

Antonella Zagaria

List of Publications by Year in descending order

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Version: 2024-02-01

80
papers

1,301
citations

331670

21
h-index

434195

31
g-index

80
all docs

80
docs citations

80
times ranked

1767
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>RUNX1</i> gene alterations characterized by allelic preference in adult acute myeloid leukemia. <i>Leukemia and Lymphoma</i> , 2023, 64, 717-721.	1.3	4
2	FLT3 mutational analysis in acute myeloid leukemia: Advantages and pitfalls with different approaches. <i>Blood Reviews</i> , 2022, 54, 100928.	5.7	4
3	Second Cancer Onset in Myeloproliferative Neoplasms: What, When, Why?. <i>International Journal of Molecular Sciences</i> , 2022, 23, 3177.	4.1	5
4	Single-Cell Sequencing: Ariadne's Thread in the Maze of Acute Myeloid Leukemia. <i>Diagnostics</i> , 2022, 12, 996.	2.6	2
5	Clonal Hematopoiesis at the Crossroads of Inflammatory Bowel Diseases and Hematological Malignancies: A Biological Link?. <i>Frontiers in Oncology</i> , 2022, 12, 873896.	2.8	12
6	Erythrocytosis with JAK2 GGCC_46/1 haplotype and without JAK2 V617F mutation is associated with CALR rs1049481_G allele. <i>Leukemia</i> , 2021, 35, 619-622.	7.2	5
7	Nanopore sequencing sheds a light on the <i>FLT3</i> gene mutations complexity in acute promyelocytic leukemia. <i>Leukemia and Lymphoma</i> , 2021, 62, 1219-1225.	1.3	5
8	Dysregulation of miRNA in Leukemia: Exploiting miRNA Expression Profiles as Biomarkers. <i>International Journal of Molecular Sciences</i> , 2021, 22, 7156.	4.1	21
9	Can the New and Old Drugs Exert an Immunomodulatory Effect in Acute Myeloid Leukemia?. <i>Cancers</i> , 2021, 13, 4121.	3.7	2
10	Nanopore sequencing approach for immunoglobulin gene analysis in chronic lymphocytic leukemia. <i>Scientific Reports</i> , 2021, 11, 17668.	3.3	6
11	IRF4 expression is low in Philadelphia negative myeloproliferative neoplasms and is associated with a worse prognosis. <i>Experimental Hematology and Oncology</i> , 2021, 10, 58.	5.0	3
12	Inside the biology of early T-cell precursor acute lymphoblastic leukemia: the perfect trick. <i>Biomarker Research</i> , 2021, 9, 89.	6.8	5
13	TP53 in Myelodysplastic Syndromes: Recent Biological and Clinical Findings. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3432.	4.1	25
14	Digital PCR: A Reliable Tool for Analyzing and Monitoring Hematologic Malignancies. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3141.	4.1	35
15	HMGA Proteins in Hematological Malignancies. <i>Cancers</i> , 2020, 12, 1456.	3.7	8
16	Nanopore Sequencing in Blood Diseases: A Wide Range of Opportunities. <i>Frontiers in Genetics</i> , 2020, 11, 76.	2.3	29
17	Molecular Complexity of Diffuse Large B-Cell Lymphoma: Can It Be a Roadmap for Precision Medicine?. <i>Cancers</i> , 2020, 12, 185.	3.7	22
18	Skin lesions in chronic myeloid leukemia patients during dasatinib treatment. <i>Cancer Management and Research</i> , 2019, Volume 11, 7991-7996.	1.9	7

