

# Antonino Neri

## List of Publications by Year in descending order

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370  
papers

18,137  
citations

11608

70  
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19690

117  
g-index

372  
all docs

372  
docs citations

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times ranked

17294  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene Expression Profiling of B Cell Chronic Lymphocytic Leukemia Reveals a Homogeneous Phenotype Related to Memory B Cells. <i>Journal of Experimental Medicine</i> , 2001, 194, 1625-1638.	4.2	823
2	p53 mutations in human lymphoid malignancies: association with Burkitt lymphoma and chronic lymphocytic leukemia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1991, 88, 5413-5417.	3.3	817
3	International Myeloma Working Group molecular classification of multiple myeloma: spotlight review. <i>Leukemia</i> , 2009, 23, 2210-2221.	3.3	775
4	BCL-6 mutations in normal germinal center B cells: Evidence of somatic hypermutation acting outside Ig loci. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 11816-11821.	3.3	495
5	B cell lymphoma-associated chromosomal translocation involves candidate oncogene <i>lyt-10</i> , homologous to NF- $\kappa$ B p50. <i>Cell</i> , 1991, 67, 1075-1087.	13.5	430
6	Rescue of Hippo coactivator YAP1 triggers DNA damage-induced apoptosis in hematological cancers. <i>Nature Medicine</i> , 2014, 20, 599-606.	15.2	250
7	CEP-18770: A novel, orally active proteasome inhibitor with a tumor-selective pharmacologic profile competitive with bortezomib. <i>Blood</i> , 2008, 111, 2765-2775.	0.6	239
8	Different regions of the immunoglobulin heavy-chain locus are involved in chromosomal translocations in distinct pathogenetic forms of Burkitt lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1988, 85, 2748-2752.	3.3	228
9	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.	0.6	224
10	Circulating tumor DNA reveals genetics, clonal evolution, and residual disease in classical Hodgkin lymphoma. <i>Blood</i> , 2018, 131, 2413-2425.	0.6	223
11	Aberrant global methylation patterns affect the molecular pathogenesis and prognosis of multiple myeloma. <i>Blood</i> , 2011, 117, 553-562.	0.6	217
12	Synthetic miR-34a Mimics as a Novel Therapeutic Agent for Multiple Myeloma: <i>In Vitro</i> and <i>In Vivo</i> Evidence. <i>Clinical Cancer Research</i> , 2012, 18, 6260-6270.	3.2	213
13	A Novel Chromosomal Translocation t(4; 14)(p16.3; q32) in Multiple Myeloma Involves the Fibroblast Growth-Factor Receptor 3 Gene. <i>Blood</i> , 1997, 90, 4062-4070.	0.6	201
14	Molecular prediction of durable remission after first-line fludarabine-cyclophosphamide-rituximab in chronic lymphocytic leukemia. <i>Blood</i> , 2015, 126, 1921-1924.	0.6	197
15	Targeting miR-21 Inhibits <i>In Vitro</i> and <i>In Vivo</i> Multiple Myeloma Cell Growth. <i>Clinical Cancer Research</i> , 2013, 19, 2096-2106.	3.2	195
16	Analysis of RAS oncogene mutations in human lymphoid malignancies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1988, 85, 9268-9272.	3.3	191
17	Drugging the lncRNA MALAT1 via LNA gapmer ASO inhibits gene expression of proteasome subunits and triggers anti-multiple myeloma activity. <i>Leukemia</i> , 2018, 32, 1948-1957.	3.3	179
18	IRTA1 and IRTA2, Novel Immunoglobulin Superfamily Receptors Expressed in B Cells and Involved in Chromosome 1q21 Abnormalities in B Cell Malignancy. <i>Immunity</i> , 2001, 14, 277-289.	6.6	176

