

Kirill D Nadezhdin

List of Publications by Year in descending order

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35
papers

1,015
citations

471477

17
h-index

434170

31
g-index

39
all docs

39
docs citations

39
times ranked

1072
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural mechanism of TRPV3 channel inhibition by the anesthetic dyclonine. <i>Nature Communications</i> , 2022, 13, 2795.	12.8	17
2	Structural snapshots of the mechanism of TRPV2 channel activation by small-molecule agonists. <i>Cell Calcium</i> , 2022, 105, 102607.	2.4	0
3	Structure and function of the calcium-selective TRP channel TRPV6. <i>Journal of Physiology</i> , 2021, 599, 2673-2697.	2.9	29
4	TRPV3 expression and purification for structure determination by Cryo-EM. <i>Methods in Enzymology</i> , 2021, 652, 31-48.	1.0	11
5	Extracellular cap domain is an essential component of the TRPV1 gating mechanism. <i>Nature Communications</i> , 2021, 12, 2154.	12.8	40
6	Structural mechanism of heat-induced opening of a temperature-sensitive TRP channel. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 564-572.	8.2	76
7	Interaction between the transmembrane domains of neurotrophin receptors p75 and TrkA mediates their reciprocal activation. <i>Journal of Biological Chemistry</i> , 2021, 297, 100926.	3.4	8
8	Structural mechanism of TRPV3 channel inhibition by the plant-derived coumarin osthole. <i>EMBO Reports</i> , 2021, 22, e53233.	4.5	26
9	Structural mechanisms of TRPV6 inhibition by ruthenium red and econazole. <i>Nature Communications</i> , 2021, 12, 6284.	12.8	30
10	All-d-Enantiomeric Peptide D3 Designed for Alzheimer's Disease Treatment Dynamically Interacts with Membrane-Bound Amyloid- β Precursors. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 16464-16479.	6.4	7
11	Structural basis of the transmembrane domain dimerization and rotation in the activation mechanism of the TRKA receptor by nerve growth factor. <i>Journal of Biological Chemistry</i> , 2020, 295, 275-286.	3.4	22
12	Inactivation-mimicking block of the epithelial calcium channel TRPV6. <i>Science Advances</i> , 2020, 6, .	10.3	22
13	Revising the mechanism of p75NTR activation: intrinsically monomeric state of death domains invokes the "helper" hypothesis. <i>Scientific Reports</i> , 2020, 10, 13686.	3.3	7
14	Purification of native CCL7 and its functional interaction with selected chemokine receptors. <i>Protein Expression and Purification</i> , 2020, 171, 105617.	1.3	6
15	Medicinal leech antimicrobial peptides lacking toxicity represent a promising alternative strategy to combat antibiotic-resistant pathogens. <i>European Journal of Medicinal Chemistry</i> , 2019, 180, 143-153.	5.5	17
16	Familial L723P Mutation Can Shift the Distribution between the Alternative APP Transmembrane Domain Cleavage Cascades by Local Unfolding of the β -Cleavage Site Suggesting a Straightforward Mechanism of Alzheimer's Disease Pathogenesis. <i>ACS Chemical Biology</i> , 2019, 14, 1573-1582.	3.4	13
17	NMR structure of a full-length single-pass membrane protein NRADD. <i>Proteins: Structure, Function and Bioinformatics</i> , 2019, 87, 786-790.	2.6	4
18	Modular toxin from the lynx spider <i>Oxyopes takobius</i> : Structure of spiderine domains in solution and membrane-mimicking environment. <i>Protein Science</i> , 2017, 26, 611-616.	7.6	10

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19	FaÅšade detergents as bicelle rim-forming agents for solution NMR spectroscopy. <i>Nanotechnology Reviews</i> , 2017, 6, 93-103.	5.8	9
20	Membrane mimetics for solution NMR studies of membrane proteins. <i>Nanotechnology Reviews</i> , 2017, 6, 15-32.	5.8	25
21	Cell-free expression of the APP transmembrane fragments with Alzheimer's disease mutations using algal amino acid mixture for structural NMR studies. <i>Protein Expression and Purification</i> , 2016, 123, 105-111.	1.3	12
22	Structural Basis of p75 Transmembrane Domain Dimerization. <i>Journal of Biological Chemistry</i> , 2016, 291, 12346-12357.	3.4	27
23	Structure of purotoxin-2 from wolf spider: modular design and membrane-assisted mode of action in arachnid toxins. <i>Biochemical Journal</i> , 2016, 473, 3113-3126.	3.7	16
24	Characterization of Small Isotropic Bicelles with Various Compositions. <i>Langmuir</i> , 2016, 32, 6624-6637.	3.5	47
25	AsLn2, a luciferin-related modified tripeptide from the bioluminescent earthworm <i>Fridericia heliota</i> . <i>Tetrahedron Letters</i> , 2014, 55, 463-465.	1.4	12
26	Bacterial and cell-free production of APP671-726 containing amyloid precursor protein transmembrane and metal-binding domains. <i>Biochemistry (Moscow)</i> , 2013, 78, 1263-1271.	1.5	6
27	Structure of Transmembrane Domain and Dimerization Mechanism of Amyloid Precursor Protein. <i>Biophysical Journal</i> , 2012, 102, 263a.	0.5	0
28	Dimeric structure of transmembrane domain of amyloid precursor protein in micellar environment. <i>FEBS Letters</i> , 2012, 586, 1687-1692.	2.8	77
29	Novel peptide from spider venom inhibits P2X3 receptors and inflammatory pain. <i>Annals of Neurology</i> , 2010, 67, 680-683.	5.3	55
30	Isolation, Structure Elucidation, and Synergistic Antibacterial Activity of a Novel Two-Component Lantibiotic Lichenicidin from <i>Bacillus licheniformis</i> VK21. <i>Biochemistry</i> , 2010, 49, 6462-6472.	2.5	67
31	Divalent cation coordination and mode of membrane interaction in cyclotides: NMR spatial structure of ternary complex Kalata B7/Mn ²⁺ /DPC micelle. <i>Journal of Inorganic Biochemistry</i> , 2008, 102, 1246-1256.	3.5	56
32	Molecular insight into mechanism of antimicrobial action of the Î²â€šhairpin peptide arenicin: Specific oligomerization in detergent micelles. <i>Biopolymers</i> , 2008, 89, 455-464.	2.4	43
33	Recombinant expression, synthesis, purification, and solution structure of arenicin. <i>Biochemical and Biophysical Research Communications</i> , 2007, 360, 156-162.	2.1	70
34	Antiamoebin I in Methanol Solution: Rapid Exchange between Right-Handed and Left-Handed310-Helical Conformations. <i>Chemistry and Biodiversity</i> , 2007, 4, 1219-1242.	2.1	23
35	Conformation and mode of membrane interaction in cyclotides. <i>FEBS Journal</i> , 2006, 273, 2658-2672.	4.7	124