

Rolf Marschalek

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9512846/publications.pdf>

Version: 2024-02-01

258
papers

7,637
citations

71102

41
h-index

71685

76
g-index

267
all docs

267
docs citations

267
times ranked

8415
citing authors

#	ARTICLE	IF	CITATIONS
1	Lineage switch to acute myeloid leukemia during induction chemotherapy for early T-cell precursor acute lymphoblastic leukemia with the translocation t(6;11)(q27;q23)/KMT2A-AFDN: A case report. <i>Leukemia Research</i> , 2022, 112, 106758.	0.8	5
2	Vaccine-induced COVID-19 mimicry syndrome. <i>ELife</i> , 2022, 11, .	6.0	41
3	Image-Based Annotation of Chemogenomic Libraries for Phenotypic Screening. <i>Molecules</i> , 2022, 27, 1439.	3.8	19
4	The differential role of the lipid raft-associated protein flotillin 2 for progression of myeloid leukemia. <i>Blood Advances</i> , 2022, 6, 3611-3624.	5.2	6
5	KMT2A-MLLT1 and the Novel SEC16A-KMT2A in a Cryptic 3-Way Translocation t(9;11;19) Present in an Infant With Acute Lymphoblastic Leukemia. <i>Journal of Pediatric Hematology/Oncology</i> , 2022, 44, e719-e722.	0.6	1
6	MLL-AF4 and a murinized pSer-variant thereof are turning on the nucleolar stress pathway. <i>Cell and Bioscience</i> , 2022, 12, 47.	4.8	2
7	Detection and Quantification of SARS-CoV-2 by Real-Time RT-PCR Assay. <i>Methods in Molecular Biology</i> , 2022, 2452, 75-98.	0.9	2
8	The immune checkpoint ICOSLG is a relapse-predicting biomarker and therapeutic target in infant t(4;11) acute lymphoblastic leukemia. <i>IScience</i> , 2022, , 104613.	4.1	6
9	MLL-r fusion transcripts in healthy individuals by induced gene proximity. <i>Klinische Padiatrie</i> , 2022, , .	0.6	0
10	The immune checkpoint ICOSLG is a relapse-predicting biomarker and therapeutic target in infant t(4;11) ALL. <i>Klinische Padiatrie</i> , 2022, , .	0.6	0
11	Designing specific chromosomal translocations of the MLL/KMT2A gene. <i>Klinische Padiatrie</i> , 2022, , .	0.6	0
12	Unravelling the recombinome of IKZF1 deletions in B-ALL. <i>Klinische Padiatrie</i> , 2022, , .	0.6	0
13	Epigenetic regulator genes direct lineage switching in <i>MLL/AF4</i> leukemia. <i>Blood</i> , 2022, 140, 1875-1890.	1.4	26
14	Outcomes for Australian children with relapsed/refractory acute lymphoblastic leukaemia treated with blinatumomab. <i>Pediatric Blood and Cancer</i> , 2021, 68, e28922.	1.5	16
15	Clinical Implications of Minimal Residual Disease Detection in Infants With <i>KMT2A</i> -Rearranged Acute Lymphoblastic Leukemia Treated on the Interfant-06 Protocol. <i>Journal of Clinical Oncology</i> , 2021, 39, 652-662.	1.6	41
16	Generation of a Sleeping Beauty Transposon-Based Cellular System for Rapid and Sensitive Screening for Compounds and Cellular Factors Limiting SARS-CoV-2 Replication. <i>Frontiers in Microbiology</i> , 2021, 12, 701198.	3.5	27
17	Anti-platelet factor 4 antibodies causing VITT do not cross-react with SARS-CoV-2 spike protein. <i>Blood</i> , 2021, 138, 1269-1277.	1.4	102
18	Therapy-related acute myeloid leukemia with KMT2A-SNX9 gene fusion associated with a hyperdiploid karyotype after hemophagocytic lymphohistiocytosis. <i>Cancer Genetics</i> , 2021, 256-257, 86-90.	0.4	3

#	ARTICLE	IF	CITATIONS
19	KMT2A-ARHGEF12, a therapy related fusion with poor prognosis. <i>Molecular Biology Reports</i> , 2021, 48, 7021-7027.	2.3	2
20	The role of reciprocal fusions in MLL-r acute leukemia: studying the chromosomal translocation t(6;11). <i>Oncogene</i> , 2021, 40, 5902-5912.	5.9	1
21	In vivo inducible reverse genetics in patients' tumors to identify individual therapeutic targets. <i>Nature Communications</i> , 2021, 12, 5655.	12.8	10
22	The role of reciprocal fusions in MLL-r acute leukemia: studying the chromosomal translocation t(4;11). <i>Oncogene</i> , 2021, 40, 6093-6102.	5.9	6
23	<i>KMT2A-CBL</i> rearrangements in acute leukemias: clinical characteristics and genetic breakpoints. <i>Blood Advances</i> , 2021, 5, 5617-5620.	5.2	1
24	Closantel is an allosteric inhibitor of human Taspase1. <i>IScience</i> , 2021, 24, 103524.	4.1	1
25	Implication of ICOSLG on Relapse in Infant T(4;11) Acute Lymphoblastic Leukemia. <i>Blood</i> , 2021, 138, 3481-3481.	1.4	0
26	Targeted Next Generation Sequencing Reveals a Third Breakpoint Cluster Region and New Partner Genes in the <i>KMT2A</i> Recombinome. <i>Blood</i> , 2021, 138, 3327-3327.	1.4	0
27	The reciprocal world of MLL fusions: A personal view. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2020, 1863, 194547.	1.9	12
28	Clinical Implications of Minimal Residual Disease Detection in Infants with <i>KMT2A</i> -Rearranged Acute Lymphoblastic Leukemia Treated on the Interfant-06 Protocol. <i>Blood</i> , 2020, 136, 41-42.	1.4	1
29	Investigating the Roles of the Yeats Domain in MLL-ENL Mediated Leukemogenesis. <i>Blood</i> , 2020, 136, 30-31.	1.4	0
30	Prenatal origin of <i>KRAS</i> mutation in a child with an acute myelomonocytic leukaemia bearing the <i>KMT2A</i> / <i>MLL</i> - <i>AFDN</i> / <i>MLLT4</i> / <i>AF6</i> fusion transcript. <i>British Journal of Haematology</i> , 2019, 185, 563-566.	2.5	6
31	Enhanced hemato-endothelial specification during human embryonic differentiation through developmental cooperation between <i>AF4-MLL</i> and <i>MLL-AF4</i> fusions. <i>Haematologica</i> , 2019, 104, 1189-1201.	3.5	15
32	Another piece of the puzzle added to understand t(4;11) leukemia better. <i>Haematologica</i> , 2019, 104, 1098-1100.	3.5	6
33	Human MLL/ <i>KMT2A</i> gene exhibits a second breakpoint cluster region for recurrent <i>MLL</i> - <i>USP2</i> fusions. <i>Leukemia</i> , 2019, 33, 2306-2340.	7.2	41
34	<i>IKZF1</i> Deletions with <i>COBL</i> Breakpoints Are Not Driven by RAG-Mediated Recombination Events in Acute Lymphoblastic Leukemia. <i>Translational Oncology</i> , 2019, 12, 726-732.	3.7	7
35	Genetically engineered CAR NK cells display selective cytotoxicity against FLT3-positive B-ALL and inhibit <i>in vivo</i> leukemia growth. <i>International Journal of Cancer</i> , 2019, 145, 1935-1945.	5.1	60
36	A New Complex Karyotype Involving a <i>KMT2A</i> -r Variant Three-Way Translocation in a Rare Clinical Presentation of a Pediatric Patient with Acute Myeloid Leukemia. <i>Cytogenetic and Genome Research</i> , 2019, 157, 213-219.	1.1	0