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19	Ras oncogene mutation in multiple myeloma.. Journal of Experimental Medicine, 1989, 170, 1715-1725.	4.2	166
20	Gene Expression Profiling Uncovers Molecular Classifiers for the Recognition of Anaplastic Large-Cell Lymphoma Within Peripheral T-Cell Neoplasms. Journal of Clinical Oncology, 2010, 28, 1583-1590.	0.8	152
21	Oct-4 Expression in Adult Human Differentiated Cells Challenges Its Role as a Pure Stem Cell Marker. Stem Cells, 2007, 25, 1675-1680.	1.4	151
22	Functional validation of the anaplastic lymphoma kinase signature identifies CEBPB and Bcl2A1 as critical target genes. Journal of Clinical Investigation, 2006, 116, 3171-3182.	3.9	139
23	DNA-demethylating and anti-tumor activity of synthetic miR-29b mimics in multiple myeloma. Oncotarget, 2012, 3, 1246-1258.	0.8	138
24	miR-29b sensitizes multiple myeloma cells to bortezomib-induced apoptosis through the activation of a feedback loop with the transcription factor Sp1. Cell Death and Disease, 2012, 3, e436-e436.	2.7	137
25	A SNP microarray and FISH-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.	1.5	134
26	The histone methyltransferase MMSET/WHSC1 activates TWIST1 to promote an epithelial-to-mesenchymal transition and invasive properties of prostate cancer. Oncogene, 2013, 32, 2882-2890.	2.6	130
27	Thalidomide Downregulates Angiogenic Genes in Bone Marrow Endothelial Cells of Patients With Active Multiple Myeloma. Journal of Clinical Oncology, 2005, 23, 5334-5346.	0.8	125
28	In Vitro and in Vivo Anti-tumor Activity of miR-221/222 Inhibitors in Multiple Myeloma. Oncotarget, 2013, 4, 242-255.	0.8	125
29	Molecular Classification of Multiple Myeloma: A Distinct Transcriptional Profile Characterizes Patients Expressing CCND1 and Negative for 14q32 Translocations. Journal of Clinical Oncology, 2005, 23, 7296-7306.	0.8	123
30	Combining Anti-Mir-155 with Chemotherapy for the Treatment of Lung Cancers. Clinical Cancer Research, 2017, 23, 2891-2904.	3.2	122
31	Canonical and noncanonical Hedgehog pathway in the pathogenesis of multiple myeloma. Blood, 2012, 120, 5002-5013.	0.6	121
32	Gene expression profiling of plasma cell dyscrasias reveals molecular patterns associated with distinct IGH translocations in multiple myeloma. Oncogene, 2005, 24, 2461-2473.	2.6	118
33	The new tumor-suppressor gene inhibitor of growth family member 4 (ING4) regulates the production of proangiogenic molecules by myeloma cells and suppresses hypoxia-inducible factor-1 (HIF-1) activity: involvement in myeloma-induced angiogenesis. Blood, 2007, 110, 4464-4475.	0.6	117
34	Anaplastic lymphoma kinase in human cancer. Journal of Molecular Endocrinology, 2011, 47, R11-R23.	1.1	116
35	Increased osteocyte death in multiple myeloma patients: role in myeloma-induced osteoclast formation. Leukemia, 2012, 26, 1391-1401.	3.3	116
36	ILF2 Is a Regulator of RNA Splicing and DNA Damage Response in 1q21-Amplified Multiple Myeloma. Cancer Cell, 2017, 32, 88-100.e6.	7.7	114

