

Nina Dusilkova

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

212
citations

1684188

5
h-index

2053705

5
g-index

10
all docs

10
docs citations

10
times ranked

510
citing authors

#	ARTICLE	IF	CITATIONS
1	Analysis of 5-Azacytidine Resistance Models Reveals a Set of Targetable Pathways. <i>Cells</i> , 2022, 11, 223.	4.1	5
2	Randomized Open-Labeled Academic Trial Comparing Standard AZA Therapy with Combination of G-CSF with AZA in High Risk MDS Patients - Interim Analysis. <i>Blood</i> , 2019, 134, 1729-1729.	1.4	0
3	Plasma miR-155, miR-203, and miR-205 are Biomarkers for Monitoring of Primary Cutaneous T-Cell Lymphomas. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2136.	4.1	33
4	Somatic mutation dynamics in MDS patients treated with azacitidine indicate clonal selection in patients-responders. <i>Oncotarget</i> , 2017, 8, 111966-111978.	1.8	8
5	Clonal Architecture of MDS Somatic Mutations Dynamically Changes during Azacitidine Therapy and Has Very Limited Potential to Predict Patient Outcome. <i>Blood</i> , 2016, 128, 4294-4294.	1.4	0
6	Myristoylated Alanine-Rich C-Kinase Substrate (MARCKS) Is a New Biomarker for Mantle Cell Lymphoma: Expression, Localization, and Phosphorylation Study. <i>Blood</i> , 2016, 128, 1767-1767.	1.4	0
7	Oncogenic microRNA-155 and its target PU.1: an integrative gene expression study in six of the most prevalent lymphomas. <i>International Journal of Hematology</i> , 2015, 102, 441-450.	1.6	17
8	Oncogenic MicroRNAs: miR-155, miR-19a, miR-181b, and miR-24 enable monitoring of early breast cancer in serum. <i>BMC Cancer</i> , 2014, 14, 448.	2.6	149
9	Somatic Mutation-Detecting Algorithm Enables Analysis of MDS Patients during Azacitidine Therapy. <i>Blood</i> , 2014, 124, 5600-5600.	1.4	0
10	Erythroid Transcription Factor GATA-1 Binds and Represses PU.1 Gene – Candidate Mechanism Of Epigenetic Repression Of PU.1 and Inefficient Erythropoiesis In MDS. <i>Blood</i> , 2013, 122, 1558-1558.	1.4	0