#	ARTICLE	IF	CITATIONS
37	Acute myeloid leukemia with t(10;11)(p11;q23.3): Results of Russian Pediatric AML registration study. <i>International Journal of Laboratory Hematology</i> , 2019, 41, 287-292.	1.3	6
38	Inhibition of PIM1 blocks the autophagic flux to sensitize glioblastoma cells to ABT-737-induced apoptosis. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2019, 1866, 175-189.	4.1	17
39	Frequent and reliable engraftment of certain adult primary acute lymphoblastic leukemias in mice. <i>Leukemia and Lymphoma</i> , 2019, 60, 848-851.	1.3	4
40	Biology of t(6;11) Fusion Proteins and Their Role in MLL-Rearranged Acute Leukemia Lineage Determination. <i>Blood</i> , 2019, 134, 5033-5033.	1.4	0
41	The MLL recombinome of acute leukemias in 2017. <i>Leukemia</i> , 2018, 32, 273-284.	7.2	527
42	How to effectively treat acute leukemia patients bearing MLL-rearrangements ?. <i>Biochemical Pharmacology</i> , 2018, 147, 183-190.	4.4	23
43	Targeted Next-Generation Sequencing for Detecting MLL Gene Fusions in Leukemia. <i>Molecular Cancer Research</i> , 2018, 16, 279-285.	3.4	27
44	A new complex rearrangement in infant ALL: t(X;11;17)(p11.2;q23;q12). <i>Cancer Genetics</i> , 2018, 228-229, 110-114.	0.4	2
45	A case of pediatric acute myeloid leukemia with t(11;16)(q23;q24) leading to a novel KMT2A-USP10 fusion gene. <i>Genes Chromosomes and Cancer</i> , 2018, 57, 522-524.	2.8	7
46	MLL-USP2: An Underestimated New Entity of MLL-Rearranged Leukemia Identified By NGS Analysis. <i>Blood</i> , 2018, 132, 3920-3920.	1.4	2
47	Quantitative Analysis of MLL Fusion Transcripts By Droplet Digital PCR to Monitor Minimal Residual Disease in MLL-Rearranged Acute Myeloid Leukemia. <i>Blood</i> , 2018, 132, 2746-2746.	1.4	0
48	Molecular characterization of KMT2A fusion partner genes in 13 cases of pediatric leukemia with complex or cryptic karyotypes. <i>Hematological Oncology</i> , 2017, 35, 760-768.	1.7	9
49	Characterization and cellular localization of human 5-lipoxygenase and its protein isoforms 5-LO ¹³ , 5-LO ⁴ and 5-LO ¹² . <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2017, 1862, 561-571.	2.4	4
50	Hypoxia Causes Downregulation of Dicer in Hepatocellular Carcinoma, Which Is Required for Upregulation of Hypoxia-Inducible Factor 1 α and Epithelial-Mesenchymal Transition. <i>Clinical Cancer Research</i> , 2017, 23, 3896-3905.	7.0	33
51	Evolution of AF6-RAS association and its implications in mixed-lineage leukemia. <i>Nature Communications</i> , 2017, 8, 1099.	12.8	21
52	TGF β /SMAD signalling modulates MLL and MLL-AF4 mediated 5-lipoxygenase promoter activation. <i>Prostaglandins and Other Lipid Mediators</i> , 2017, 133, 60-67.	1.9	6
53	A Case of Acute Myeloid Leukemia with Novel Translocation t(6;11)(p22.2;q23) and Concurrent Insertion ins(11;9)(q23;p21.3p21.3). <i>Advances in Experimental Medicine and Biology</i> , 2017, 1021, 93-98.	1.6	1
54	MLL α 1, 2017, . .		0