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37	Cyclin D1 overexpression is a favorable prognostic variable for newly diagnosed multiple myeloma patients treated with high-dose chemotherapy and single or double autologous transplantation. <i>Blood</i> , 2003, 102, 1588-1594.	0.6	113
38	Low bone marrow oxygen tension and hypoxia-inducible factor-1 $\alpha$ overexpression characterize patients with multiple myeloma: role on the transcriptional and proangiogenic profiles of CD138+ cells. <i>Leukemia</i> , 2010, 24, 1967-1970.	3.3	107
39	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.	0.7	104
40	Hypoxia-inducible factor (HIF)-1 $\alpha$ suppression in myeloma cells blocks tumoral growth in vivo inhibiting angiogenesis and bone destruction. <i>Leukemia</i> , 2013, 27, 1697-1706.	3.3	104
41	Selective targeting of IRF4 by synthetic microRNA-125b-5p mimics induces anti-multiple myeloma activity in vitro and in vivo. <i>Leukemia</i> , 2015, 29, 2173-2183.	3.3	104
42	The histone deacetylase inhibitor ITF2357 has anti-leukemic activity in vitro and in vivo and inhibits IL-6 and VEGF production by stromal cells. <i>Leukemia</i> , 2007, 21, 1892-1900.	3.3	102
43	Identification of a 3-gene model as a powerful diagnostic tool for the recognition of ALK-negative anaplastic large-cell lymphoma. <i>Blood</i> , 2012, 120, 1274-1281.	0.6	101
44	Differential repetitive DNA methylation in multiple myeloma molecular subgroups. <i>Carcinogenesis</i> , 2009, 30, 1330-1335.	1.3	99
45	Deregulated FGFR3 mutants in multiple myeloma cell lines with t(4;14): comparative analysis of Y373C, K650E and the novel G384D mutations. <i>Oncogene</i> , 2001, 20, 3553-3562.	2.6	98
46	Identification of a new subclass of ALK-negative ALCL expressing aberrant levels of ERBB4 transcripts. <i>Blood</i> , 2016, 127, 221-232.	0.6	97
47	miR-29b induces SOCS-1 expression by promoter demethylation and negatively regulates migration of multiple myeloma and endothelial cells. <i>Cell Cycle</i> , 2013, 12, 3650-3662.	1.3	96
48	Therapeutic Targeting of miR-29b/HDAC4 Epigenetic Loop in Multiple Myeloma. <i>Molecular Cancer Therapeutics</i> , 2016, 15, 1364-1375.	1.9	94
49	Autoimmune cytopenias in chronic lymphocytic leukemia. <i>American Journal of Hematology</i> , 2014, 89, 1055-1062.	2.0	93
50	Kaposi's Sarcoma-Associated Herpesvirus Infection and Multiple Myeloma. <i>Science</i> , 1997, 278, 1969-1973.	6.0	92
51	Targeting of multiple myeloma-related angiogenesis by miR-199a-5p mimics: <i>in vitro</i> and <i>in vivo</i> anti-tumor activity. <i>Oncotarget</i> , 2014, 5, 3039-3054.	0.8	92
52	Gene Expression Profiling of Bone Marrow Endothelial Cells in Patients with Multiple Myeloma. <i>Clinical Cancer Research</i> , 2009, 15, 5369-5378.	3.2	91
53	Microvessel density, a surrogate marker of angiogenesis, is significantly related to survival in multiple myeloma patients. <i>British Journal of Haematology</i> , 2002, 118, 817-820.	1.2	87
54	Biological and prognostic impact of APOBEC-induced mutations in the spectrum of plasma cell dyscrasias and multiple myeloma cell lines. <i>Leukemia</i> , 2018, 32, 1043-1047.	3.3	87

