structure of the CLC-1 chloride channel from. <i>ELife</i> , 2018 , 7,	8.9	52

92	Structural Titration of Slo2.2, a Na-Dependent K Channel. Cell, 2017, 168, 390-399.e11	56.2	74
91	Structures of the Human HCN1 Hyperpolarization-Activated Channel. <i>Cell</i> , 2017 , 168, 111-120.e11	56.2	185
90	Cryo-EM Structure of the Open Human Ether-Ego-go-Related K Channel hERG. <i>Cell</i> , 2017 , 169, 422-430.	e 56 .2	262
89	Cryo-EM Structure of a KCNQ1/CaM Complex Reveals Insights into Congenital Long QT Syndrome. <i>Cell</i> , 2017 , 169, 1042-1050.e9	56.2	182
88	Structure of a CLC chloride ion channel by cryo-electron microscopy. <i>Nature</i> , 2017 , 541, 500-505	50.4	87
87	Cryo-EM structure of the open high-conductance Ca-activated K channel. <i>Nature</i> , 2017 , 541, 46-51	50.4	134
86	Structural basis for gating the high-conductance Ca-activated K channel. <i>Nature</i> , 2017 , 541, 52-57	50.4	103
85	Structure-based membrane dome mechanism for Piezo mechanosensitivity. ELife, 2017, 6,	8.9	175
84	Molecular structure of human KATP in complex with ATP and ADP. ELife, 2017, 6,	8.9	89
83	Author response: Structure-based membrane dome mechanism for Piezo mechanosensitivity 2017,		3
83	Author response: Structure-based membrane dome mechanism for Piezo mechanosensitivity 2017 , The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5,	8.9	7
	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> ,	8.9	
82	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5,		7
82	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5, Cooperative regulation by G proteins and Na(+) of neuronal GIRK2 K(+) channels. <i>ELife</i> , 2016 , 5, Novel cell-free high-throughput screening method for pharmacological tools targeting K+ channels.	8.9	7 26
82 81 80	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5, Cooperative regulation by G proteins and Na(+) of neuronal GIRK2 K(+) channels. <i>ELife</i> , 2016 , 5, Novel cell-free high-throughput screening method for pharmacological tools targeting K+ channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 5748-53 Structure of the voltage-gated K+ channel Eag1 reveals an alternative voltage sensing mechanism.	8.9	7 26 62 190
82 81 80	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5, Cooperative regulation by G proteins and Na(+) of neuronal GIRK2 K(+) channels. <i>ELife</i> , 2016 , 5, Novel cell-free high-throughput screening method for pharmacological tools targeting K+ channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 5748-53 Structure of the voltage-gated K+ channel Eag1 reveals an alternative voltage sensing mechanism. <i>Science</i> , 2016 , 353, 664-9	8.9	7 26 62 190
82 81 80 79 78	The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. <i>ELife</i> , 2016 , 5, Cooperative regulation by G proteins and Na(+) of neuronal GIRK2 K(+) channels. <i>ELife</i> , 2016 , 5, Novel cell-free high-throughput screening method for pharmacological tools targeting K+ channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 5748-53 Structure of the voltage-gated K+ channel Eag1 reveals an alternative voltage sensing mechanism. <i>Science</i> , 2016 , 353, 664-9 Cryo-electron microscopy structure of the Slo2.2 Na(+)-activated K(+) channel. <i>Nature</i> , 2015 , 527, 198-20.	8.9 11.5 33.3	7 26 62 190 83

74	Phosphatidic acid modulation of Kv channel voltage sensor function. <i>ELife</i> , 2014 , 3,	8.9	41
73	Author response: Phosphatidic acid modulation of Kv channel voltage sensor function 2014,		2
72	X-ray structure of the mammalian GIRK2-IG-protein complex. <i>Nature</i> , 2013 , 498, 190-7	50.4	218
71	Domain-swapped chain connectivity and gated membrane access in a Fab-mediated crystal of the human TRAAK K+ channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 2129-34	11.5	91
70	Structure of a pore-blocking toxin in complex with a eukaryotic voltage-dependent K(+) channel. <i>ELife</i> , 2013 , 2, e00594	8.9	139
69	Crystal structure of the human K2P TRAAK, a lipid- and mechano-sensitive K+ ion channel. <i>Science</i> , 2012 , 335, 436-41	33.3	307
68	Functional and structural analysis of the human SLO3 pH- and voltage-gated K+ channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 19274-9	11.5	42
67	Molecular mechanism of proton transport in CLC Cl-/H+ exchange transporters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 11699-704	11.5	47
66	Open structure of the Ca2+ gating ring in the high-conductance Ca2+-activated K+ channel. <i>Nature</i> , 2011 , 481, 94-7	50.4	109
65	Structural basis of PIP2 activation of the classical inward rectifier K+ channel Kir2.2. <i>Nature</i> , 2011 , 477, 495-8	50.4	456
64	Crystal structure of the mammalian GIRK2 K+ channel and gating regulation by G proteins, PIP2, and sodium. <i>Cell</i> , 2011 , 147, 199-208	56.2	340
63	Structure of the human BK channel Ca2+-activation apparatus at 3.0 A resolution. <i>Science</i> , 2010 , 329, 182-6	33.3	249
62	A gating charge transfer center in voltage sensors. <i>Science</i> , 2010 , 328, 67-73	33.3	357
61	Solution structure and phospholipid interactions of the isolated voltage-sensor domain from KvAP. <i>Journal of Molecular Biology</i> , 2010 , 403, 591-606	6.5	81
60	Structure of a eukaryotic CLC transporter defines an intermediate state in the transport cycle. <i>Science</i> , 2010 , 330, 635-41	33.3	207
59	Crystal structure of the eukaryotic strong inward-rectifier K+ channel Kir2.2 at 3.1 A resolution. <i>Science</i> , 2009 , 326, 1668-74	33.3	274
58	Functional reconstitution of purified human Hv1 H+ channels. <i>Journal of Molecular Biology</i> , 2009 , 387, 1055-60	6.5	85
57	A gating model for the archeal voltage-dependent K(+) channel KvAP in DPhPC and POPE:POPG decane lipid bilayers. <i>Journal of Molecular Biology</i> , 2009 , 390, 902-12	6.5	58