#	ARTICLE	IF	CITATIONS
55	The AF4-MLL fusion transiently augments multilineage hematopoietic engraftment but is not sufficient to initiate leukemia in cord blood CD34+ cells. <i>Oncotarget</i> , 2017, 8, 81936-81941.	1.8	13
56	Genetic characteristics determine lentiviral transduction rates in patient-derived ALL cells. , 2017, 229, .		0
57	Systematic Classification of Mixed-Lineage Leukemia Fusion Partners Predicts Additional Cancer Pathways. <i>Annals of Laboratory Medicine</i> , 2016, 36, 85-100.	2.5	57
58	Identification of a Cryptic Insertion ins(11;X)(q23;q28q12) Resulting in a <i>KMT2A</i>-<i>FLNA</i> Fusion in a 13-Month-Old Child with Acute Myelomonocytic Leukemia. <i>Cytogenetic and Genome Research</i> , 2016, 150, 281-286.	1.1	2
59	Identification and characterization of the elusive mutation causing the historical von Willebrand Disease type IIC Miami. <i>Journal of Thrombosis and Haemostasis</i> , 2016, 14, 1725-1735.	3.8	9
60	Activated <i>KRAS</i> Cooperates with MLL-AF4 to Promote Extramedullary Engraftment and Migration of Cord Blood CD34+ HSPC But Is Insufficient to Initiate Leukemia. <i>Cancer Research</i> , 2016, 76, 2478-2489.	0.9	37
61	Comparison of MRD Levels and Gene Expression Patterns in MLL-R Versus Non-MLL Infant ALL. <i>Blood</i> , 2016, 128, 1740-1740.	1.4	1
62	<i>COBL</i> is a novel hotspot for <i>IKZF1</i> deletions in childhood acute lymphoblastic leukemia. <i>Oncotarget</i> , 2016, 7, 53064-53073.	1.8	9
63	Expression of MLL-AF4 or AF4-MLL fusions does not impact the efficiency of DNA damage repair. <i>Oncotarget</i> , 2016, 7, 30440-30452.	1.8	19
64	The IRX1/HOXA connection: insights into a novel t(4;11)- specific cancer mechanism. <i>Oncotarget</i> , 2016, 7, 35341-35352.	1.8	22
65	Identification of Multiple, Patient-Specific MLL Fusion Transcript Isoforms in Childhood Leukemia Using Anchored Multiplex PCR-Based Enrichment (AMP-E). <i>Blood</i> , 2016, 128, 2908-2908.	1.4	0
66	DDX6 transfers P-TEFb kinase to the AF4/AF4N (AFF1) super elongation complex. <i>American Journal of Blood Research</i> , 2016, 6, 28-45.	0.6	17
67	Revisiting the biology of infant t(4;11)/MLL-AF4+ B-cell acute lymphoblastic leukemia. <i>Blood</i> , 2015, 126, 2676-2685.	1.4	100
68	MLL Leukemia and Future Treatment Strategies. <i>Archiv Der Pharmazie</i> , 2015, 348, 221-228.	4.1	29
69	Long-term remission of therapy-related acute myeloid leukemia with a new t(11;18)(q23;q21.2) translocation and KMT2A-ME2 (MLL-ME2) fusion gene. <i>Cancer Genetics</i> , 2015, 208, 610-614.	0.4	1
70	Unraveling the Activation Mechanism of Taspase1 which Controls the Oncogenic AF4-MLL Fusion Protein. <i>EBioMedicine</i> , 2015, 2, 386-395.	6.1	9
71	Optimized Sleeping Beauty transposons rapidly generate stable transgenic cell lines. <i>Biotechnology Journal</i> , 2015, 10, 647-653.	3.5	352
72	Subclonality and prenatal origin of <i>RAS</i> mutations in <i>KMT2A</i> (<i>MLL</i>)-rearranged infant acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , 2015, 170, 268-271.	2.5	23

#	ARTICLE	IF	CITATIONS
73	Effective Targeting of the P53-MDM2 Axis in Preclinical Models of Infant <i>MLL</i> -Rearranged Acute Lymphoblastic Leukemia. <i>Clinical Cancer Research</i> , 2015, 21, 1395-1405.	7.0	43
74	Evidence-based RT-PCR methods for the detection of the 8 most common <i>MLL</i> aberrations in acute leukemias. <i>Leukemia Research</i> , 2015, 39, 242-247.	0.8	22
75	Molecular studies reveal a <i>MLL</i> - <i>MLLT3</i> gene fusion displaced in a case of childhood acute lymphoblastic leukemia with complex karyotype. <i>Cancer Genetics</i> , 2015, 208, 143-147.	0.4	6
76	A new variant of <i>KMT2A(MLL)</i> - <i>FLNA</i> fusion transcript in acute myeloid leukemia with <i>ins(X;11)(q28;q23q23)</i> . <i>Cancer Genetics</i> , 2015, 208, 148-151.	0.4	9
77	<i>AF4</i> and <i>AF4-MLL</i> mediate transcriptional elongation of 5-lipoxygenase mRNA by 1, 25-dihydroxyvitamin D3. <i>Oncotarget</i> , 2015, 6, 25784-25800.	1.8	11
78	<i>AF4</i> and <i>AF4N</i> protein complexes: recruitment of P-TEFb kinase, their interactome and potential functions. <i>American Journal of Blood Research</i> , 2015, 5, 10-24.	0.6	9
79	Inhibition of class I HDACs abrogates the dominant effect of <i>MLL</i> - <i>AF4</i> by activation of wild-type <i>MLL</i> . <i>Oncogenesis</i> , 2014, 3, e127-e127.	4.9	22
80	<i>MLL</i> - <i>SEPT5</i> fusion transcript in infant acute myeloid leukemia with <i>t(11;22)(q23;q11)</i> . <i>Leukemia and Lymphoma</i> , 2014, 55, 662-667.	1.3	10
81	Ceramide synthases <i>CerS4</i> and <i>CerS5</i> are upregulated by 17 β -estradiol and <i>GPBR1</i> via AP-1 in human breast cancer cells. <i>Biochemical Pharmacology</i> , 2014, 92, 577-589.	4.4	37
82	Complex <i>MLL</i> rearrangement in non-infiltrated bone marrow in an infant with stage II precursor B-lymphoblastic lymphoma. <i>European Journal of Haematology</i> , 2014, 93, 349-353.	2.2	5
83	<i>MLL</i> partner genes in secondary acute lymphoblastic leukemia: Report of a new partner <i>PRRC1</i> and review of the literature. <i>Leukemia Research</i> , 2014, 38, 1316-1319.	0.8	7
84	Unique <i>BHLHB3</i> overexpression in pediatric acute myeloid leukemia with <i>t(6;11)(q27;q23)</i> . <i>Leukemia</i> , 2014, 28, 1564-1568.	7.2	3
85	Functional characterisation of different <i>MLL</i> fusion proteins by using inducible Sleeping Beauty vectors. <i>Cancer Letters</i> , 2014, 352, 196-202.	7.2	34
86	Ligand-independent <i>FLT3</i> activation does not cooperate with <i>MLL</i> - <i>AF4</i> to immortalize/transform cord blood CD34+ cells. <i>Leukemia</i> , 2014, 28, 666-674.	7.2	27
87	Secondary acute monocytic leukemia positive for 11q23 rearrangement in Nijmegen breakage syndrome. <i>Pediatric Blood and Cancer</i> , 2014, 61, 1469-1471.	1.5	4
88	Molecular characterization and clinical course of <i>MLL</i> - <i>ACTN4</i> rearrangement in therapy-related hematologic malignancies. <i>Haematologica</i> , 2014, 99, e49-e51.	3.5	8
89	Molecular characterization and clinical impact of <i>t(11;15)(q23;q14-15)</i> <i>MLL</i> - <i>CASC5</i> rearrangement. <i>Haematologica</i> , 2014, 99, e11-e13.	3.5	8
90	From a Multipotent Stilbene to Soluble Epoxide Hydrolase Inhibitors with Antiproliferative Properties. <i>ChemMedChem</i> , 2013, 8, 919-923.	3.2	25