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55	Biological and Clinical Relevance of miRNA Expression Signatures in Primary Plasma Cell Leukemia. <i>Clinical Cancer Research</i> , 2013, 19, 3130-3142.	3.2	86
56	A p53-Dependent Tumor Suppressor Network Is Induced by Selective miR-125a Inhibition in Multiple Myeloma Cells. <i>Journal of Cellular Physiology</i> , 2014, 229, 2106-2116.	2.0	86
57	International prognostic score for asymptomatic early-stage chronic lymphocytic leukemia. <i>Blood</i> , 2020, 135, 1859-1869.	0.6	86
58	Promises and Challenges of MicroRNA-based Treatment of Multiple Myeloma. <i>Current Cancer Drug Targets</i> , 2012, 12, 838-846.	0.8	84
59	The Kr��ppel-like factor 2 transcription factor gene is recurrently mutated in splenic marginal zone lymphoma. <i>Leukemia</i> , 2015, 29, 503-507.	3.3	84
60	Rearranged <i>NFKB-2</i> Genes in Lymphoid Neoplasms Code for Constitutively Active Nuclear Transactivators. <i>Molecular and Cellular Biology</i> , 1995, 15, 5180-5187.	1.1	83
61	Structural and functional characterization of the promoter regions of the <i>NFKB2</i> gene. <i>Nucleic Acids Research</i> , 1995, 23, 2328-2336.	6.5	82
62	Molecular Analysis of 11q13 Breakpoints in Multiple Myeloma. <i>Blood</i> , 1999, 93, 1330-1337.	0.6	80
63	Long non-coding RNA NEAT1 targeting impairs the DNA repair machinery and triggers anti-tumor activity in multiple myeloma. <i>Leukemia</i> , 2020, 34, 234-244.	3.3	80
64	Distinct lncRNA transcriptional fingerprints characterize progressive stages of multiple myeloma. <i>Oncotarget</i> , 2016, 7, 14814-14830.	0.8	79
65	Lenalidomide and low-dose dexamethasone for newly diagnosed primary plasma cell leukemia. <i>Leukemia</i> , 2014, 28, 222-225.	3.3	77
66	Acquired CYP19A1 amplification is an early specific mechanism of aromatase inhibitor resistance in ER± metastatic breast cancer. <i>Nature Genetics</i> , 2017, 49, 444-450.	9.4	77
67	miR-451a is underexpressed and targets AKT/mTOR pathway in papillary thyroid carcinoma. <i>Oncotarget</i> , 2016, 7, 12731-12747.	0.8	77
68	Clinical Relevance of Expression of the CIP/KIP Cell-Cycle Inhibitors p21 and p27 in Laryngeal Cancer. <i>Journal of Clinical Oncology</i> , 1999, 17, 3150-3159.	0.8	75
69	Small nucleolar RNAs as new biomarkers in chronic lymphocytic leukemia. <i>BMC Medical Genomics</i> , 2013, 6, 27.	0.7	73
70	Genotypic Monoclonality in Immunophenotypically Polyclonal Orbital Lymphoid Tumors. <i>Ophthalmology</i> , 1987, 94, 980-994.	2.5	72
71	Immunohistochemical Analysis of Cyclin D1 Shows Deregulated Expression in Multiple Myeloma with the t(11;14). <i>American Journal of Pathology</i> , 2000, 156, 1505-1513.	1.9	72
72	Treatment and prognosis in a series of primary extranodal lymphomas of the ocular adnexa. <i>Annals of Oncology</i> , 1998, 9, 779-781.	0.6	70

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73	Clinical relevance of cyclin D1 protein overexpression in laryngeal squamous cell carcinoma.. Journal of Clinical Oncology, 1998, 16, 3069-3077.	0.8	70
74	The expression pattern of small nucleolar and small Cajal body-specific RNAs characterizes distinct molecular subtypes of multiple myeloma. Blood Cancer Journal, 2012, 2, e96-e96.	2.8	70
75	Non-coding RNA: a novel opportunity for the personalized treatment of multiple myeloma. Expert Opinion on Biological Therapy, 2013, 13, S125-S137.	1.4	70
76	Upregulation of translational machinery and distinct genetic subgroups characterise hyperdiploidy in multiple myeloma. British Journal of Haematology, 2007, 136, 565-573.	1.2	66
77	The Reconstruction of Transcriptional Networks Reveals Critical Genes with Implications for Clinical Outcome of Multiple Myeloma. Clinical Cancer Research, 2011, 17, 7402-7412.	3.2	65
78	Immunomodulatory drugs lenalidomide and pomalidomide inhibit multiple myeloma-induced osteoclast formation and the RANKL/OPG ratio in the myeloma microenvironment targeting the expression of adhesion molecules. Experimental Hematology, 2013, 41, 387-397.e1.	0.2	65
79	Molecular spectrum of <i>BRAF</i> , <i>NRAS</i> and <i>KRAS</i> gene mutations in plasma cell dyscrasias: implication for MEK-ERK pathway activation. Oncotarget, 2015, 6, 24205-24217.	0.8	65
80	Frequent p53 gene involvement in splenic B-cell leukemia/lymphomas of possible marginal zone origin. Blood, 1994, 84, 270-278.	0.6	63
81	Molecular characterization of human multiple myeloma cell lines by integrative genomics: Insights into the biology of the disease. Genes Chromosomes and Cancer, 2007, 46, 226-238.	1.5	62
82	Analysis of CD20-dependent cellular cytotoxicity by G-CSF-stimulated neutrophils. Leukemia, 2002, 16, 693-699.	3.3	60
83	Integrative high-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.	1.5	60
84	Genome-wide analysis of primary plasma cell leukemia identifies recurrent imbalances associated with changes in transcriptional profiles. American Journal of Hematology, 2013, 88, 16-23.	2.0	60
85	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.	3.2	60
86	Therapeutic Targeting of miR-29b/HDAC4 Epigenetic Loop in Multiple Myeloma. Molecular Cancer Therapeutics, 2016, 15, 1364-1375.	1.9	60
87	Inhibition of EZH2 triggers the tumor suppressive miR-29b network in multiple myeloma. Oncotarget, 2017, 8, 106527-106537.	0.8	60
88	Analysis of FGFR3 gene mutations in multiple myeloma patients with t(4;14). British Journal of Haematology, 2001, 114, 362-364.	1.2	59
89	Molecular and transcriptional characterization of 17p loss in B-cell chronic lymphocytic leukemia. Genes Chromosomes and Cancer, 2008, 47, 781-793.	1.5	59
90	Detection of t(4;14)(p16.3;q32) Chromosomal Translocation in Multiple Myeloma by Double-Color Fluorescent In Situ Hybridization. Blood, 1999, 94, 724-732.	0.6	58

