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	Whole-exome sequencing of primary plasma cell leukemia discloses heterogeneous mutational patterns. Oncotarget, 2015, 6, 17543-17558.	1.8	55
2	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
3	Association between gene and miRNA expression profiles and stereotyped subset #4 B-cell receptor in chronic lymphocytic leukemia. Leukemia and Lymphoma, 2015, 56, 3150-3158.	1.3	23
4	Insulin Growth Factor 1 Receptor Expression Is Associated with NOTCH1 Mutation, Trisomy 12 and Aggressive Clinical Course in Chronic Lymphocytic Leukaemia. PLoS ONE, 2015, 10, e0118801.	2.5	15
5	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€CLL1â€GISL study. American Journal of Hematology, 2014, 89, 743-750.	4.1	14
6	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
7	Distinct patterns of global promoter methylation in early stage chronic lymphocytic leukemia. Genes Chromosomes and Cancer, 2014, 53, 264-273.	2.8	10
8	Highâ€throughput sequencing for the identification of <scp>NOTCH</scp>1 mutations in early stage chronic lymphocytic leukaemia: biological and clinical implications. British Journal of Haematology, 2014, 165, 629-639.	2.5	52
9	Small nucleolar RNAs as new biomarkers in chronic lymphocytic leukemia. BMC Medical Genomics, 2013, 6, 27.	1.5	73
10	Genomeâ€wide analysis of primary plasma cell leukemia identifies recurrent imbalances associated with changes in transcriptional profiles. American Journal of Hematology, 2013, 88, 16-23.	4.1	60
11	Chromosome 2p gain in monoclonal Bâ€cell lymphocytosis and in early stage chronic lymphocytic leukemia. American Journal of Hematology, 2013, 88, 24-31.	4.1	27
12	Clinical Monoclonal B Lymphocytosis versus Rai 0 Chronic Lymphocytic Leukemia: A Comparison of Cellular, Cytogenetic, Molecular, and Clinical Features. Clinical Cancer Research, 2013, 19, 5890-5900.	7.0	60
13	Transcriptional Characterization of a Prospective Series of Primary Plasma Cell Leukemia Revealed Signatures Associated with Tumor Progression and Poorer Outcome. Clinical Cancer Research, 2013, 19, 3247-3258.	7.0	50
14	Biological and Clinical Relevance of miRNA Expression Signatures in Primary Plasma Cell Leukemia. Clinical Cancer Research, 2013, 19, 3130-3142.	7.0	86
15	Prognostic Significance of Telomere Length in Chronic Lymphocytic Leukemia Patients in Early Stage Disease,. Blood, 2011, 118, 3890-3890.	1.4	7
16	Integrative Genomics Analyses Reveal Molecularly Distinct Subgroups of B-Cell Chronic Lymphocytic Leukemia Patients with 13q14 Deletion. Clinical Cancer Research, 2010, 16, 5641-5653.	7.0	52
17	Integrative highâ€resolution microarray analysis of human myeloma cell lines reveals deregulated miRNA expression associated with allelic imbalances and gene expression profiles. Genes Chromosomes and Cancer, 2009, 48, 521-531.	2.8	60
18	A SNP microarray and FISHâ€based procedure to detect allelic imbalances in multiple myeloma: An integrated genomics approach reveals a wide gene dosage effect. Genes Chromosomes and Cancer, 2009, 48, 603-614.	2.8	134

#	Article	IF	Citations
19	Identification of microRNA expression patterns and definition of a microRNA/mRNA regulatory network in distinct molecular groups of multiple myeloma. Blood, 2009, 114, e20-e26.	1.4	224
20	An integrative genomic approach reveals coordinated expression of intronic miR-335, miR-342, and miR-561 with deregulated host genes in multiple myeloma. BMC Medical Genomics, 2008, 1, 37.	1.5	104
21	Molecular and transcriptional characterization of 17p loss in Bâ€cell chronic lymphocytic leukemia. Genes Chromosomes and Cancer, 2008, 47, 781-793.	2.8	59
22	Repetitive DNA Hypomethylation in Multiple Myeloma. Blood, 2008, 112, 2703-2703.	1.4	16
23	Molecular and transcriptional characterization of the novel 17p11.2â€p12 amplicon in multiple myeloma. Genes Chromosomes and Cancer, 2007, 46, 1109-1118.	2.8	20