Anna Weinzinger

List of Publications by Citations

Source: https://exaly.com/author-pdf/6798723/anna-weinzinger-publications-by-citations.pdf

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

60 28 954 20 h-index g-index citations papers 80 1,123 4.07 3.7 L-index avg, IF ext. papers ext. citations

#	Paper	IF	Citations
60	Grayanotoxin poisoning: Thad honey diseaseTand beyond. Cardiovascular Toxicology, 2012, 12, 208-15	3.4	77
59	Toward a consensus model of the HERG potassium channel. <i>ChemMedChem</i> , 2010 , 5, 455-67	3.7	57
58	New potential binding determinant for hERG channel inhibitors. <i>Scientific Reports</i> , 2016 , 6, 24182	4.9	55
57	Structural basis of control of inward rectifier Kir2 channel gating by bulk anionic phospholipids. <i>Journal of General Physiology</i> , 2016 , 148, 227-37	3.4	45
56	The anti-protozoal drug pentamidine blocks KIR2.x-mediated inward rectifier current by entering the cytoplasmic pore region of the channel. <i>British Journal of Pharmacology</i> , 2010 , 159, 1532-41	8.6	39
55	Computer simulations of structure-activity relationships for HERG channel blockers. <i>Biochemistry</i> , 2011 , 50, 6146-56	3.2	36
54	Efficient and specific cardiac IKIInhibition by a new pentamidine analogue. <i>Cardiovascular Research</i> , 2013 , 99, 203-14	9.9	32
53	Distinct interactions of Na+ and Ca2+ ions with the selectivity filter of the bacterial sodium channel Na(V)Ab. <i>Biochemical and Biophysical Research Communications</i> , 2013 , 430, 1272-6	3.4	29
52	Molecular determinants for activation of human ether-Ego-go-related gene 1 potassium channels by 3-nitro-n-(4-phenoxyphenyl) benzamide. <i>Molecular Pharmacology</i> , 2011 , 80, 630-7	4.3	29
51	Probing the energy landscape of activation gating of the bacterial potassium channel KcsA. <i>PLoS Computational Biology</i> , 2013 , 9, e1003058	5	28
50	Timothy mutation disrupts the link between activation and inactivation in Ca(V)1.2 protein. <i>Journal of Biological Chemistry</i> , 2011 , 286, 31557-64	5.4	27
49	Mechanism of hERG channel block by the psychoactive indole alkaloid ibogaine. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2014 , 348, 346-58	4.7	25
48	Conserved functional consequences of disease-associated mutations in the slide helix of Kir6.1 and Kir6.2 subunits of the ATP-sensitive potassium channel. <i>Journal of Biological Chemistry</i> , 2017 , 292, 1738	37 ⁵ 473	98 ³
47	Different inward and outward conduction mechanisms in NaVMs suggested by molecular dynamics simulations. <i>PLoS Computational Biology</i> , 2014 , 10, e1003746	5	23
46	Conduction through a narrow inward-rectifier K channel pore. <i>Journal of General Physiology</i> , 2019 , 151, 1231-1246	3.4	22
45	In silico analysis of conformational changes induced by mutation of aromatic binding residues: consequences for drug binding in the hERG K+ channel. <i>PLoS ONE</i> , 2011 , 6, e28778	3.7	22
44	Probing the architecture of an L-type calcium channel with a charged phenylalkylamine: evidence for a widely open pore and drug trapping. <i>Journal of Biological Chemistry</i> , 2007 , 282, 3864-70	5.4	21

(2019-2007)