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19	RARG Gene Dysregulation in Acute Myeloid Leukemia. <i>Frontiers in Molecular Biosciences</i> , 2019, 6, 114.	3.5	18
20	A complex and cryptic intrachromosomal rearrangement generating the FIP1L1_PDGFR A in adult acute myeloid leukemia. <i>Cancer Genetics</i> , 2019, 239, 8-12.	0.4	0
21	Next-Generation Sequencing in Acute Lymphoblastic Leukemia. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2929.	4.1	60
22	The Pleiotropic Role of Retinoic Acid/Retinoic Acid Receptors Signaling: From Vitamin A Metabolism to Gene Rearrangements in Acute Promyelocytic Leukemia. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2921.	4.1	16
23	Nanopore Targeted Sequencing for Rapid Gene Mutations Detection in Acute Myeloid Leukemia. <i>Genes</i> , 2019, 10, 1026.	2.4	26
24	Droplet Digital PCR Is a Robust Tool for Monitoring Minimal Residual Disease in Adult Philadelphia-Positive Acute Lymphoblastic Leukemia. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 474-482.	2.8	41
25	RARA and RARG gene downregulation associated with EZH2 mutation in acute promyelocytic-like morphology leukemia. <i>Human Pathology</i> , 2018, 80, 82-86.	2.0	19
26	A novel t(3;9)(q21.2; p24.3) associated with SMARCA2 and ZNF148 genes rearrangement in myelodysplastic syndrome. <i>Leukemia and Lymphoma</i> , 2018, 59, 996-999.	1.3	4
27	Droplet digital PCR for the quantification of Alu methylation status in hematological malignancies. <i>Diagnostic Pathology</i> , 2018, 13, 98.	2.0	7
28	Genomic BCR-ABL1 breakpoint characterization by a multi-strategy approach for personalized monitoring of residual disease in chronic myeloid leukemia patients. <i>Oncotarget</i> , 2018, 9, 10978-10986.	1.8	29
29	The JAK2 GGCC (46/1) Haplotype in Myeloproliferative Neoplasms: Causal or Random?. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1152.	4.1	16
30	Design and MinION testing of a nanopore targeted gene sequencing panel for chronic lymphocytic leukemia. <i>Scientific Reports</i> , 2018, 8, 11798.	3.3	34
31	Droplet Digital PCR Is a Reliable Tool for Monitoring Minimal Residual Disease in Acute Promyelocytic Leukemia. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 437-444.	2.8	34
32	CPX-351 in acute myeloid leukemia: can a new formulation maximize the efficacy of old compounds?. <i>Expert Review of Hematology</i> , 2017, 10, 853-862.	2.2	21
33	Mutational analysis in BCR - ABL1 positive leukemia by deep sequencing based on nanopore MinION technology. <i>Experimental and Molecular Pathology</i> , 2017, 103, 33-37.	2.1	36
34	Monosomal karyotype in myeloid neoplasias: a literature review. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 2163-2171.	2.0	13
35	<i>SETBP1</i> dysregulation in congenital disorders and myeloid neoplasms. <i>Oncotarget</i> , 2017, 8, 51920-51935.	1.8	26
36	Droplet digital PCR analysis of <i>NOTCH1</i> gene mutations in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2016, 7, 86469-86479.	1.8	23