(2004-2009)

56	Two separate interfaces between the voltage sensor and pore are required for the function of voltage-dependent K(+) channels. <i>PLoS Biology</i> , 2009 , 7, e47	9.7	125
55	Inferred motions of the S3a helix during voltage-dependent K+ channel gating. <i>Journal of Molecular Biology</i> , 2008 , 381, 569-80	6.5	23
54	Functional analysis of Kv1.2 and paddle chimera Kv channels in planar lipid bilayers. <i>Journal of Molecular Biology</i> , 2008 , 382, 24-33	6.5	38
53	Voltage-dependent K+ channel gating and voltage sensor toxin sensitivity depend on the mechanical state of the lipid membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 19276-81	11.5	133
52	Potassium Channels and the Atomic Basis of Selective Ion Conduction 2008, 431-461		
51	Crystal structure of a Kir3.1-prokaryotic Kir channel chimera. <i>EMBO Journal</i> , 2007 , 26, 4005-15	13	257
50	Atomic structure of a voltage-dependent K+ channel in a lipid membrane-like environment. <i>Nature</i> , 2007 , 450, 376-82	50.4	1164
49	Structural and thermodynamic properties of selective ion binding in a K+ channel. <i>PLoS Biology</i> , 2007 , 5, e121	9.7	183
48	Ion selectivity in a semisynthetic K+ channel locked in the conductive conformation. <i>Science</i> , 2006 , 314, 1004-7	33.3	113
47	Structural and functional consequences of an amide-to-ester substitution in the selectivity filter of a potassium channel. <i>Journal of the American Chemical Society</i> , 2006 , 128, 11591-9	16.4	47
46	Phospholipids and the origin of cationic gating charges in voltage sensors. <i>Nature</i> , 2006 , 444, 775-9	50.4	336
45	Principles of selective ion transport in channels and pumps. <i>Science</i> , 2005 , 310, 1461-5	33.3	672
44	Crystal structure of a mammalian voltage-dependent Shaker family K+ channel. <i>Science</i> , 2005 , 309, 897-	·903 33	1835
43	Voltage sensor of Kv1.2: structural basis of electromechanical coupling. <i>Science</i> , 2005 , 309, 903-8	33.3	819
42	Calibrated measurement of gating-charge arginine displacement in the KvAP voltage-dependent K+ channel. <i>Cell</i> , 2005 , 123, 463-75	56.2	182
41	Structural biology. Membrane protein insertion and stability. <i>Science</i> , 2005 , 307, 1425-6	33.3	25
40	Structure of the KvAP voltage-dependent K+ channel and its dependence on the lipid membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 15441-6	11.5	265
39	Structural biology. Voltage sensor meets lipid membrane. <i>Science</i> , 2004 , 306, 1304-5	33.3	30

