Joanna Szczepanek

List of Publications by Year in descending order

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759233 839539 34 384 12 18 citations h-index g-index papers 37 37 37 490 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Analysis of profibrogenic microRNAs (miRNAs) expression in urine and serum of chronic kidney disease (CKD) stage 1–4 patients and their relationship with proteinuria and kidney function. International Urology and Nephrology, 2022, 54, 937-947.	1.4	13
2	Anti-SARS-CoV-2 IgG against the S Protein: A Comparison of BNT162b2, mRNA-1273, ChAdOx1 nCoV-2019 and Ad26.COV2.S Vaccines. Vaccines, 2022, 10, 99.	4.4	13
3	The Dynamics of Changes in the Concentration of IgG against the S1 Subunit in Polish Healthcare Workers in the Period from 1 to 12 Months after Injection, Including Four COVID-19 Vaccines. Vaccines, 2022, 10, 506.	4.4	4
4	MicroRNA as a Potential Therapeutic Molecule in Cancer. Cells, 2022, 11, 1008.	4.1	44
5	The Effect of Platelet-Rich Plasma on the Intra-Articular Microenvironment in Knee Osteoarthritis. International Journal of Molecular Sciences, 2021, 22, 5492.	4.1	54
6	Salt stress and salt shock differently affect DNA methylation in salt-responsive genes in sugar beet and its wild, halophytic ancestor. PLoS ONE, 2021, 16, e0251675.	2.5	14
7	Differences in the Concentration of Anti-SARS-CoV-2 IgG Antibodies Post-COVID-19 Recovery or Post-Vaccination. Cells, 2021, 10, 1952.	4.1	58
8	Genetics in Cartilage Lesions: Basic Science and Therapy Approaches. International Journal of Molecular Sciences, 2020, 21, 5430.	4.1	21
9	Role of microRNA dysregulation in childhood acute leukemias: diagnostics, monitoring and therapeutics: A comprehensive review. World Journal of Clinical Oncology, 2020, 11, 348-369.	2.3	18
10	Data set for transcriptome analysis of liver in cattle breeds. Translational Research in Veterinary Science, 2020, 2, 51.	0.1	0
11	New diagnostic and therapeutic options in canine osteoarthritis. Medycyna Weterynaryjna, 2020, 76, 6372-2020.	0.1	0
12	Data set for transcriptome analysis of pituitary galnd in cattle breeds. Translational Research in Veterinary Science, 2020, 2, 57.	0.1	0
13	Comparative Analysis of the Liver Transcriptome among Cattle Breeds Using RNA-seq. Veterinary Sciences, 2019, 6, 36.	1.7	8
14	RNA-seq based SNP discovery in gluteus medius muscle of Polish Landrace pigs. Translational Research in Veterinary Science, 2019, 2, 51.	0.1	0
15	RNA-seq based SNP discovery in liver transcriptome of Polish Landrace pigs. Translational Research in Veterinary Science, 2019, 2, 67.	0.1	0
16	The role of microRNAs in animal physiology and pathology. Translational Research in Veterinary Science, 2018, 1, 13.	0.1	2
17	New Candidate Genes for Lack of Sensitivity to Therapy in Pediatric Leukemias. Current Cancer Drug Targets, 2017, 17, 333-343.	1.6	4
18	Genomic and transcriptomic profiles and <i>in vitro</i> resistance to mitoxantrone and idarubicin in pediatric acute leukemias. Journal of Gene Medicine, 2016, 18, 165-179.	2.8	6

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19	An analysis of the expression of collagen I and III genes in the fascia of obese patients. Journal of Surgical Research, 2015, 195, 475-480.	1.6	12
20	The expression patterns of plasma membrane aquaporins in leaves of sugar beet and its halophyte relative, Beta vulgaris ssp. maritima, in response to salt stress. Biologia (Poland), 2015, 70, 467-477.	1.5	13
21	Whole Genome and Transcriptome Analysis of Genetic Alterations in Context of Busulfan in Vitro Resistance in Pediatric Acute Leukemias. Blood, 2015, 126, 4849-4849.	1.4	1
22	Identification of the Genomic Rearrangements Associated with the Ex Vivo Resistance to Anthracyclines in Childhood Acute Leukemias. Blood, 2015, 126, 4997-4997.	1.4	1
23	Differential in Vitro Drug Resistance Profile Between First and Second Relapsed Acute Lymphoblastic Leukemia and Acute Myeloblastic Leukemia in Children. Blood, 2015, 126, 3696-3696.	1.4	0
24	Array-Based Genomic and Transcriptomic Profiling in Pediatric Acute Leukemias in Search of Genes Responsible for Cyclophosphamide Resistance. Blood, 2014, 124, 3802-3802.	1.4	1
25	Array Comparative Genomic Hybridization in Pediatric Acute Leukemias. Pediatric Hematology and Oncology, 2013, 30, 677-687.	0.8	2
26	Analysis of Relative Expression Level of VEGF (Vascular Endothelial Growth Factor), HIF-1a (Hypoxia) Tj ETQqO 0 Glomerulonephritis (CGN) Patients. Kidney and Blood Pressure Research, 2013, 38, 83-91.	0 rgBT /Ov 2.0	erlock 10 Tf 5 14
27	Genetic Mechanisms Of Cytarabine, Etoposide and Daunorubicin Resistance In Pediatric Acute Leukemias. Blood, 2013, 122, 4938-4938.	1.4	0
28	Gene expression signatures and ex vivo drug sensitivity profiles in children with acute lymphoblastic leukemia. Journal of Applied Genetics, 2012, 53, 83-91.	1.9	8
29	Identification of the genes expression profile associated with the ex vivo resistance to etoposide in childhood acute leukemias. Postepy Higieny I Medycyny Doswiadczalnej, 2012, 66, 401-408.	0.1	4
30	Expression profiles of signal transduction genes in ex vivo drug-resistant pediatric acute lymphoblastic leukemia. Anticancer Research, 2012, 32, 503-6.	1.1	28
31	Relapse of Acute Lymphoblastic Leukemia in Children in the Context of Microarray Analyses. Archivum Immunologiae Et Therapiae Experimentalis, 2011, 59, 61-68.	2.3	22
32	Differential ex vivo activity of bortezomib in newly diagnosed paediatric acute lymphoblastic and myeloblastic leukaemia. Anticancer Research, 2010, 30, 2119-24.	1.1	13
33	Gene Expression and Ex Vivo Drug Sensitivity Profiles in Children with Acute Leukemia: An Insight for High-Dose Therapy and Role of Etoposide and Fludarabine Blood, 2009, 114, 2747-2747.	1.4	0
34	Prognostic impact of combined fludarabine, treosulfan and mitoxantrone resistance profile in childhood acute myeloid leukemia. Anticancer Research, 2008, 28, 1927-31.	1.1	3