

Laura Mosca

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

1,219
citations

516710

16
h-index

642732

23
g-index

23
all docs

23
docs citations

23
times ranked

2240
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of microRNA expression patterns and definition of a microRNA/mRNA regulatory network in distinct molecular groups of multiple myeloma. <i>Blood</i> , 2009, 114, e20-e26.	1.4	224
2	A SNP microarray and FISH-based procedure to detect allelic imbalances in multiple myeloma: An integrated genomics approach reveals a wide gene dosage effect. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 603-614.	2.8	134
3	An integrative genomic approach reveals coordinated expression of intronic miR-335, miR-342, and miR-561 with deregulated host genes in multiple myeloma. <i>BMC Medical Genomics</i> , 2008, 1, 37.	1.5	104
4	Biological and Clinical Relevance of miRNA Expression Signatures in Primary Plasma Cell Leukemia. <i>Clinical Cancer Research</i> , 2013, 19, 3130-3142.	7.0	86
5	Small nucleolar RNAs as new biomarkers in chronic lymphocytic leukemia. <i>BMC Medical Genomics</i> , 2013, 6, 27.	1.5	73
6	Integrative high-resolution microarray analysis of human myeloma cell lines reveals deregulated miRNA expression associated with allelic imbalances and gene expression profiles. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 521-531.	2.8	60
7	Genome-wide analysis of primary plasma cell leukemia identifies recurrent imbalances associated with changes in transcriptional profiles. <i>American Journal of Hematology</i> , 2013, 88, 16-23.	4.1	60
8	Clinical Monoclonal B Lymphocytosis versus Rai 0 Chronic Lymphocytic Leukemia: A Comparison of Cellular, Cytogenetic, Molecular, and Clinical Features. <i>Clinical Cancer Research</i> , 2013, 19, 5890-5900.	7.0	60
9	Molecular and transcriptional characterization of 17p loss in B-cell chronic lymphocytic leukemia. <i>Genes Chromosomes and Cancer</i> , 2008, 47, 781-793.	2.8	59
10	Whole-exome sequencing of primary plasma cell leukemia discloses heterogeneous mutational patterns. <i>Oncotarget</i> , 2015, 6, 17543-17558.	1.8	55
11	Integrative Genomics Analyses Reveal Molecularly Distinct Subgroups of B-Cell Chronic Lymphocytic Leukemia Patients with 13q14 Deletion. <i>Clinical Cancer Research</i> , 2010, 16, 5641-5653.	7.0	52
12	High-throughput sequencing for the identification of NOTCH1 mutations in early stage chronic lymphocytic leukaemia: biological and clinical implications. <i>British Journal of Haematology</i> , 2014, 165, 629-639.	2.5	52
13	Transcriptional Characterization of a Prospective Series of Primary Plasma Cell Leukemia Revealed Signatures Associated with Tumor Progression and Poorer Outcome. <i>Clinical Cancer Research</i> , 2013, 19, 3247-3258.	7.0	50
14	Chromosome 2p gain in monoclonal B-cell lymphocytosis and in early stage chronic lymphocytic leukemia. <i>American Journal of Hematology</i> , 2013, 88, 24-31.	4.1	27
15	Association between gene and miRNA expression profiles and stereotyped subset #4 B-cell receptor in chronic lymphocytic leukemia. <i>Leukemia and Lymphoma</i> , 2015, 56, 3150-3158.	1.3	23
16	Molecular and transcriptional characterization of the novel 17p11.2-p12 amplicon in multiple myeloma. <i>Genes Chromosomes and Cancer</i> , 2007, 46, 1109-1118.	2.8	20
17	Repetitive DNA Hypomethylation in Multiple Myeloma. <i>Blood</i> , 2008, 112, 2703-2703.	1.4	16
18	Insulin Growth Factor 1 Receptor Expression Is Associated with NOTCH1 Mutation, Trisomy 12 and Aggressive Clinical Course in Chronic Lymphocytic Leukaemia. <i>PLoS ONE</i> , 2015, 10, e0118801.	2.5	15

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19	Prospective validation of a risk score based on biological markers for predicting progression free survival in Binet stage A chronic lymphocytic leukemia patients: Results of the multicenter Oâ€CCLL1â€CIGSL study. American Journal of Hematology, 2014, 89, 743-750.	4.1	14
20	Surrogate molecular markers for IGHV mutational status in chronic lymphocytic leukemia for predicting time to first treatment. Leukemia Research, 2015, 39, 840-845.	0.8	12
21	Distinct patterns of global promoter methylation in early stage chronic lymphocytic leukemia. Genes Chromosomes and Cancer, 2014, 53, 264-273.	2.8	10
22	Prognostic Significance of Telomere Length in Chronic Lymphocytic Leukemia Patients in Early Stage Disease,. Blood, 2011, 118, 3890-3890.	1.4	7
23	Molecular events underlying interleukinâ€6 independence in a subclone of the CMAâ€03 multiple myeloma cell line. Genes Chromosomes and Cancer, 2014, 53, 154-167.	2.8	6