38	A membrane-access mechanism of ion channel inhibition by voltage sensor toxins from spider venom. <i>Nature</i> , 2004 , 430, 232-5	50.4	239
37	Electron microscopic analysis of KvAP voltage-dependent K+ channels in an open conformation. <i>Nature</i> , 2004 , 430, 806-10	50.4	116
36	Nobel Lecture. Potassium channels and the atomic basis of selective ion conduction. <i>Bioscience Reports</i> , 2004 , 24, 75-100	4.1	63
35	Potassium channels and the atomic basis of selective ion conduction (Nobel Lecture). <i>Angewandte Chemie - International Edition</i> , 2004 , 43, 4265-77	16.4	293
34	Kaliumkan [e und die atomare Basis der selektiven Ionenleitung (Nobel-Vortrag). <i>Angewandte Chemie</i> , 2004 , 116, 4363-4376	3.6	47
33	Ion binding affinity in the cavity of the KcsA potassium channel. <i>Biochemistry</i> , 2004 , 43, 4978-82	3.2	74
32	Localization of the voltage-sensor toxin receptor on KvAP. <i>Biochemistry</i> , 2004 , 43, 10071-9	3.2	76
31	A mutant KcsA K(+) channel with altered conduction properties and selectivity filter ion distribution. <i>Journal of Molecular Biology</i> , 2004 , 338, 839-46	6.5	104
30	Functional analysis of an archaebacterial voltage-dependent K+ channel. <i>Nature</i> , 2003 , 422, 180-5	50.4	193
29	X-ray structure of a voltage-dependent K+ channel. <i>Nature</i> , 2003 , 423, 33-41	50.4	1630
28	The principle of gating charge movement in a voltage-dependent K+ channel. <i>Nature</i> , 2003 , 423, 42-8	50.4	691
27	The occupancy of ions in the K+ selectivity filter: charge balance and coupling of ion binding to a protein conformational change underlie high conduction rates. <i>Journal of Molecular Biology</i> , 2003 , 333, 965-75	6.5	340
26	Potassium channels. FEBS Letters, 2003, 555, 62-5	3.8	348
25	Gating the selectivity filter in ClC chloride channels. <i>Science</i> , 2003 , 300, 108-12	33.3	659
24	X-ray structure of a ClC chloride channel at 3.0 A reveals the molecular basis of anion selectivity. <i>Nature</i> , 2002 , 415, 287-94		1364
24			1364
	Nature, 2002 , 415, 287-94	50.4	

20	Potassium channel receptor site for the inactivation gate and quaternary amine inhibitors. <i>Nature</i> , 2001 , 411, 657-61	50.4	497
19	Energetic optimization of ion conduction rate by the K+ selectivity filter. <i>Nature</i> , 2001 , 414, 37-42	50.4	660
18	Chemistry of ion coordination and hydration revealed by a K+ channel-Fab complex at 2.0 A resolution. <i>Nature</i> , 2001 , 414, 43-8	50.4	1713
17	Structure of the RCK domain from the E. coli K+ channel and demonstration of its presence in the human BK channel. <i>Neuron</i> , 2001 , 29, 593-601	13.9	264
16	The cavity and pore helices in the KcsA K+ channel: electrostatic stabilization of monovalent cations. <i>Science</i> , 1999 , 285, 100-2	33.3	389
15	Crystal structure and functional analysis of the HERG potassium channel N terminus: a eukaryotic PAS domain. <i>Cell</i> , 1998 , 95, 649-55	56.2	385
14	A snake toxin inhibitor of inward rectifier potassium channel ROMK1. <i>Biochemistry</i> , 1998 , 37, 14867-74	3.2	47
13	The structure of the potassium channel: molecular basis of K+ conduction and selectivity. <i>Science</i> , 1998 , 280, 69-77	33.3	5694
12	Purification, characterization, and synthesis of an inward-rectifier K+ channel inhibitor from scorpion venom. <i>Biochemistry</i> , 1997 , 36, 6936-40	3.2	42
11	Hanatoxin modifies the gating of a voltage-dependent K+ channel through multiple binding sites. <i>Neuron</i> , 1997 , 18, 665-73	13.9	226
10	Mapping the receptor site for hanatoxin, a gating modifier of voltage-dependent K+ channels. <i>Neuron</i> , 1997 , 18, 675-82	13.9	215
9	Contribution of the S4 segment to gating charge in the Shaker K+ channel. <i>Neuron</i> , 1996 , 16, 1169-77	13.9	600
8	Solution structure of the potassium channel inhibitor agitoxin 2: caliper for probing channel geometry. <i>Protein Science</i> , 1995 , 4, 1478-89	6.3	112
7	An inhibitor of the Kv2.1 potassium channel isolated from the venom of a Chilean tarantula. <i>Neuron</i> , 1995 , 15, 941-9	13.9	228
6	Electrostatic tuning of Mg2+ affinity in an inward-rectifier K+ channel. <i>Nature</i> , 1994 , 371, 243-6	50.4	288
5	Purification and characterization of three inhibitors of voltage-dependent K+ channels from Leiurus quinquestriatus var. hebraeus venom. <i>Biochemistry</i> , 1994 , 33, 6834-9	3.2	243
4	Determination of the subunit stoichiometry of a voltage-activated potassium channel. <i>Nature</i> , 1991 , 350, 232-5	50.4	853
3	The mechanosensitive ion channel TRAAK is localized to the mammalian node of Ranvier		1

2 Analysis of the Mechanosensor Channel Functionality of TACAN

3

Correlation between structure and function in phosphatidylinositol lipid-dependent Kir2.2 gating

1