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List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

432
citations

759233

12
h-index

794594

19
g-index

35
all docs

35
docs citations

35
times ranked

777
citing authors

#	ARTICLE	IF	CITATIONS
1	Expression Profiles of Long Non-Coding RNA GAS5 and MicroRNA-222 in Younger AML Patients. <i>Diagnostics</i> , 2022, 12, 86.	2.6	4
2	Prognostic significance of combined BAALC and MN1 gene expression level in acute myeloid leukemia with normal karyotype. <i>International Journal of Laboratory Hematology</i> , 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. <i>Srpski Arhiv Za Celokupno Lekarstvo</i> , 2021, 149, 415-421.	0.2	0
4	Association of SLC28A3 Gene Expression and CYP2B6*6 Allele with the Response to Fludarabine Plus Cyclophosphamide in Chronic Lymphocytic Leukemia Patients. <i>Pathology and Oncology Research</i> , 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. <i>Indian Journal of Hematology and Blood Transfusion</i> , 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. <i>International Journal of Laboratory Hematology</i> , 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. <i>Archives of Medical Science</i> , 2020, , .	0.9	3
8	Complex transcriptional regulation of the BCL2L12 gene: Novel, active promoter in K562 cells. <i>Gene</i> , 2020, 750, 144723.	2.2	4
9	Genomic profiling of thymoma using a targeted high-throughput approach. <i>Archives of Medical Science</i> , 2020, , .	0.9	2
10	Genetic and Epigenetic Profiling in Personalized Medicine: Advances in Treatment of Acute Myeloid Leukemia. <i>Europeanization and Globalization</i> , 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 Ib. <i>Gene</i> , 2019, 703, 17-25.	2.2	5
12	Application of targeted next generation sequencing for the mutational profiling of patients with acute lymphoblastic leukemia. <i>Journal of Medical Biochemistry</i> , 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. <i>Leukemia Research</i> , 2018, 67, 32-38.	0.8	17
14	Impact of alterations in X-linked IRAK1 gene and miR-146a on susceptibility and clinical manifestations in patients with systemic sclerosis. <i>Immunology Letters</i> , 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. <i>Leukemia Research</i> , 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. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 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. <i>International Journal of Molecular Sciences</i> , 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. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016, 128, 275-279.	2.8	6