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91	Biological and clinical relevance of quantitative global methylation of repetitive DNA sequences in chronic lymphocytic leukemia. <i>Epigenetics</i> , 2011, 6, 188-194.	1.3	58
92	The chronic lymphocytic leukemia international prognostic index predicts time to first treatment in early CLL: Independent validation in a prospective cohort of early stage patients. <i>American Journal of Hematology</i> , 2016, 91, 1090-1095.	2.0	58
93	Distinct transcriptional profiles characterize bone microenvironment mesenchymal cells rather than osteoblasts in relationship with multiple myeloma bone disease. <i>Experimental Hematology</i> , 2010, 38, 141-153.	0.2	57
94	Bendamustine in combination with Ofatumumab in relapsed or refractory chronic lymphocytic leukemia: a GIMEMA Multicenter Phase II Trial. <i>Leukemia</i> , 2014, 28, 642-648.	3.3	57
95	Whole-exome sequencing of primary plasma cell leukemia discloses heterogeneous mutational patterns. <i>Oncotarget</i> , 2015, 6, 17543-17558.	0.8	55
96	Improved risk stratification in myeloma using a microRNA-based classifier. <i>British Journal of Haematology</i> , 2013, 162, 348-359.	1.2	53
97	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.	3.2	52
98	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.	1.2	52
99	microRNAome Expression in Chronic Lymphocytic Leukemia: Comparison with Normal B-cell Subsets and Correlations with Prognostic and Clinical Parameters. <i>Clinical Cancer Research</i> , 2014, 20, 4141-4153.	3.2	52
100	Validation of the CLL-IPI and comparison with the MDACC prognostic index in newly diagnosed patients. <i>Blood</i> , 2016, 128, 2093-2095.	0.6	52
101	Therapeutic vulnerability of multiple myeloma to MIR17PTi, a first-in-class inhibitor of pri-miR-17-92. <i>Blood</i> , 2018, 132, 1050-1063.	0.6	52
102	Molecular and immunohistochemical analysis of the bcl-1/cyclin D1 gene in laryngeal squamous cell carcinomas. , 1997, 79, 1114-1121.		50
103	Definition of progression risk based on combinations of cellular and molecular markers in patients with Binet stage A chronic lymphocytic leukaemia. <i>British Journal of Haematology</i> , 2009, 146, 44-53.	1.2	50
104	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.	3.2	50
105	miR-23b/SP1/c-myc forms a feed-forward loop supporting multiple myeloma cell growth. <i>Blood Cancer Journal</i> , 2016, 6, e380-e380.	2.8	50
106	Heterogeneity of TP53 Mutations and P53 Protein Residual Function in Cancer: Does It Matter?. <i>Frontiers in Oncology</i> , 2020, 10, 593383.	1.3	50
107	Long non-coding RNAs in normal and malignant hematopoiesis. <i>Oncotarget</i> , 2016, 7, 50666-50681.	0.8	50
108	Characterization of oncogene dysregulation in multiple myeloma by combined FISH and DNA microarray analyses. <i>Genes Chromosomes and Cancer</i> , 2005, 42, 117-127.	1.5	49

