NataÅja M ToÅjić

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Expression Profiles of Long Non-Coding RNA GAS5 and MicroRNA-222 in Younger AML Patients. Diagnostics, 2022, 12, 86.	2.6	4
2	Prognostic significance of combined BAALC and MN1 gene expression level in acute myeloid leukemia with normal karyotype. International Journal of Laboratory Hematology, 2021, 43, 433-440.	1.3	1
3	Comparative analysis of international prognostic index for chronic lymphocytic leukemia, progression-risk score, and MD Anderson Cancer Center 2011 score - a single center experience. Srpski Arhiv Za Celokupno Lekarstvo, 2021, 149, 415-421.	0.2	Ο
4	Association of SLC28A3 Gene Expression and CYP2B6*6 Allele with the Response to Fludarabine Plus Cyclophosphamide in Chronic Lymphocytic Leukemia Patients. Pathology and Oncology Research, 2020, 26, 743-752.	1.9	2
5	Expression Pattern and Prognostic Significance of EVI1 Gene in Adult Acute Myeloid Leukemia Patients with Normal Karyotype. Indian Journal of Hematology and Blood Transfusion, 2020, 36, 292-299.	0.6	6
6	The influence of Wilms' tumor 1 gene expression level on prognosis and risk stratification of acute promyelocytic leukemia patients. International Journal of Laboratory Hematology, 2020, 42, 82-87.	1.3	7
7	Expression pattern of circulating long non-coding RNA GAS5 as a novel biomarker in non-small cell lung cancer patients. Archives of Medical Science, 2020, , .	0.9	3
8	Complex transcriptional regulation of the BCL2L12 gene: Novel, active promoter in K562 cells. Gene, 2020, 750, 144723.	2.2	4
9	Genomic profiling of thymoma using a targeted high-throughput approach. Archives of Medical Science, 2020, , .	0.9	2
10	Genetic and Epigenetic Profiling in Personalized Medicine: Advances in Treatment of Acute Myeloid Leukemia. Europeanization and Globalization, 2019, , 341-374.	0.1	0
11	CRISPR/Cas9 genome editing of SLC37A4 gene elucidates the role of molecular markers of endoplasmic reticulum stress and apoptosis in renal involvement in glycogen storage disease type lb. Gene, 2019, 703, 17-25.	2.2	5
12	Application of targeted next generation sequencing for the mutational profiling of patients with acute lymphoblastic leukemia. Journal of Medical Biochemistry, 2019, 39, 72-82.	1.7	0
13	Prognostic significance of SOX2, SOX3, SOX11, SOX14 and SOX18 gene expression in adult de novo acute myeloid leukemia. Leukemia Research, 2018, 67, 32-38.	0.8	17
14	Impact of alterations in X-linked IRAK1gene and miR-146a on susceptibility and clinical manifestations in patients with systemic sclerosis. Immunology Letters, 2018, 204, 1-8.	2.5	12
15	The predictive value of morphological findings in early diagnosis of acute myeloid leukemia with recurrent cytogenetic abnormalities. Leukemia Research, 2018, 75, 23-28.	0.8	7
16	Use of Wilms Tumor 1 Gene Expression as a Reliable Marker for Prognosis and Minimal Residual Disease Monitoring in Acute Myeloid Leukemia With Normal Karyotype Patients. Clinical Lymphoma, Myeloma and Leukemia, 2017, 17, 312-319.	0.4	26
17	Gene Mutation Profiles in Primary Diffuse Large B Cell Lymphoma of Central Nervous System: Next Generation Sequencing Analyses. International Journal of Molecular Sciences, 2016, 17, 683.	4.1	29
18	Effects of DMSO, glycerol, betaine and their combinations in detecting single nucleotide polymorphisms of epidermal growth factor receptor (EGFR) gene promoter sequence in non-small-cell lung cancer (NSCLC) patients. Journal of Pharmaceutical and Biomedical Analysis, 2016, 128, 275-279.	2.8	6

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19	Somatic mutations of isocitrate dehydrogenases 1 and 2 are prognostic and follow-up markers in patients with acute myeloid leukaemia with normal karyotype. Radiology and Oncology, 2016, 50, 385-393.	1.7	16
20	Parallel targeted next generation sequencing of childhood and adult acute myeloid leukemia patients reveals uniform genomic profile of the disease. Tumor Biology, 2016, 37, 13391-13401.	1.8	13
21	Prognostic Significance of Cereblon Expression in Patients With Multiple Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2016, 16, 610-615.	0.4	12
22	Frequencies of EGFR single nucleotide polymorphisms in non-small cell lung cancer patients and healthy individuals in the Republic of Serbia: a preliminary study. Tumor Biology, 2016, 37, 10479-10486.	1.8	13
23	Association of Bax Expression and Bcl2/Bax Ratio with Clinical and Molecular Prognostic Markers in Chronic Lymphocytic Leukemia. Journal of Medical Biochemistry, 2016, 35, 150-157.	1.7	46
24	Expression Of The <i>Bcl2</i> Gene In Chronic Lymphocytic Leukaemia Patients. Serbian Journal of Experimental and Clinical Research, 2015, 16, 187-191.	0.1	1
25	Expression of bacteriocin LsbB is dependent on a transcription terminator. Microbiological Research, 2015, 179, 45-53.	5.3	19
26	<i>TPMT</i> gene expression is increased during maintenance therapy in childhood acute lymphoblastic leukemia patients in a <i>TPMT</i> gene promoter variable number of tandem repeat-dependent manner. Pharmacogenomics, 2015, 16, 1701-1712.	1.3	21
27	Cancer Monitoring Methods. BioMed Research International, 2014, 2014, 1-1.	1.9	0
28	Expression of Bcl2L12 in chronic lymphocytic leukemia patients: association with clinical and molecular prognostic markers. Medical Oncology, 2013, 30, 405.	2.5	16
29	Molecular Genetics and Genotype-Based Estimation of BH4-Responsiveness in Serbian PKU Patients: Spotlight on Phenotypic Implications of p.L48S. JIMD Reports, 2012, 9, 49-58.	1.5	22
30	Prognostic Impact of <i>NPM1</i> Mutations in Serbian Adult Patients with Acute Myeloid Leukemia. Acta Haematologica, 2012, 128, 203-212.	1.4	12
31	Overexpression of the novel member of the BCL2 gene family, BCL2L12, is associated with the disease outcome in patients with acute myeloid leukemia. Clinical Biochemistry, 2012, 45, 1362-1367.	1.9	10
32	Distinct Genetic Lesions Drive Leukemogenesis in Secondary Acute Myeloid Leukemia,. Blood, 2011, 118, 3559-3559.	1.4	15
33	Incidence of FLT3 and nucleophosmin gene mutations in childhood acute myeloid leukemia: Serbian experience and the review of the literature. Medical Oncology, 2010, 27, 640-645.	2.5	14
34	Functional analysis of the role of the <i>TPMT</i> gene promoter VNTR polymorphism in <i>TPMT</i> gene transcription. Pharmacogenomics, 2010, 11, 547-557.	1.3	40
35	Importance of early detection and follow-up of FLT3 mutations in patients with acute myeloid leukemia. Annals of Hematology, 2007, 86, 741-747.	1.8	31