#	ARTICLE	IF	CITATIONS
91	RE: Acute myeloid leukemia associated with FGFR1 abnormalities. International Journal of Hematology, 2013, 98, 139-140.	1.6	0
92	Complex Three-Way Translocation Involving <i>MLL</i> , <i>ELL</i> , <i>RREB1</i> , and <i>CMAHP</i> Genes in an Infant with Acute Myeloid Leukemia and <i>t(6;19;11)(p22.2;p13.1;q23.3)</i> . Cytogenetic and Genome Research, 2013, 141, 7-15.	1.1	11
93	Functional analysis of the two reciprocal fusion genes <i>MLL-NEBL</i> and <i>NEBL-MLL</i> reveal their oncogenic potential. Cancer Letters, 2013, 332, 30-34.	7.2	23
94	The <i>MLL</i> recombinome of acute leukemias in 2013. Leukemia, 2013, 27, 2165-2176.	7.2	393
95	Analyzing acute leukemia patients with complex <i>MLL</i> rearrangements by a sequential LDI-PCR approach. Cancer Letters, 2013, 338, 249-254.	7.2	4
96	<i>MLL</i> , 2013, , 452-455.		0
97	The distribution of <i>MLL</i> breakpoints correlates with outcome in infant acute leukaemia. British Journal of Haematology, 2013, 161, 224-236.	2.5	46
98	Acute Promyelocytic Leukemia with a Rare <i>PML</i> ; Exon 4/ <i>RARA</i> ; Exon 3 Fusion Transcript Variant. Acta Haematologica, 2013, 130, 176-180.	1.4	2
99	Secondary mutations in <i>t(4;11)</i> leukemia patients. Leukemia, 2013, 27, 1425-1427.	7.2	26
100	Genomic breakpoints and clinical features of <i>MLL-TET1</i> rearrangement in acute leukemias. Haematologica, 2013, 98, e55-e57.	3.5	17
101	Clone-specific secondary aberrations are not detected in neonatal blood spots of children with <i>ETV6-RUNX1</i> -positive leukemia. Haematologica, 2013, 98, e108-e110.	3.5	4
102	An alternative splice process renders <i>MLL</i> either into an transcriptional activator or repressor. Klinische Padiatrie, 2013, 225, 601-7.	0.6	15
103	The Genomic Landscape Of Lineage Switch Acute Leukemia. Blood, 2013, 122, 2552-2552.	1.4	2
104	<i>MLL</i> genomic DNA Breakpoints In Infant Acute Leukemia. Blood, 2013, 122, 1350-1350.	1.4	3
105	Backtracking Of The Genetic Events Preceding Acute Leukemia Diagnosis. Blood, 2013, 122, 2562-2562.	1.4	2
106	Refinement of <i>IKZF1</i> recombination hotspots in pediatric BCP-ALL patients. American Journal of Blood Research, 2013, 3, 165-73.	0.6	16
107	Do Non-Genomically Encoded Fusion Transcripts Cause Recurrent Chromosomal Translocations?. Cancers, 2012, 4, 1036-1049.	3.7	12
108	Long chain ceramides and very long chain ceramides have opposite effects on human breast and colon cancer cell growth. International Journal of Biochemistry and Cell Biology, 2012, 44, 620-628.	2.8	178

#	ARTICLE	IF	CITATIONS
109	A three-way translocation of MLL, MLLT11, and the novel reciprocal partner gene MYO18A in a child with acute myeloid leukemia. <i>Cancer Genetics</i> , 2012, 205, 261-265.	0.4	14
110	Involvement of the <i>MLL</i> gene in adult T-lymphoblastic leukemia. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 1114-1124.	2.8	14
111	<i>CBL</i> mutations do not frequently occur in paediatric acute myeloid leukaemia. <i>British Journal of Haematology</i> , 2012, 159, 577-584.	2.5	7
112	Genetic and clinical characterization of 45 acute leukemia patients with <i>MLL</i> gene rearrangements from a single institution. <i>Molecular Oncology</i> , 2012, 6, 553-564.	4.6	19
113	Submicroscopic Deletion of <i>FGFR1</i> Gene Is Recurrently Detected in Myeloid and Lymphoid Neoplasms Associated with <i>ZMYM2-FGFR1</i> Rearrangements: A Case Study. <i>Acta Haematologica</i> , 2012, 127, 119-123.	1.4	12
114	Acute promyelocytic leukemia with trisomy 8 showing normal PML-RARA FISH signal patterns: diagnostic application of long-distance polymerase chain reaction in molecularly discrepant leukemia cases. <i>Annals of Hematology</i> , 2012, 91, 1645-1648.	1.8	6
115	<i>MLL</i> - <i>ELL</i> fusion gene in two infants with acute monoblastic leukemia and myeloid sarcoma. <i>Leukemia and Lymphoma</i> , 2012, 53, 1222-1224.	1.3	7
116	Molecular methods for genomic analyses of variant <i>PML-RARA</i> or other <i>RARA</i> -related chromosomal translocations in acute promyelocytic leukemia. <i>The Korean Journal of Hematology</i> , 2012, 47, 307.	0.7	5
117	Diagnostic Usefulness of Genomic Breakpoint Analysis of Various Gene Rearrangements in Acute Leukemias: A Perspective of Long Distance or Long Distance Inverse-PCR based Approaches. <i>Annals of Laboratory Medicine</i> , 2012, 32, 316-318.	2.5	15
118	Letter to the Editor: Detection of <i>EML4-ALK</i> and Other <i>ALK</i> Fusion Genes in Lung Cancer: A Lesson from the Leukemia Fusion Gene Analysis and Future Application. <i>Journal of Korean Medical Science</i> , 2012, 27, 576.	2.5	0
119	Editorial: Pharmazie in unserer Zeit 3/2012. <i>Pharmazie in Unserer Zeit</i> , 2012, 41, 179-179.	0.0	0
120	Genomic analysis of a four-way t(4;11;22;10) associated with MLL-AF4 in an adult acute lymphoblastic leukemia. <i>Annals of Hematology</i> , 2012, 91, 977-979.	1.8	4
121	Abl-interactor 2 (ABI2): A novel MLL translocation partner in acute myeloid leukemia. <i>Leukemia Research</i> , 2012, 36, e113-e115.	0.8	6
122	<i>TOP3A</i> , a new partner gene fused to <i>MLL</i> in an adult patient with de novo acute myeloid leukaemia. <i>British Journal of Haematology</i> , 2012, 157, 128-131.	2.5	5
123	Epigenetics and blood disorders. <i>British Journal of Haematology</i> , 2012, 158, 307-322.	2.5	11
124	Molecular Genetic Characterization of 3'-Deletion of MLL Gene in Infant Acute Leukemia. <i>Blood</i> , 2012, 120, 2498-2498.	1.4	0
125	MLL Gene Rearrangements in 174 Infants with Acute Leukemia. <i>Blood</i> , 2012, 120, 2537-2537.	1.4	0
126	A complex MLL rearrangement identified five years after initial MDS diagnosis results in out-of-frame fusions without progression to acute leukemia. <i>Cancer Genetics</i> , 2011, 204, 557-562.	0.4	10

