Monika Dzieciatkowska

List of Publications by Year in descending order

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39 papers 1,741 citations

394421 19 h-index 302126 39 g-index

44 all docs

44 docs citations

44 times ranked 2827 citing authors

#	Article	IF	CITATIONS
1	Erythrocyte transglutaminase-2 combats hypoxia and chronic kidney disease by promoting oxygen delivery and carnitine homeostasis. Cell Metabolism, 2022, 34, 299-316.e6.	16.2	28
2	Specialized interferon action in COVID-19. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119 , .	7.1	56
3	DNA damage contributes to neurotoxic inflammation in Aicardi-Goutières syndrome astrocytes. Journal of Experimental Medicine, 2022, 219, .	8.5	35
4	p97 dysfunction underlies a loss of quality control of damaged membrane proteins and promotes oxidative stress and sickling in sickle cell disease. FASEB Journal, 2022, 36, e22246.	0.5	5
5	A Designer Nanoparticle Platform for Controlled Intracellular Delivery of Bioactive Macromolecules: Inhibition of Ubiquitin-Specific Protease 7 in Breast Cancer Cells. ACS Chemical Biology, 2022, 17, 1853-1865.	3.4	3
6	Matrix reverses immortalization-mediated stem cell fate determination. Biomaterials, 2021, 265, 120387.	11.4	15
7	Seroconversion stages COVID19 into distinct pathophysiological states. ELife, 2021, 10, .	6.0	40
8	Targeted Intracellular Delivery of Trastuzumab Using Designer Phage Lambda Nanoparticles Alters Cellular Programs in Human Breast Cancer Cells. ACS Nano, 2021, 15, 11789-11805.	14.6	18
9	Red Blood Cell Proteasome in Beta-Thalassemia Trait: Topology of Activity and Networking in Blood Bank Conditions. Membranes, 2021, 11, 716.	3.0	11
10	The Post-Storage Performance of RBCs from Beta-Thalassemia Trait Donors Is Related to Their Storability Profile. International Journal of Molecular Sciences, 2021, 22, 12281.	4.1	8
11	Cytoplasmic dynein-1 cargo diversity is mediated by the combinatorial assembly of FTS–Hook–FHIP complexes. ELife, 2021, 10, .	6.0	27
12	Extracellular vesicles from young women's breast cancer patients drive increased invasion of non-malignant cells via the Focal Adhesion Kinase pathway: a proteomic approach. Breast Cancer Research, 2020, 22, 128.	5.0	21
13	Serum Proteomics in COVID-19 Patients: Altered Coagulation and Complement Status as a Function of IL-6 Level. Journal of Proteome Research, 2020, 19, 4417-4427.	3.7	155
14	Evidence of Structural Protein Damage and Membrane Lipid Remodeling in Red Blood Cells from COVID-19 Patients. Journal of Proteome Research, 2020, 19, 4455-4469.	3.7	189
15	Site-Dependent Lineage Preference of Adipose Stem Cells. Frontiers in Cell and Developmental Biology, 2020, 8, 237.	3.7	13
16	Red cell proteasome modulation by storage, redox metabolism and transfusion. Blood Transfusion, 2020, , .	0.4	7
17	Hook3 is a scaffold for the opposite-polarity microtubule-based motors cytoplasmic dynein-1 and KIF1C. Journal of Cell Biology, 2019, 218, 2982-3001.	5.2	57
18	Methylation of protein aspartates and deamidated asparagines as a function of blood bank storage and oxidative stress in human red blood cells. Transfusion, 2018, 58, 2978-2991.	1.6	71

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19	A comparison of different methods of red blood cell leukoreduction and additive solutions on the accumulation of neutrophilâ€priming activity during storage. Transfusion, 2018, 58, 2003-2012.	1.6	7
20	Correlation of preâ€operative plasma protein concentrations in cardiac surgery patients with bleeding outcomes using a targeted quantitative proteomics approach. Proteomics - Clinical Applications, 2017, 11, 1600175.	1.6	5
21	The Immunosuppressant Mycophenolic Acid Alters Nucleotide and Lipid Metabolism in an Intestinal Cell Model. Scientific Reports, 2017, 7, 45088.	3.3	19
22	Hydroxylamine Chemical Digestion for Insoluble Extracellular Matrix Characterization. Journal of Proteome Research, 2017, 16, 4177-4184.	3.7	52
23	Blood and Plasma Proteomics: Targeted Quantitation and Posttranslational Redox Modifications. Methods in Molecular Biology, 2017, 1619, 353-371.	0.9	7
24	Biophysical analysis of the effect of chemical modification by 4-oxononenal on the structure, stability, and function of binding immunoglobulin protein (BiP). PLoS ONE, 2017, 12, e0183975.	2.5	4
25	CD147: a small molecule transporter ancillary protein at the crossroad of multiple hallmarks of cancer and metabolic reprogramming. Oncotarget, 2017, 8, 6742-6762.	1.8	36
26	Red blood cell proteomics update: is there more to discover?. Blood Transfusion, 2017, 15, 182-187.	0.4	76
27	Oxidative modifications of glyceraldehyde 3-phosphate dehydrogenase regulate metabolic reprogramming of stored red blood cells. Blood, 2016, 128, e32-e42.	1.4	183
28	Hypertonic Saline Primes Activation of the p53–p21 Signaling Axis in Human Small Airway Epithelial Cells That Prevents Inflammation Induced by Pro-inflammatory Cytokines. Journal of Proteome Research, 2016, 15, 3813-3826.	3.7	11
29	Monoubiquitination of survival motor neuron regulates its cellular localization and Cajal body integrity. Human Molecular Genetics, 2016, 25, 1392-1405.	2.9	18
30	Preserved Proteins from Extinct Bison latifrons Identified by Tandem Mass Spectrometry; Hydroxylysine Glycosides are a Common Feature of Ancient Collagen. Molecular and Cellular Proteomics, 2015, 14, 1946-1958.	3.8	73
31	Plasma QconCATs reveal a gender-specific proteomic signature in apheresis platelet plasma supernatants. Journal of Proteomics, 2015, 120, 1-6.	2.4	32
32	Lin28 promotes the proliferative capacity of neural progenitor cells in brain development. Development (Cambridge), 2015, 142, 1616-1627.	2 . 5	109
33	The cerebrospinal fluid immunoglobulin transcriptome and proteome in neuromyelitis optica reveals central nervous system-specific B cell populations. Journal of Neuroinflammation, 2015, 12, 19.	7.2	48
34	Quantification of Extracellular Matrix Proteins from a Rat Lung Scaffold to Provide a Molecular Readout for Tissue Engineering. Molecular and Cellular Proteomics, 2015, 14, 961-973.	3.8	131
35	Phosphorylation of the Scc2 cohesin deposition complex subunit regulates chromosome condensation through cohesin integrity. Molecular Biology of the Cell, 2015, 26, 3754-3767.	2.1	8
36	Proteomics of apheresis platelet supernatants during routine storage: Gender-related differences. Journal of Proteomics, 2015, 112, 190-209.	2.4	23

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37	Cell cycle-specific cleavage of Scc2 regulates its cohesin deposition activity. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 7060-7065.	7.1	13
38	Toward the identification of a subset of unexplained infertility: a sperm proteomic approach. Fertility and Sterility, 2014, 102, 692-699.	1.0	57
39	GeLC-MS/MS Analysis of Complex Protein Mixtures. Methods in Molecular Biology, 2014, 1156, 53-66.	0.9	64