Angelo Lonoce

List of Publications by Year
in descending order

21 papers	936 citations	840776 11 h-index	752698 20 g-index
22 all docs	22 docs citations	22 times ranked	1682 citing authors

#	Article	IF	Citations
1	1q23.1 homozygous deletion and downregulation of Fc receptor-like family genes confer poor prognosis in chronic lymphocytic leukemia. Clinical and Experimental Medicine, 2019, 19, 261-267.	3.6	4
2	MYC-containing amplicons in acute myeloid leukemia: genomic structures, evolution, and transcriptional consequences. Leukemia, 2018, 32, 2152-2166.	7.2	70
3	RALE051: a novel established cell line of sporadic Burkitt lymphoma. Leukemia and Lymphoma, 2018, 59, 1252-1255.	1.3	0
4	Epigenetically induced ectopic expression of UNCX impairs the proliferation and differentiation of myeloid cells. Haematologica, 2017, 102, 1204-1214.	3.5	8
5	MYC-containing amplicons in acute myeloid leukemia: Genomic structures, evolution, and transcriptional consequences. Leukemia, 2017, , .	7.2	2
6	A rare but recurrent t(8;13)(q24;q14) translocation in Bâ€cell chronic lymphocytic leukaemia causing <scp>MYC</scp> upâ€regulation and concomitant loss of <scp>PVT</scp>1, miRâ€15/16 and <scp>DLEU</scp>7. British Journal of Haematology, 2016, 172, 296-299.	2.5	7
7	t(15;21) translocations leading to the concurrent downregulation of RUNX1 and its transcription factor partner genes SIN3A and TCF12 in myeloid disorders. Molecular Cancer, 2015, 14, 211.	19.2	12
8	A New Entity of Acute Myeloid Leukemia Driven By Epigenetic and Somatic Dis-Regulation of Uncx, a Novel Homeobox Transcription Factor Gene. Blood, 2015, 126, 1356-1356.	1.4	0
9	Genomic organization and evolution of double minutes/homogeneously staining regions withMYCamplification in human cancer. Nucleic Acids Research, 2014, 42, 9131-9145.	14.5	91
10	Two alternatively spliced 5â€²BCR/3â€²JAK2 fusion transcripts in a myeloproliferative neoplasm with a three-way t(9;18;22)(p23;p11.3;q11.2) translocation. Cancer Genetics, 2011, 204, 512-515.	0.4	11
11	Gene amplification as double minutes or homogeneously staining regions in solid tumors: Origin and structure. Genome Research, 2010, 20, 1198-1206.	5.5	194
12	Characterization of a hotspot region on chromosome 12 for amplification in ring chromosomes in atypical lipomatous tumors. Genes Chromosomes and Cancer, 2009, 48, 993-1001.	2.8	10
13	Identification and molecular characterization of recurrent genomic deletions on 7p12 in the IKZF1 gene in a large cohort of BCR-ABL1â€“positive acute lymphoblastic leukemia patients: on behalf of Gruppo Italiano Malattie Ematologiche dell'Adulto Acute Leukemia Working Party (GIMEMA AL WP). Blood, 2009, 114, 2159-2167.	1.4	201
14	Extramedullary molecular evidence of the 5â€²KIAA1509/3â€²PDGFRB fusion gene in chronic eosinophilic leukemia. Leukemia Research, 2008, 32, 347-351.	0.8	5
15	Bone marrow ectopic expression of a non-coding RNA in childhood T-cell acute lymphoblastic leukemia with a novel t(2;11)(q11.2;p15.1) translocation. Molecular Cancer, 2008, 7, 80.	19.2	3
16	MYC-containing double minutes in hematologic malignancies: evidence in favor of the episome model and exclusion of MYC as the target gene. Human Molecular Genetics, 2006, 15, 933-942.	2.9	116
17	Molecular cytogenetic characterization of a complex rearrangement involving chromosomes 9 and 22 in a case of Ph-negative chronic myeloid leukemia. Cancer Genetics and Cytogenetics, 2002, 136, 141-145.	1.0	12
18	Structural Organization of Multiple Alphoid Subsets Coexisting on Human Chromosomes 1, 4, 5, 7, 9, 15, 18, and 19. Genomics, 1996, 38, 325-330.	2.9	45

#	Article	IF	Citations
19	Comparative mapping of human alphoid sequences in great apes using fluorescence in situ hybridization. Genomics, 1995, 25, 477-484.	2.9	110
20	Linkage studies in Italian families with familial adenomatous polyposis. Human Genetics, 1993, 90, 545-550.	3.8	4
21	Familial adenomatous polyposis: Identification of a new frameshift mutation of the APC gene in an Italian family. Biochemical and Biophysical Research Communications, 1992, 184, 1357-1363.	2.1	24