#	ARTICLE	IF	CITATIONS
127	Characterisation of two novel large F8 deletions in patients with severe haemophilia A and factor VIII inhibitors. <i>Thrombosis and Haemostasis</i> , 2011, 105, 279-284.	3.4	3
128	Bioassays to Monitor Taspase1 Function for the Identification of Pharmacogenetic Inhibitors. <i>PLoS ONE</i> , 2011, 6, e18253.	2.5	25
129	Nerve Injury Evoked Loss of Latexin Expression in Spinal Cord Neurons Contributes to the Development of Neuropathic Pain. <i>PLoS ONE</i> , 2011, 6, e19270.	2.5	9
130	ETV6/RUNX1-positive relapses evolve from an ancestral clone and frequently acquire deletions of genes implicated in glucocorticoid signaling. <i>Blood</i> , 2011, 117, 2658-2667.	1.4	83
131	High IGSF4 expression in pediatric M5 acute myeloid leukemia with t(9;11)(p22;q23). <i>Blood</i> , 2011, 117, 928-935.	1.4	17
132	Evaluation of gene expression signatures predictive of cytogenetic and molecular subtypes of pediatric acute myeloid leukemia. <i>Haematologica</i> , 2011, 96, 221-230.	3.5	98
133	Mechanisms of leukemogenesis by MLL fusion proteins. <i>British Journal of Haematology</i> , 2011, 152, 141-154.	2.5	158
134	The leukemogenic AF4-MLL fusion protein causes P-TEFb kinase activation and altered epigenetic signatures. <i>Leukemia</i> , 2011, 25, 135-144.	7.2	73
135	The heterodimerization domains of MLL-FYRN and FYRC are potential target structures in t(4;11) leukemia. <i>Leukemia</i> , 2011, 25, 663-670.	7.2	31
136	Backtracking to birth of the NUP98-HOXD13 gene fusion in an infant acute myeloid leukemia. <i>Leukemia</i> , 2011, 25, 1192-1194.	7.2	8
137	Oncogenic FAM131B-BRAF fusion resulting from 7q34 deletion comprises an alternative mechanism of MAPK pathway activation in pilocytic astrocytoma. <i>Acta Neuropathologica</i> , 2011, 121, 763-774.	7.7	211
138	TEL/AML1-positive patients lacking TEL exon 5 resemble canonical TEL/AML1 cases. <i>Pediatric Blood and Cancer</i> , 2011, 56, 217-225.	1.5	9
139	KIAA1524: A novel MLL translocation partner in acute myeloid leukemia. <i>Leukemia Research</i> , 2011, 35, 133-135.	0.8	21
140	It takes two-to-leukemia: About addictions and requirements. <i>Leukemia Research</i> , 2011, 35, 424-425.	0.8	5
141	A novel PML-ADAMTS17-RARA gene rearrangement in a patient with pregnancy-related acute promyelocytic leukemia. <i>Leukemia Research</i> , 2011, 35, e106-e110.	0.8	15
142	De novo Acute Myeloid Leukemia Associated with t(11;17)(q23;q25) and MLL-SEPT9 Rearrangement in an Elderly Patient: A Case Study and Review of the Literature. <i>Acta Haematologica</i> , 2011, 126, 195-198.	1.4	11
143	Cell-based Analysis of Structure-Function Activity of Threonine Aspartase 1. <i>Journal of Biological Chemistry</i> , 2011, 286, 3007-3017.	3.4	45
144	What Is New? An Update of the MLL Recombinome Including the Three Novel Partner Genes ABI2, PDS5A, and TOP3A. <i>Blood</i> , 2011, 118, 1351-1351.	1.4	0

