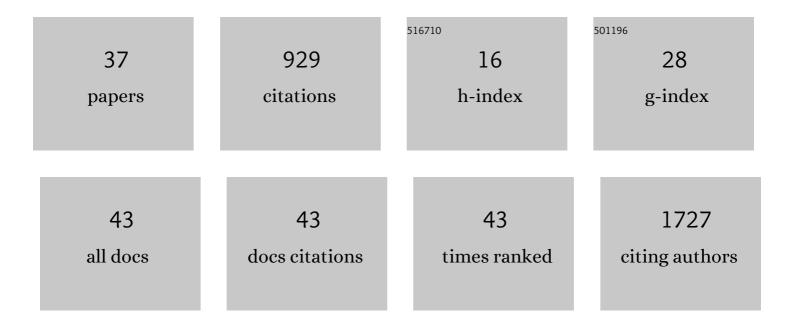
Dominik Cysewski

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Lipid droplets in skeletal muscle during grass snake (Natrix natrix L.) development. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2022, 1867, 159086.	2.4	3
2	4-1BBL–containing leukemic extracellular vesicles promote immunosuppressive effector regulatory T cells. Blood Advances, 2022, 6, 1879-1894.	5.2	13
3	Osteopontin—A Potential Biomarker for IgA Nephropathy: Machine Learning Application. Biomedicines, 2022, 10, 734.	3.2	1
4	Mechanisms of Resistance to Photodynamic Therapy (PDT) in Vulvar Cancer. International Journal of Molecular Sciences, 2022, 23, 4117.	4.1	4
5	A heterotypic assembly mechanism regulates <scp>CHIP E3</scp> ligase activity. EMBO Journal, 2022, 41,	7.8	9
6	Mass-spectrometric identification of oxidative modifications in plasma-purified plasminogen: Association with hypofibrinolysis in patients with acute pulmonary embolism. Biochemical and Biophysical Research Communications, 2022, 621, 53-58.	2.1	3
7	Cytoplasmic polyadenylation by TENT5A is required for proper bone formation. Cell Reports, 2021, 35, 109015.	6.4	24
8	Proteome-Wide Analysis of Protein Lysine <i>N</i> -Homocysteinylation in <i>Saccharomyces cerevisiae</i> . Journal of Proteome Research, 2021, 20, 2458-2476.	3.7	4
9	Diamond Nanofilm Normalizes Proliferation and Metabolism in Liver Cancer Cells. Nanotechnology, Science and Applications, 2021, Volume 14, 115-137.	4.6	3
10	Butyrylcholinesterase–Protein Interactions in Human Serum. International Journal of Molecular Sciences, 2021, 22, 10662.	4.1	8
11	Distinct characteristics of multisystem inflammatory syndrome in children in Poland. Scientific Reports, 2021, 11, 23562.	3.3	24
12	Keap1 controls protein S-nitrosation and apoptosis-senescence switch in endothelial cells. Redox Biology, 2020, 28, 101304.	9.0	22
13	Identification of glycated and acetylated lysine residues in human α2-antiplasmin. Biochemical and Biophysical Research Communications, 2020, 521, 19-23.	2.1	3
14	Functional analysis and cryo-electron microscopy of <i>Campylobacter jejuni</i> serine protease HtrA. Gut Microbes, 2020, 12, 1810532.	9.8	12
15	Mitochondrial protein biogenesis in the synapse is supported by local translation. EMBO Reports, 2020, 21, e48882.	4.5	63
16	Similar but Not Identical—Binding Properties of LSU (Response to Low Sulfur) Proteins From Arabidopsis thaliana. Frontiers in Plant Science, 2020, 11, 1246.	3.6	15
17	P0489URINARY PROTEOMIC MARKERS OF MEMBRANOUS NEPHROPATHY. Nephrology Dialysis Transplantation, 2020, 35, .	0.7	0
18	Keap1 governs ageing-induced protein aggregation in endothelial cells. Redox Biology, 2020, 34, 101572.	9.0	16

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19	Overexpression of the Selective Autophagy Cargo Receptor NBR1 Modifies Plant Response to Sulfur Deficit. Cells, 2020, 9, 669.	4.1	18
20	HaDeX: an R package and web-server for analysis of data from hydrogen–deuterium exchange mass spectrometry experiments. Bioinformatics, 2020, 36, 4516-4518.	4.1	13
21	Effects of Graphene Oxide Nanofilm and Chicken Embryo Muscle Extract on Muscle Progenitor Cell Differentiation and Contraction. Molecules, 2020, 25, 1991.	3.8	11
22	Tunneling nanotube-mediated intercellular vesicle and protein transfer in the stroma-provided imatinib resistance in chronic myeloid leukemia cells. Cell Death and Disease, 2019, 10, 817.	6.3	59
23	Quantitative proteomics revealed C6orf203/MTRES1 as a factor preventing stress-induced transcription deficiency in human mitochondria. Nucleic Acids Research, 2019, 47, 7502-7517.	14.5	21
24	Inhibition of proteasome rescues a pathogenic variant of respiratory chain assembly factor COA7. EMBO Molecular Medicine, 2019, 11, .	6.9	59
25	Synergy between the alteration in the N-terminal region of butyrylcholinesterase K variant and apolipoprotein E4 in late-onset Alzheimer's disease. Scientific Reports, 2019, 9, 5223.	3.3	14
26	Human dihydrofolate reductase is a substrate of protein kinase CK2α. Biochemical and Biophysical Research Communications, 2019, 513, 368-373.	2.1	6
27	Nrf2 Sequesters Keap1 Preventing Podosome Disassembly: A Quintessential Duet Moonlights in Endothelium. Antioxidants and Redox Signaling, 2019, 30, 1709-1730.	5.4	16
28	ISG'ylation increases stability of numerous proteins including Stat1, which prevents premature termination of immune response in LPS-stimulated microglia. Neurochemistry International, 2018, 112, 227-233.	3.8	20
29	Controlling the mitochondrial antisense – role of the SUV3-PNPase complex and its co-factor GRSF1 in mitochondrial RNA surveillance. Molecular and Cellular Oncology, 2018, 5, e1516452.	0.7	14
30	Dedicated surveillance mechanism controls G-quadruplex forming non-coding RNAs in human mitochondria. Nature Communications, 2018, 9, 2558.	12.8	67
31	The non-canonical poly(A) polymerase FAM46C acts as an onco-suppressor in multiple myeloma. Nature Communications, 2017, 8, 619.	12.8	77
32	A short splicing isoform of HBS1L links the cytoplasmic exosome and SKI complexes in humans. Nucleic Acids Research, 2016, 45, gkw862.	14.5	40
33	hUTP24 is essential for processing of the human rRNA precursor at site A ₁ , but not at site A ₀ . RNA Biology, 2015, 12, 1010-1029.	3.1	24
34	Linear mtDNA fragments and unusual mtDNA rearrangements associated with pathological deficiency of MGME1 exonuclease. Human Molecular Genetics, 2014, 23, 6147-6162.	2.9	64
35	Identification of Protein Partners in Mycobacteria Using a Single-Step Affinity Purification Method. PLoS ONE, 2014, 9, e91380.	2.5	20
36	Exonuclease hDIS3L2 specifies an exosome-independent 3′-5′ degradation pathway of human cytoplasmic mRNA. EMBO Journal, 2013, 32, 1855-1868.	7.8	136

#	Article	IF	CITATIONS
37	A new strategy for gene targeting and functional proteomics using the DT40 cell line. Nucleic Acids Research, 2013, 41, e167-e167.	14.5	17