## Justyna Iwaszkiewicz

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Dominant monoallelic variant in the PAK2 gene causes Knobloch syndrome type 2. Human Molecular Genetics, 2021, 31, 1-9.	2.9	6
2	Influenza A viruses balance ER stress with host protein synthesis shutoff. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	17
3	Taurine treatment of retinal degeneration and cardiomyopathy in a consanguineous family with SLC6A6 taurine transporter deficiency. Human Molecular Genetics, 2020, 29, 618-623.	2.9	29
4	De Novo and Bi-allelic Pathogenic Variants in NARS1 Cause Neurodevelopmental Delay Due to Toxic Gain-of-Function and Partial Loss-of-Function Effects. American Journal of Human Genetics, 2020, 107, 311-324.	6.2	32
5	Biallelic variants in PSMB1 encoding the proteasome subunit β6 cause impairment of proteasome function, microcephaly, intellectual disability, developmental delay and short stature. Human Molecular Genetics, 2020, 29, 1132-1143.	2.9	30
6	Allosteric activation of MALT1 by its ubiquitin-binding Ig3 domain. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 3093-3102.	7.1	14
7	Disulfide-Linked Peptides for Blocking BTLA/HVEM Binding. International Journal of Molecular Sciences, 2020, 21, 636.	4.1	15
8	Bi-allelic Variants in DYNC112 Cause Syndromic Microcephaly with Intellectual Disability, Cerebral Malformations, and Dysmorphic Facial Features. American Journal of Human Genetics, 2019, 104, 1073-1087.	6.2	19
9	Mutations in the palm domain disrupt modulation of acid-sensing ion channel 1a currents by neuropeptides. Scientific Reports, 2019, 9, 2599.	3.3	19
10	Biallelic variants in FBXL3 cause intellectual disability, delayed motor development and short stature. Human Molecular Genetics, 2019, 28, 972-979.	2.9	17
11	Molecular Dynamics Studies on Amyloidogenic Proteins. Springer Series on Bio- and Neurosystems, 2019, , 467-499.	0.2	0
12	Biallelic variants in KIF14 cause intellectual disability with microcephaly. European Journal of Human Genetics, 2018, 26, 330-339.	2.8	52
13	Biallelic variants in LINGO1 are associated with autosomal recessive intellectual disability, microcephaly, speech and motor delay. Genetics in Medicine, 2018, 20, 778-784.	2.4	21
14	Design of short peptides to block BTLA/HVEM interactions for promoting anticancer T-cell responses. PLoS ONE, 2017, 12, e0179201.	2.5	28
15	Activating mutations in genes related to TCR signaling in angioimmunoblastic and other follicular helper T-cell–derived lymphomas. Blood, 2016, 128, 1490-1502.	1.4	255
16	Design and Validation of a Novel Generic Platform for the Production of Tetravalent IgG1-like Bispecific Antibodies. Journal of Immunology, 2016, 196, 3199-3211.	0.8	30
17	Bicc1 Polymerization Regulates the Localization and Silencing of Bound mRNA. Molecular and Cellular Biology, 2015, 35, 3339-3353.	2.3	27
18	The CAP1/Prss8 catalytic triad is not involved in PAR2 activation and protease nexinâ€1 (PNâ€1) inhibition. FASEB lournal, 2014, 28, 4792-4805.	0.5	13

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19	Molecular Dynamics Studies on Amyloidogenic Proteins. Springer Series in Bio-/neuroinformatics, 2014, , 445-481.	0.1	2
20	RNA pentaloop structures as effective targets of regulators belonging to the RsmA/CsrA protein family. RNA Biology, 2013, 10, 1030-1041.	3.1	37
21	Monoubiquitination and Activity of the Paracaspase MALT1 Requires Glutamate 549 in the Dimerization Interface. PLoS ONE, 2013, 8, e72051.	2.5	25
22	Mutations of the Serine Protease CAP1/Prss8 Lead to Reduced Embryonic Viability, Skin Defects, and Decreased ENaC Activity. American Journal of Pathology, 2012, 181, 605-615.	3.8	36
23	Identification of human IKK-2 inhibitors of natural origin (Part II): In Silico prediction of IKK-2 inhibitors in natural extracts with known anti-inflammatory activity. European Journal of Medicinal Chemistry, 2011, 46, 6098-6103.	5.5	22
24	The V-ATPase proteolipid cylinder promotes the lipid-mixing stage of SNARE-dependent fusion of yeast vacuoles. EMBO Journal, 2011, 30, 4126-4141.	7.8	78
25	Identification of Human IKK-2 Inhibitors of Natural Origin (Part I): Modeling of the IKK-2 Kinase Domain, Virtual Screening and Activity Assays. PLoS ONE, 2011, 6, e16903.	2.5	23
26	A Combined Computational and Functional Approach Identifies New Residues Involved in pH-dependent Gating of ASIC1a. Journal of Biological Chemistry, 2010, 285, 16315-16329.	3.4	66
27	The role of the Val57 aminoâ€acid residue in the hinge loop of the human cystatin C. Conformational studies of the beta2â€L1â€beta3 segments of wildâ€type human cystatin C and its mutants. Biopolymers, 2009, 91, 373-383.	2.4	24
28	The Fourth Extracellular Loop of the α Subunit of Na,K-ATPase. Journal of Biological Chemistry, 2008, 283, 27850-27858.	3.4	6