43	Differences in (-)citronellal binding to various odorant receptors. <i>Biochemical and Biophysical Research Communications</i> , 2007 , 361, 941-5	3.4	21
42	Coupled and independent contributions of residues in IS6 and IIS6 to activation gating of CaV1.2. Journal of Biological Chemistry, 2009 , 284, 12276-84	5.4	20
41	Structural model of the Ca V 1.2 pore. <i>Channels</i> , 2008 , 2, 210-5	3	20
40	Molecular Dynamics Simulations of KirBac1.1 Mutants Reveal Global Gating Changes of Kir Channels. <i>Journal of Chemical Information and Modeling</i> , 2015 , 55, 814-22	6.1	19
39	The hERG potassium channel and drug trapping: insight from docking studies with propafenone derivatives. <i>ChemMedChem</i> , 2010 , 5, 436-42	3.7	18
38	Surprises from an unusual CLC homolog. <i>Biophysical Journal</i> , 2012 , 103, L44-6	2.9	16
37	ICA-105574 interacts with a common binding site to elicit opposite effects on inactivation gating of EAG and ERG potassium channels. <i>Molecular Pharmacology</i> , 2013 , 83, 805-13	4.3	16
36	Structure-activity relationships of pentamidine-affected ion channel trafficking and dofetilide mediated rescue. <i>British Journal of Pharmacology</i> , 2013 , 169, 1322-34	8.6	15
35	Dehydroevodiamine and hortiamine, alkaloids from the traditional Chinese herbal drug Evodia rutaecarpa, are I blockers with proarrhythmic effects in vitro and in vivo. <i>Pharmacological Research</i> , 2018 , 131, 150-163	10.2	14
34	Molecular dynamics and mutational analysis of a channelopathy mutation in the IIS6 helix of Ca V 1.2. <i>Channels</i> , 2008 , 2, 216-23	3	14
33	LUF7244, an allosteric modulator/activator of K 11.1 channels, counteracts dofetilide-induced torsades de pointes arrhythmia in the chronic atrioventricular block dog model. <i>British Journal of Pharmacology</i> , 2019 , 176, 3871-3885	8.6	12
32	Cysteines in the loop between IS5 and the pore helix of Ca(V)3.1 are essential for channel gating. <i>Pflugers Archiv European Journal of Physiology</i> , 2010 , 460, 1015-28	4.6	12
31	Neutralisation of a single voltage sensor affects gating determinants in all four pore-forming S6 segments of Ca(V)1.2: a cooperative gating model. <i>Pflugers Archiv European Journal of Physiology</i> , 2012 , 464, 391-401	4.6	11
30	Pore stability and gating in voltage-activated calcium channels. <i>Channels</i> , 2008 , 2, 61-9	3	11
29	Atomistic basis of opening and conduction in mammalian inward rectifier potassium (Kir2.2) channels. <i>Journal of General Physiology</i> , 2020 , 152,	3.4	11
28	PA-6 inhibits inward rectifier currents carried by V93I and D172N gain-of-function K2.1 channels, but increases channel protein expression. <i>Journal of Biomedical Science</i> , 2017 , 24, 44	13.3	10
27	Molecular Insights into hERG Potassium Channel Blockade by Lubeluzole. <i>Cellular Physiology and Biochemistry</i> , 2018 , 45, 2233-2245	3.9	10
26	A structural model of the human serotonin transporter in an outward-occluded state. <i>PLoS ONE</i> , 2019 , 14, e0217377	3.7	10