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37	Droplet digital PCR assay for quantifying of CALR mutant allelic burden in myeloproliferative neoplasms. <i>Annals of Hematology</i> , 2016, 95, 1559-1560.	1.8	16
38	TP53 gene mutation analysis in chronic lymphocytic leukemia by nanopore MinION sequencing. <i>Diagnostic Pathology</i> , 2016, 11, 96.	2.0	51
39	MYEOV gene overexpression in primary plasma cell leukemia with t(11;14)(q13;q32). <i>Oncology Letters</i> , 2016, 12, 1460-1464.	1.8	7
40	Absolute quantification of the pretreatment <i>PML-RARA</i> transcript defines the relapse risk in acute promyelocytic leukemia. <i>Oncotarget</i> , 2015, 6, 13269-13277.	1.8	35
41	Overexpression of the LSAMP and TUSC7 genes in acute myeloid leukemia following microdeletion/duplication of chromosome 3. <i>Cancer Genetics</i> , 2015, 208, 517-522.	0.4	8
42	<i>BCR-ABL1 e6a2</i> transcript in chronic myeloid leukemia: biological features and molecular monitoring by droplet digital PCR. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2015, 467, 357-363.	2.8	24
43	Centromeric fragment of chromosome 7 in atypical chronic myeloid leukemia with the SET binding protein 1 gene mutation. <i>Leukemia and Lymphoma</i> , 2015, 56, 826-828.	1.3	1
44	Myelodysplastic syndrome with 5q deletion following IgM monoclonal gammopathy, showing gene mutation MYD88 L265P. <i>Blood Cells, Molecules, and Diseases</i> , 2015, 54, 51-52.	1.4	5
45	Droplet Digital PCR Analysis for Diagnosis and Minimal Residual Disease Monitoring in Adult Ph+ Acute Lymphoblastic Leukemia. <i>Blood</i> , 2015, 126, 4989-4989.	1.4	1
46	5' <i>RUNX1-3' UTR</i> <i>UASP42</i> chimeric gene in acute myeloid leukemia can occur through an insertion mechanism rather than translocation and may be mediated by genomic segmental duplications. <i>Molecular Cytogenetics</i> , 2014, 7, 66.	0.9	11
47	<i>BCL6</i> corepressor gene dysregulation due to chromosomal translocation in acute myeloid leukemia: a new mechanism based on long non-coding RNA dislocation?. <i>Leukemia and Lymphoma</i> , 2014, 55, 2199-2201.	1.3	2
48	<i>ADAMTS2</i> gene dysregulation in T/myeloid mixed phenotype acute leukemia. <i>BMC Cancer</i> , 2014, 14, 963.	2.6	6
49	IgG-lymphoplasmacytic lymphoma following polycythemia vera: <i>JAK2 V617F</i> and <i>MYD88 L265P</i> mutations separated in the same house. <i>Annals of Hematology</i> , 2014, 93, 1605-1607.	1.8	8
50	Acute myeloid leukemia with t(16;16) (p13;q22) showing a new <i>CBFB-MYH11</i> fusion transcript associated with an atypical leukemic blasts morphology. <i>Human Pathology</i> , 2014, 45, 643-647.	2.0	10
51	Gene expression profiling of chronic myeloid leukemia with variant t(9;22) reveals a different signature from cases with classic translocation. <i>Molecular Cancer</i> , 2013, 12, 36.	19.2	14
52	A novel t(4;16)(q25;q23.1) associated with <i>EGF</i> and <i>ELOVL6</i> deregulation in acute myeloid leukemia. <i>Gene</i> , 2013, 529, 144-147.	2.2	8
53	<i>SETBP1</i> and <i>miR_4319</i> dysregulation in primary myelofibrosis progression to acute myeloid leukemia. <i>Journal of Hematology and Oncology</i> , 2012, 5, 48.	17.0	27
54	A new recurrent chromosomal translocation t(3;11)(q13;q14) in myelodysplastic syndromes associated with overexpression of the <i>ILDR1</i> gene. <i>Leukemia Research</i> , 2012, 36, 852-856.	0.8	6

