

Julia Starkova

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

477
citations

687363

13
h-index

713466

21
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all docs

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docs citations

27
times ranked

1024
citing authors

#	ARTICLE	IF	CITATIONS
1	PTEN/PI3K/Akt pathway alters sensitivity of T-cell acute lymphoblastic leukemia to L-asparaginase. <i>Scientific Reports</i> , 2022, 12, 4043.	3.3	8
2	Pancreatitis-associated protein as an early marker of asparaginase-associated pancreatitis. <i>Leukemia and Lymphoma</i> , 2021, , 1-5.	1.3	0
3	Transmembrane adaptor protein WBP1L regulates CXCR4 signalling and murine haematopoiesis. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 1980-1992.	3.6	6
4	Metabolic profile of leukemia cells influences treatment efficacy of L-asparaginase. <i>BMC Cancer</i> , 2020, 20, 526.	2.6	18
5	<i>ERG</i> deletions in childhood acute lymphoblastic leukemia with <i>DUX4</i> rearrangements are mostly polyclonal, prognostically relevant and their detection rate strongly depends on screening method sensitivity. <i>Haematologica</i> , 2019, 104, 1407-1416.	3.5	34
6	Genomic landscape of pediatric B-other acute lymphoblastic leukemia in a consecutive European cohort. <i>Haematologica</i> , 2019, 104, 1396-1406.	3.5	78
7	Two novel fusion genes, <i>AIF1</i> and <i>ABL1</i> , result together with <i>ETV6</i> from a single chromosomal rearrangement in acute lymphoblastic leukemia with prenatal origin. <i>Genes Chromosomes and Cancer</i> , 2018, 57, 471-477.	2.8	5
8	Low HOX gene expression in PML-RAR α -positive leukemia results from suppressed histone demethylation. <i>Epigenetics</i> , 2018, 13, 73-84.	2.7	16
9	Assessment of the Metabolic Profile of Primary Leukemia Cells. <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	1
10	Integrative analysis of transcriptomics and clinical data uncovers the tumor-suppressive activity of MITF in prostate cancer. <i>Cell Death and Disease</i> , 2018, 9, 1041.	6.3	14
11	Altered Metabolism of Leukemic Cells: New Therapeutic Opportunity. <i>International Review of Cell and Molecular Biology</i> , 2018, 336, 93-147.	3.2	8
12	A triad of NRP2, DLX and p53 proteins in lung cancer metastasis. <i>Oncotarget</i> , 2017, 8, 96464-96465.	1.8	4
13	Characterization of leukemias with ETV6-ABL1 fusion. <i>Haematologica</i> , 2016, 101, 1082-1093.	3.5	66
14	Distinct bilineal leukemia immunophenotypes are not genetically determined. <i>Blood</i> , 2016, 128, 2263-2266.	1.4	23
15	Intragenic ERG Deletions Do Not Explain the Biology of ERG-Related Acute Lymphoblastic Leukemia. <i>PLoS ONE</i> , 2016, 11, e0160385.	2.5	7
16	Characterization of Leukemias with ETV6-ABL1 Fusion. <i>Blood</i> , 2015, 126, 84-84.	1.4	1
17	Homeobox gene expression in acute myeloid leukemia is linked to typical underlying molecular aberrations. <i>Journal of Hematology and Oncology</i> , 2014, 7, 94.	17.0	14
18	The Role of Histone Demethylases in the Transcription Regulation of HOX Genes in PML-RAR α + AML Patients. <i>Blood</i> , 2014, 124, 876-876.	1.4	0

#	ARTICLE	IF	CITATIONS
19	Molecular Background of BCP-ALL Cases with an Early Switch to Monocytic Lineage. <i>Blood</i> , 2014, 124, 3562-3562.	1.4	0
20	Low expression of asparagine synthetase in lymphoid blasts precludes its role in sensitivity to L-asparaginase. <i>Experimental Hematology</i> , 2012, 40, 657-665.	0.4	36
21	Valproic acid triggers differentiation and apoptosis in AML1/ETO-positive leukemic cells specifically. <i>Cancer Letters</i> , 2012, 319, 144-153.	7.2	22
22	ETV6/RUNX1 (TEL/AML1) is a frequent prenatal first hit in childhood leukemia. <i>Blood</i> , 2011, 117, 368-369.	1.4	52
23	Up-regulation of homeodomain genes, DLX1 and DLX2, by FLT3 signaling. <i>Haematologica</i> , 2011, 96, 820-828.	3.5	19
24	<i>HOX</i> gene expression in phenotypic and genotypic subgroups and low <i>HOXA</i> gene expression as an adverse prognostic factor in pediatric ALL. <i>Pediatric Blood and Cancer</i> , 2010, 55, 1072-1082.	1.5	25
25	Expression Profile of Selected HOX Genes in Different Childhood ALL Subtypes. <i>Blood</i> , 2008, 112, 4462-4462.	1.4	0
26	The Identification of (ETV6)/RUNX1-Regulated Genes in Lymphopoiesis Using Histone Deacetylase Inhibitors in ETV6/RUNX1-Positive Lymphoid Leukemic Cells. <i>Clinical Cancer Research</i> , 2007, 13, 1726-1735.	7.0	19
27	Histone Deacetylase Inhibitors Are Capable to Modify Leukaemia-Specific Phenotype of TEL/AML1-Positive Leukaemic Cells.. <i>Blood</i> , 2004, 104, 1891-1891.	1.4	1