25	Disease Associated Mutations in K Proteins Linked to Aberrant Inward Rectifier Channel Trafficking. <i>Biomolecules</i> , 2019 , 9,	5.9	10
24	Inhibition of cardiac inward rectifier currents by cationic amphiphilic drugs. <i>Current Molecular Medicine</i> , 2013 , 13, 1284-98	2.5	10
23	Computational Insights Into Voltage Dependence of Polyamine Block in a Strong Inwardly Rectifying K Channel. <i>Frontiers in Pharmacology</i> , 2020 , 11, 721	5.6	8
22	Glibenclamide and HMR1098 normalize Cant yndrome-associated gain-of-function currents. <i>Journal of Cellular and Molecular Medicine</i> , 2019 , 23, 4962-4969	5.6	7
21	Exploring the structure of the voltage-gated Na+ channel by an engineered drug access pathway to the receptor site for local anesthetics. <i>Journal of Biological Chemistry</i> , 2014 , 289, 21770-81	5.4	7
20	Structural insights into trapping and dissociation of small molecules in K+ channels. <i>Journal of Chemical Information and Modeling</i> , 2014 , 54, 3218-28	6.1	6
19	Insights in KIR2.1 channel structure and function by an evolutionary approach; cloning and functional characterization of the first reptilian inward rectifier channel KIR2.1, derived from the California kingsnake (Lampropeltis getula californiae). <i>Biochemical and Biophysical Research</i>	3.4	6
18	Communications, 2014 , 452, 992-7 Physicochemical properties of pore residues predict activation gating of Ca V1.2: a correlation mutation analysis. <i>Pflugers Archiv European Journal of Physiology</i> , 2011 , 461, 53-63	4.6	6
17	LUF7244 plus Dofetilide Rescues Aberrant K11.1 Trafficking and Produces Functional I. <i>Molecular Pharmacology</i> , 2020 , 97, 355-364	4.3	5
16	Distinct modulation of inactivation by a residue in the pore domain of voltage-gated Na channels: mechanistic insights from recent crystal structures. <i>Scientific Reports</i> , 2018 , 8, 631	4.9	5
15	Toward a Structural View of hERG Activation by the Small-Molecule Activator ICA-105574. <i>Journal of Chemical Information and Modeling</i> , 2020 , 60, 360-371	6.1	5
14	Molecular Basis of Altered hERG1 Channel Gating Induced by Ginsenoside Rg3. <i>Molecular Pharmacology</i> , 2017 , 92, 437-450	4.3	4
13	Evolutionary trace of human odorant receptors of chromosome 17. <i>Flavour and Fragrance Journal</i> , 2009 , 24, 192-197	2.5	4
12	Dynamics of the EAG1 K channel selectivity filter assessed by molecular dynamics simulations. <i>Biochemical and Biophysical Research Communications</i> , 2017 , 484, 107-112	3.4	3
11	Drug trapping in hERG K channels: (not) a matter of drug size?. <i>MedChemComm</i> , 2016 , 7, 512-518	5	3
10	Computational Identification of Novel Kir6 Channel Inhibitors. <i>Frontiers in Pharmacology</i> , 2019 , 10, 549	5.6	2
9	Histidine at position 462 determines the low quinine sensitivity of ether-Ego-go channel superfamily member K 12.1. <i>British Journal of Pharmacology</i> , 2019 , 176, 2708-2723	8.6	2
8	Genomics of selected human odorant receptors. <i>Monatshefte Fa Chemie</i> , 2008 , 139, 1537-1544	1.4	2

LIST OF PUBLICATIONS

7	Atomistic basis of opening and conduction in mammalian inward rectifier potassium (Kir2.2) channels		2
6	Simulating PIP-Induced Gating Transitions in Kir6.2 Channels. <i>Frontiers in Molecular Biosciences</i> , 2021 , 8, 711975	5.6	2
5	Development of I Ion Channel Blockers Targeting Sulfonylurea Resistant Mutant K6.2 Based Channels for Treating DEND Syndrome <i>Frontiers in Pharmacology</i> , 2021 , 12, 814066	5.6	1
4	Computational identification of novel Kir6 channel inhibitors		1
3	A selectivity filter mutation provides insights into gating regulation of a K channel <i>Communications Biology</i> , 2022 , 5, 345	6.7	1
2	The Bradycardic Agent Ivabradine Acts as an Atypical Inhibitor of Voltage-Gated Sodium Channels <i>Frontiers in Pharmacology</i> , 2022 , 13, 809802	5.6	1
1	MD Simulations of KirBac1.1 Mutants Reveal Gating Changes at the Bundle Crossing Region. Biophysical Journal, 2013, 104, 213a	2.9	