#	ARTICLE	IF	CITATIONS
145	Premature transcript termination, trans-splicing and DNA repair: a vicious path to cancer. American Journal of Blood Research, 2011, 1, 1-12.	0.6	18
146	The AF4-MLL fusion protein is capable of inducing ALL in mice without requirement of MLL-AF4. Blood, 2010, 115, 3570-3579.	1.4	133
147	Three-way translocation involving MLL, MLLT1, and a novel third partner, NRXN1, in a patient with acute lymphoblastic leukemia and t(2;19;11) (p12;p13.3;q23). Cancer Genetics and Cytogenetics, 2010, 197, 32-38.	1.0	13
148	Nebulette is the second member of the nebulin family fused to the MLL gene in infant leukemia. Cancer Genetics and Cytogenetics, 2010, 198, 151-154.	1.0	11
149	Acute leukemias with <i>ETV6/ABL1</i> (<i>TEL/ABL</i>) fusion: Poor prognosis and prenatal origin. Genes Chromosomes and Cancer, 2010, 49, 873-884.	2.8	36
150	BTBD18: A novel MLL partner gene in an infant with acute lymphoblastic leukemia and inv(11)(q13;q23). Leukemia Research, 2010, 34, e294-e296.	0.8	8
151	Mixed lineage leukemia: roles in human malignancies and potential therapy. FEBS Journal, 2010, 277, 1822-1831.	4.7	55
152	Acute monocytic leukaemia originating from <i>MLL-MLLT3</i> -positive pre-B cells. British Journal of Haematology, 2010, 150, 621-623.	2.5	10
153	Transcriptional properties of human NANOG1 and NANOG2 in acute leukemic cells. Nucleic Acids Research, 2010, 38, 5384-5395.	14.5	55
154	Therapy-related acute myeloid leukemia with t(2;11)(q37;q23) after treatment for osteosarcoma. Cancer Genetics and Cytogenetics, 2010, 203, 288-291.	1.0	9
155	FISH-negative cryptic PML-RARA rearrangement detected by long-distance polymerase chain reaction and sequencing analyses: a case study and review of the literature. Cancer Genetics and Cytogenetics, 2010, 203, 278-283.	1.0	59
156	High BRE expression in pediatric MLL-rearranged AML is associated with favorable outcome. Leukemia, 2010, 24, 2048-2055.	7.2	27
157	Complex and cryptic chromosomal rearrangements involving the MLL gene in acute leukemia: A study of 7 patients and review of the literature. Blood Cells, Molecules, and Diseases, 2010, 44, 268-274.	1.4	27
158	Identification of a <i>MLL-MLLT4</i> fusion gene resulting from a t(6;11)(q27;q23) presenting as a del(11q) in a child with T-cell acute lymphoblastic leukemia. Leukemia and Lymphoma, 2010, 51, 1570-1573.	1.3	7
159	High IGSF4 expression In Pediatric Acute Monoblastic Leukemia with t(9;11). Blood, 2010, 116, 3614-3614.	1.4	0
160	An Interstitial Deletion at 3p21.3 Results in the Genetic Fusion of <i>MLH1</i> and <i>ITGA9</i> in a Lynch Syndrome Family. Clinical Cancer Research, 2009, 15, 762-769.	7.0	14
161	CASP8AP2 is a novel partner gene of MLL rearrangement with t(6;11)(q15;q23) in acute myeloid leukemia. Cancer Genetics and Cytogenetics, 2009, 195, 94-95.	1.0	10
162	Covert preleukemia driven by <i>MLL</i> gene fusion. Genes Chromosomes and Cancer, 2009, 48, 98-107.	2.8	14

#	ARTICLE	IF	CITATIONS
163	A complex 1;19;11 translocation involving the MLL gene in a patient with congenital acute monoblastic leukemia identified by molecular and cytogenetic techniques. <i>Annals of Hematology</i> , 2009, 88, 795-797.	1.8	8
164	<i>FLNA</i> , a new partner gene fused to <i>MLL</i> in a patient with acute myelomonoblastic leukaemia. <i>British Journal of Haematology</i> , 2009, 146, 693-695.	2.5	10
165	Prognostic significance of minimal residual disease in infants with acute lymphoblastic leukemia treated within the Interfant-99 protocol. <i>Leukemia</i> , 2009, 23, 1073-1079.	7.2	137
166	New insights to the MLL recombinome of acute leukemias. <i>Leukemia</i> , 2009, 23, 1490-1499.	7.2	363
167	Infant acute bilineal leukemia. <i>Leukemia Research</i> , 2009, 33, 1005-1008.	0.8	9
168	NRIP3: a novel translocation partner of MLL detected in a pediatric acute myeloid leukemia with complex chromosome 11 rearrangements. <i>Haematologica</i> , 2009, 94, 1033-1033.	3.5	12
169	LDI-PCR: Identification of Known and Unknown Gene Fusions of the Human MLL Gene. <i>Methods in Molecular Biology</i> , 2009, 538, 71-83.	0.9	12
170	Two independent gene signatures in pediatric t(4;11) acute lymphoblastic leukemia patients. <i>European Journal of Haematology</i> , 2009, 83, 406-419.	2.2	51
171	The MLL recombinome of adult CD10-negative B-cell precursor acute lymphoblastic leukemia: results from the GMALL study group. <i>Blood</i> , 2009, 113, 4011-4015.	1.4	85
172	WT1 Gene Expression Limited Application for MRD Monitoring in AML Patients Carrying MLL Rearrangements.. <i>Blood</i> , 2009, 114, 4700-4700.	1.4	0
173	The Central Region of TEL (ETV6) Is Dispensable for the TEL/AML1 (ETV6/RUNX1) Leukemogenesis.. <i>Blood</i> , 2009, 114, 1601-1601.	1.4	0
174	Acute Leukaemias with TEL/ABL Fusion: a Subgroup with Poor Prognosis and Prenatal Origin.. <i>Blood</i> , 2009, 114, 2596-2596.	1.4	0
175	Hypomethylation of Tumor Suppressor Genes in Acute Myeloid Leukemia: Characteristic of Cell Lines with MLL Abnormalities?.. <i>Blood</i> , 2009, 114, 365-365.	1.4	0
176	Inv(11)(q21q23) fuses MLL to the Notch co-activator mastermind-like 2 in secondary T-cell acute lymphoblastic leukemia. <i>Leukemia</i> , 2008, 22, 1807-1811.	7.2	23
177	An Alu-mediated novel large deletion is the most frequent cause of type A3 von Willebrand disease in Hungary. <i>Journal of Thrombosis and Haemostasis</i> , 2008, 6, 1729-1735.	3.8	28
178	MLL insertion with MLL-MLLT3 gene fusion in acute leukemia: case report and review of the literature. <i>Cancer Genetics and Cytogenetics</i> , 2008, 183, 53-59.	1.0	5
179	A MLL-KIAA0284 fusion gene in a patient with secondary acute myeloid leukemia and t(11;14)(q23;q32). <i>Blood Cells, Molecules, and Diseases</i> , 2008, 41, 210-214.	1.4	3
180	MRD Monitoring Reveals a Specific Biology of BCR/ABL-Positive ALL. <i>Blood</i> , 2008, 112, 2529-2529.	1.4	2