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109	Consensus statement from European experts on the diagnosis, management, and treatment of multiple myeloma: from standard therapy to novel approaches. <i>Leukemia and Lymphoma</i> , 2010, 51, 1424-1443.	0.6	49
110	Myeloma cells inhibit non-canonical wnt co-receptor ror2 expression in human bone marrow osteoprogenitor cells: effect of wnt5a/ror2 pathway activation on the osteogenic differentiation impairment induced by myeloma cells. <i>Leukemia</i> , 2013, 27, 451-463.	3.3	48
111	Depletion of SIRT6 enzymatic activity increases acute myeloid leukemia cells' vulnerability to DNA-damaging agents. <i>Haematologica</i> , 2018, 103, 80-90.	1.7	48
112	lncRNA profiling in early-stage chronic lymphocytic leukemia identifies transcriptional fingerprints with relevance in clinical outcome. <i>Blood Cancer Journal</i> , 2016, 6, e468-e468.	2.8	47
113	Notch signaling deregulation in multiple myeloma: A rational molecular target. <i>Oncotarget</i> , 2015, 6, 26826-26840.	0.8	47
114	Clinical relevance of p53 and bcl-2 protein over-expression in laryngeal squamous-cell carcinoma. , 1998, 79, 263-268.		46
115	CD26 expression in mature B-cell neoplasia: its possible role as a new prognostic marker in B-CLL. <i>Hematological Oncology</i> , 2009, 27, 140-147.	0.8	46
116	Analysis of p53 and ras Gene Mutations in Endometriosis. <i>Gynecologic and Obstetric Investigation</i> , 1994, 38, 70-71.	0.7	45
117	Transcriptional features of multiple myeloma patients with chromosome 1q gain. <i>Leukemia</i> , 2007, 21, 1113-1116.	3.3	45
118	Identification of primary MAFB target genes in multiple myeloma. <i>Experimental Hematology</i> , 2009, 37, 78-86.	0.2	45
119	The cumulative amount of serum-free light chain is a strong prognosticator in chronic lymphocytic leukemia. <i>Blood</i> , 2011, 118, 6353-6361.	0.6	45
120	Notch signaling drives multiple myeloma induced osteoclastogenesis. <i>Oncotarget</i> , 2014, 5, 10393-10406.	0.8	45
121	MicroRNAs in the Pathobiology of Multiple Myeloma. <i>Current Cancer Drug Targets</i> , 2012, 12, 823-837.	0.8	44
122	Constitutive expression of lymphoma-associated NFKB-2/Lyt-10 proteins is tumorigenic in murine fibroblasts. <i>Oncogene</i> , 1997, 14, 1805-1810.	2.6	42
123	A novel patient-derived tumorgraft model with TRAF1-ALK anaplastic large-cell lymphoma translocation. <i>Leukemia</i> , 2015, 29, 1390-1401.	3.3	42
124	Minimal residual disease in acute lymphoblastic leukemia detected by immune selection and gene rearrangement analysis.. <i>Journal of Clinical Oncology</i> , 1989, 7, 338-343.	0.8	41
125	Notch pathway promotes ovarian cancer growth and migration via CXCR4/SDF1 chemokine system. <i>International Journal of Biochemistry and Cell Biology</i> , 2015, 66, 134-140.	1.2	41
126	Disentangling the microRNA regulatory milieu in multiple myeloma: integrative genomics analysis outlines mixed miRNA-TF circuits and pathway-derived networks modulated in t(4;14) patients. <i>Oncotarget</i> , 2016, 7, 2367-2378.	0.8	41