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55	Decreased TET2 gene expression during chronic myeloid leukemia progression. <i>Leukemia Research</i> , 2011, 35, e220-e222.	0.8	30
56	Genomic segmental duplications on the basis of the t(9;22) rearrangement in chronic myeloid leukemia. <i>Oncogene</i> , 2010, 29, 2509-2516.	5.9	35
57	Non random distribution of genomic features in breakpoint regions involved in chronic myeloid leukemia cases with variant t(9;22) or additional chromosomal rearrangements. <i>Molecular Cancer</i> , 2010, 9, 120.	19.2	37
58	<i>MIRN199B</i> downregulation in chronic myeloid leukaemia is associated with deletions on der(9). <i>British Journal of Haematology</i> , 2009, 144, 271-273.	2.5	7
59	Downregulated expression of genes mapping on chromosome 9 in chronic myeloid leukemia cases bearing genomic deletions on der(9). <i>Leukemia</i> , 2009, 23, 813-816.	7.2	3
60	The double deceit generated by an insertion mechanism in chronic myeloid leukemia with t(9;9;22). <i>Annals of Hematology</i> , 2008, 87, 923-926.	1.8	1
61	Extramedullary molecular evidence of the 5'KIAA1509/3'PDGFRB fusion gene in chronic eosinophilic leukemia. <i>Leukemia Research</i> , 2008, 32, 347-351.	0.8	5
62	Genomic and molecular switching in relapsed acute promyelocytic leukemia. <i>Leukemia</i> , 2008, 22, 1469-1472.	7.2	1
63	Home-brew-FISH assay shows higher efficiency than BCR-ABL dual color, dual fusion probe in detecting microdeletions and complex rearrangements associated with t(9;22) in chronic myeloid leukemia. <i>Cancer Genetics and Cytogenetics</i> , 2007, 174, 121-126.	1.0	23
64	Molecular cytogenetic findings supporting the evidence of a biclonal origin in acute myeloid leukemia. <i>Annals of Hematology</i> , 2006, 85, 129-131.	1.8	0
65	Molecular cytogenetic characterization of deletions on der(9) in chronic myelocytic leukemia. <i>Cancer Genetics and Cytogenetics</i> , 2006, 167, 97-102.	1.0	21
66	Submicroscopic deletions in an acute myeloid leukemia case with a four-way t(8;11;16;21). <i>Leukemia Research</i> , 2005, 29, 855-858.	0.8	12
67	Molecular cytogenetic study of instability at 1q21-q32 in adult acute lymphoblastic leukemia. <i>Cancer Genetics and Cytogenetics</i> , 2005, 156, 54-58.	1.0	4
68	Pericentric chromosome 8 inversion associated with the 5'RUNX1/3'CBFA2T1 gene in acute myeloid leukemia cases. <i>Annals of Hematology</i> , 2005, 84, 245-249.	1.8	7
69	A chronic myelocytic leukemia case bearing deletions on the three chromosomes involved in a variant t(9;22;11). <i>Cancer Genetics and Cytogenetics</i> , 2004, 148, 137-140.	1.0	7
70	A fluorescence in situ hybridization study of complex t(9;22) in two chronic myelocytic leukemia cases with a masked Philadelphia chromosome. <i>Cancer Genetics and Cytogenetics</i> , 2004, 150, 81-85.	1.0	25
71	A novel chromosomal translocation t(3;7)(q26;q21) in myeloid leukemia resulting in overexpression of EVI1. <i>Annals of Hematology</i> , 2004, 83, 78-83.	1.8	26
72	Insertions generating the 5'RUNX1/3'CBFA2T1 gene in acute myeloid leukemia cases show variable breakpoints. <i>Genes Chromosomes and Cancer</i> , 2004, 41, 86-91.	2.8	19

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73	Derivative Chromosome 9 Deletions in Chronic Myeloid Leukemia are Associated with Loss of Tumor Suppressor Genes. <i>Leukemia and Lymphoma</i> , 2004, 45, 689-694.	1.3	17
74	Genomic deletions on other chromosomes involved in variant t(9;22) chronic myeloid leukemia cases. <i>Genes Chromosomes and Cancer</i> , 2003, 36, 353-360.	2.8	17
75	Deletions on der(9) chromosome in adult Ph-positive acute lymphoblastic leukemia occur with a frequency similar to that observed in chronic myeloid leukemia. <i>Leukemia</i> , 2003, 17, 528-531.	7.2	8
76	A novel translocation t(14;15)(q32;q24) bearing deletion on der(14) in Philadelphia-positive chronic myeloid leukemia. <i>Haematologica</i> , 2003, 88, 1076-7.	3.5	1
77	A 76-kb duplicon maps close to the BCR gene on chromosome 22 and the ABL gene on chromosome 9: Possible involvement in the genesis of the Philadelphia chromosome translocation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 9882-9887.	7.1	71
78	Molecular cytogenetic characterization of a complex rearrangement involving chromosomes 9 and 22 in a case of Ph-negative chronic myeloid leukemia. <i>Cancer Genetics and Cytogenetics</i> , 2002, 136, 141-145.	1.0	12
79	Breakpoint characterization of der(9) deletions in chronic myeloid leukemia patients. <i>Genes Chromosomes and Cancer</i> , 2002, 35, 271-276.	2.8	48
80	Molecular cytogenetics characterization of a novel translocation involving chromosomes 17 and 19 in a Ph+ adult acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , 2002, 119, 488-491.	2.5	1