#	ARTICLE	IF	CITATIONS
181	MRD Monitoring in AML Patients with MLL-MLLT4 by Quantification of Fusion Gene Transcripts and Genomic Chromosomal Breakpoint Sequences. <i>Blood</i> , 2008, 112, 4888-4888.	1.4	1
182	New Insights to the MLL Recombinome Including 8 Novel Partner Genes ACTN4, C2CD3, DCP1A, FLNA, LAMC3, LOC100128568, NRIP3, and TNRC18.. <i>Blood</i> , 2008, 112, 2047-2047.	1.4	1
183	Incidence and Spectrum of MLL Gene Rearrangements in Pediatric Acute Leukemias in Poland. <i>Blood</i> , 2008, 112, 4851-4851.	1.4	0
184	AF4-MLL Expression Is Necessary and Sufficient for Leukemia Onset in Mice. <i>Blood</i> , 2008, 112, 685-685.	1.4	1
185	Transcription linked to recombination: a gene-internal promoter coincides with the recombination hot spot II of the human MLL gene. <i>Oncogene</i> , 2007, 26, 1361-1371.	5.9	49
186	Combined effects of the two reciprocal t(4;11) fusion proteins MLL- \hat{A} AF4 and AF4- \hat{A} MLL confer resistance to apoptosis, cell cycling capacity and growth transformation. <i>Oncogene</i> , 2007, 26, 3352-3363.	5.9	51
187	Spliced MLL fusions: a novel mechanism to generate functional chimeric MLL-MLLT1 transcripts in t(11;19)(q23;p13.3) leukemia. <i>Leukemia</i> , 2007, 21, 588-590.	7.2	32
188	Immunobiological diversity in infant acute lymphoblastic leukemia is related to the occurrence and type of MLL gene rearrangement. <i>Leukemia</i> , 2007, 21, 633-641.	7.2	102
189	Complex MLL rearrangements in t(4;11) leukemia patients with absent AF4 \hat{A} MLL fusion allele. <i>Leukemia</i> , 2007, 21, 1232-1238.	7.2	40
190	Childhood secondary ALL after ALL treatment. <i>Leukemia</i> , 2007, 21, 1431-1435.	7.2	26
191	A common 253 \hat{A} kb deletion involving VWF and TMEM16B in German and Italian patients with severe von Willebrand disease type 3. <i>Journal of Thrombosis and Haemostasis</i> , 2007, 5, 722-728.	3.8	37
192	C/EBP \hat{I} 2 suppression by interruption of CUGBP1 resulting from a complex rearrangement of MLL. <i>Cancer Genetics and Cytogenetics</i> , 2007, 177, 108-114.	1.0	15
193	Three-Way Translocations and Insertions Are the Molecular Basis for Complex MLL Rearrangements.. <i>Blood</i> , 2007, 110, 979-979.	1.4	7
194	Novel Spliced MLL Fusions Have Been Identified Involving the MLL Partner Genes ELL, EPS15, MLLT3, and SEPT5.. <i>Blood</i> , 2007, 110, 978-978.	1.4	0
195	The MLL Recombinome of Adult ALL - Results from the GMALL Study Group.. <i>Blood</i> , 2007, 110, 977-977.	1.4	0
196	A Novel Approach for Analyzing Gene Expression Profiles Defines Two Distinct Subgroups of t(4;11) Positive Infant Acute Lymphoblastic Leukemia Patients.. <i>Blood</i> , 2007, 110, 4282-4282.	1.4	0
197	Prognostic Factors in Adult Patients up to 60 Years with AML and Translocations of Chromosome 11q23: Individual Patient Data Based Analysis of the German AML-Intergroup.. <i>Blood</i> , 2007, 110, 759-759.	1.4	1
198	Monitoring minimal residual disease by quantification of genomic chromosomal breakpoint sequences in acute leukemias with MLL aberrations. <i>Leukemia</i> , 2006, 20, 451-457.	7.2	57

#	ARTICLE	IF	CITATIONS
199	The MLL recombinome of acute leukemias. <i>Leukemia</i> , 2006, 20, 777-784.	7.2	196
200	Genomic DNA of leukemic patients: Target for clinical diagnosis of MLL rearrangements. <i>Biotechnology Journal</i> , 2006, 1, 656-663.	3.5	21
201	Molecular dissection of t(11;17) in acute myeloid leukemia reveals a variety of gene fusions with heterogeneous fusion transcripts and multiple splice variants. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 1041-1049.	2.8	33
202	Mixed Lineage Leukemia—Rearranged Childhood Pro-B and CD10-Negative Pre-B Acute Lymphoblastic Leukemia Constitute a Distinct Clinical Entity. <i>Clinical Cancer Research</i> , 2006, 12, 2988-2994.	7.0	40
203	Prognosis of Adult Patients \geq 60 Years with AML and Aberrations of Chromosome 11q23: Pooled Data Analysis of the German AML-Intergroup. <i>Blood</i> , 2006, 108, 16-16.	1.4	2
204	Inv(11)(q21q23) Fuses MLL to the NOTCH Co-Activator Mastermind-Like 2 in Secondary T Cell Acute Lymphoblastic Leukemia. <i>Blood</i> , 2006, 108, 4284-4284.	1.4	2
205	C/EBP Suppression by Interruption of CUGBP1 Resulting from a Complex Rearrangement of MLL. <i>Blood</i> , 2006, 108, 2080-2080.	1.4	0
206	Clonal expansion of a new MLL rearrangement in the absence of leukemia. <i>Blood</i> , 2005, 105, 4151-4152.	1.4	20
207	Diagnostic tool for the identification of MLL rearrangements including unknown partner genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 449-454.	7.1	175
208	In-Utero Origin of Childhood Acute Leukaemias. <i>Blood</i> , 2005, 106, 1445-1445.	1.4	0
209	A LDI-PCR Based Method Allows the Identification of Any MLL Rearrangement. <i>Blood</i> , 2005, 106, 2848-2848.	1.4	0
210	Childhood Secondary ALL after ALL Treatment. <i>Blood</i> , 2005, 106, 852-852.	1.4	49
211	Recombination at chromosomal sequences involved in leukaemogenic rearrangements is differentially regulated by p53. <i>Carcinogenesis</i> , 2004, 25, 1305-1313.	2.8	18
212	Interaction of AF4 wild-type and AF4-MLL fusion protein with SIAH proteins: indication for t(4;11) pathobiology?. <i>Oncogene</i> , 2004, 23, 6237-6249.	5.9	68
213	Analyzing the MLL Recombinome. <i>Blood</i> , 2004, 104, 4267-4267.	1.4	0
214	NQO1 C609T polymorphism in distinct entities of pediatric hematologic neoplasms. <i>Haematologica</i> , 2004, 89, 1492-7.	3.5	27
215	Multidimensional gas chromatography—mass spectrometry for tracer studies of fatty acid metabolism via stable isotopes in cultured human trophoblast cells. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2003, 791, 235-244.	2.3	12
216	MLL-mediated transcriptional gene regulation investigated by gene expression profiling. <i>Oncogene</i> , 2003, 22, 3655-3668.	5.9	28