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127	Integrated analysis of microRNAs, transcription factors and target genes expression discloses a specific molecular architecture of hyperdiploid multiple myeloma. <i>Oncotarget</i> , 2015, 6, 19132-19147.	0.8	41
128	p63 in Laryngeal Squamous Cell Carcinoma: Evidence for a Role of TA-p63 Down-Regulation in Tumorigenesis and Lack of Prognostic Implications of p63 Immunoreactivity. <i>Laboratory Investigation</i> , 2002, 82, 1327-1334.	1.7	40
129	IDH2 inhibition enhances proteasome inhibitor responsiveness in hematological malignancies. <i>Blood</i> , 2019, 133, 156-167.	0.6	40
130	A compendium of <i>DIS3</i> mutations and associated transcriptional signatures in plasma cell dyscrasias. <i>Oncotarget</i> , 2015, 6, 26129-26141.	0.8	40
131	Molecular spectrum of <i>TP53</i> mutations in plasma cell dyscrasias by next generation sequencing: an Italian cohort study and overview of the literature. <i>Oncotarget</i> , 2016, 7, 21353-21361.	0.8	40
132	The transactivating isoforms of p63 are overexpressed in high-grade follicular lymphomas independent of the occurrence of p63 gene amplification. <i>Journal of Pathology</i> , 2005, 206, 337-345.	2.1	39
133	HOXB7 expression by myeloma cells regulates their pro-angiogenic properties in multiple myeloma patients. <i>Leukemia</i> , 2011, 25, 527-537.	3.3	39
134	ALK signaling and target therapy in anaplastic large cell lymphoma. <i>Frontiers in Oncology</i> , 2012, 2, 41.	1.3	39
135	miR-22 suppresses DNA ligase III addiction in multiple myeloma. <i>Leukemia</i> , 2019, 33, 487-498.	3.3	39
136	Variability of polymerase chain reaction detection of the bcl-2-IgH translocation in an international multicentre study. <i>Annals of Oncology</i> , 1999, 10, 1349-1354.	0.6	38
137	Relevance of telomere/telomerase system impairment in early stage chronic lymphocytic leukemia. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 612-621.	1.5	38
138	The Involvement of the Candidate Proto-Oncogene NFKB2/lyt-10 in Lymphoid Malignancies. <i>Leukemia and Lymphoma</i> , 1996, 23, 43-48.	0.6	37
139	Identification of a tumor-associated mutant form of the NF- $\kappa$ B RelA gene with reduced DNA-binding and transactivating activities. <i>Oncogene</i> , 1997, 14, 791-799.	2.6	37
140	The oral protein-kinase C $\beta$ inhibitor enzastaurin (LY317615) suppresses signalling through the AKT pathway, inhibits proliferation and induces apoptosis in multiple myeloma cell lines. <i>Leukemia and Lymphoma</i> , 2008, 49, 1374-1383.	0.6	37
141	Predictive value of $\beta$ 2-microglobulin ( $\beta$ 2-m) levels in chronic lymphocytic leukemia since Binet A stages. <i>Haematologica</i> , 2009, 94, 887-888.	1.7	37
142	Impact of Host Genes and Strand Selection on miRNA and miRNA* Expression. <i>PLoS ONE</i> , 2011, 6, e23854.	1.1	37
143	Analysis of p53 gene mutations in acute myeloid leukemia. <i>American Journal of Hematology</i> , 1994, 46, 304-309.	2.0	36
144	FGFR3 Gene Mutations Associated With Human Skeletal Disorders Occur Rarely in Multiple Myeloma. <i>Blood</i> , 1998, 92, 2987-2989.	0.6	36

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145	Low-dose subcutaneous alemtuzumab in refractory chronic lymphocytic leukaemia (CLL): results of a prospective, single-arm multicentre study. <i>Leukemia</i> , 2009, 23, 2027-2033.	3.3	36
146	Pleiotropic anti-myeloma activity of ITF2357: inhibition of interleukin-6 receptor signaling and repression of miR-19a and miR-19b. <i>Haematologica</i> , 2010, 95, 260-269.	1.7	36
147	Relevance of Stereotyped B-Cell Receptors in the Context of the Molecular, Cytogenetic and Clinical Features of Chronic Lymphocytic Leukemia. <i>PLoS ONE</i> , 2011, 6, e24313.	1.1	36
148	The HDAC inhibitor Givinostat modulates the hematopoietic transcription factors NFE2 and C-MYB in JAK2V617F myeloproliferative neoplasm cells. <i>Experimental Hematology</i> , 2012, 40, 634-645.e10.	0.2	36
149	Bâ€cell receptor configuration and adverse cytogenetics are associated with autoimmune hemolytic anemia in chronic lymphocytic leukemia. <i>American Journal of Hematology</i> , 2013, 88, 32-36.	2.0	36
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