#	ARTICLE	IF	CITATIONS
217	A highly specific and sensitive fluorescence in situ hybridization assay for the detection of t(4;11)(q21;q23) and concurrent submicroscopic deletions in acute leukaemias. <i>British Journal of Haematology</i> , 2002, 116, 758-764.	2.5	42
218	Low frequency of clonotypic Ig and T-cell receptor gene rearrangements in t(4;11) infant acute lymphoblastic leukaemia and its implication for the detection of minimal residual disease. <i>British Journal of Haematology</i> , 2002, 117, 315-321.	2.5	24
219	Presence of N regions in the clonotypic DJ rearrangements of the immunoglobulin heavy-chain genes indicates an exquisitely short latency in t(4;11)-positive infant acute lymphoblastic leukemia. <i>Blood</i> , 2001, 98, 2272-2274.	1.4	24
220	Breakpoints of t(4;11) translocations in the human MLL and AF4 genes in ALL patients are preferentially clustered outside of high-affinity matrix attachment regions. <i>Journal of Cellular Biochemistry</i> , 2001, 82, 299-309.	2.6	24
221	Rapid isolation of chromosomal breakpoints from patients with t(4;11) acute lymphoblastic leukemia: implications for basic and clinical research. <i>Leukemia</i> , 2001, 15, 286-288.	7.2	9
222	Cryptic t(4;11) encoding MLL-AF4 due to insertion of 5' MLL sequences in chromosome 4. <i>Leukemia</i> , 2001, 15, 595-600.	7.2	29
223	Biased distribution of chromosomal breakpoints involving the MLL gene in infants versus children and adults with t(4;11) ALL. <i>Oncogene</i> , 2001, 20, 2900-2907.	5.9	76
224	Fatal connections: When DNA ends meet on the nuclear matrix. <i>Journal of Cellular Biochemistry</i> , 2000, 79, 3-22.	2.6	1
225	Presence of clone-specific antigen receptor gene rearrangements at birth indicates an in utero origin of diverse types of early childhood acute lymphoblastic leukemia. <i>Blood</i> , 2000, 95, 2722-4.	1.4	18
226	A DNA damage repair mechanism is involved in the origin of chromosomal translocations t(4;11) in primary leukemic cells. <i>Oncogene</i> , 1999, 18, 4663-4671.	5.9	106
227	Rapid isolation of chromosomal breakpoints from patients with t(4;11) acute lymphoblastic leukemia: implications for basic and clinical research. <i>Cancer Research</i> , 1999, 59, 3357-62.	0.9	31
228	Fine structure of translocation breakpoints in leukemic blasts with chromosomal translocation t(4;11): the DNA damage-repair model of translocation. <i>Oncogene</i> , 1998, 17, 3035-3044.	5.9	90
229	Formation of autocrine loops in human cerebral meningioma tissue by leukemia inhibitor factor, interleukin-6, and oncostatin M: inhibition of meningioma cell growth in vitro by recombinant oncostatin M. <i>Journal of Neurosurgery</i> , 1998, 88, 541-548.	1.6	36
230	The Structure of the Human ALL-1/MLL/HRX Gene. <i>Leukemia and Lymphoma</i> , 1997, 27, 417-428.	1.3	32
231	A novel, negative selectable marker for gene disruption in <i>Dictyostelium</i> . <i>Gene</i> , 1997, 202, 171-176.	2.2	6
232	Exon/intron structure of the human AF4 gene, a member of the AF4/LAF4/FMR2 gene family coding for a nuclear protein with structural alterations in acute leukaemia. <i>British Journal of Haematology</i> , 1997, 98, 157-169.	2.5	86
233	The human ALL-1/MLL/HRX antigen is predominantly localized in the nucleus of resting and proliferating peripheral blood mononuclear cells. <i>Cancer Research</i> , 1997, 57, 2035-41.	0.9	18
234	Exon/intron structure of the human ALL-1 (MLL) gene involved in translocations to chromosomal region 11q23 and acute leukaemias. <i>British Journal of Haematology</i> , 1996, 93, 966-972.	2.5	119

#	ARTICLE	IF	CITATIONS
235	A specific deletion in the breakpoint cluster region of the ALL-1 gene is associated with acute lymphoblastic T-cell leukemias. <i>Cancer Research</i> , 1996, 56, 2171-7.	0.9	41
236	Molecular analysis of the chromosomal breakpoint and fusion transcripts in the acute lymphoblastic SEM cell line with chromosomal translocation t(4;11). <i>British Journal of Haematology</i> , 1995, 90, 308-320.	2.5	21
237	The acute lymphoblastic leukaemia cell line SEM with t(4;11) chromosomal rearrangement is biphenotypic and responsive to interleukin-7. <i>British Journal of Haematology</i> , 1994, 86, 275-283.	2.5	106
238	Internally Located and Oppositely Oriented Polymerase II Promoters Direct Convergent Transcription of a LINE-Like Retroelement, the <i>Dictyostelium</i> Repetitive Element, from <i>Dictyostelium discoideum</i> . <i>Molecular and Cellular Biology</i> , 1994, 14, 3074-3084.	2.3	18
239	Characterization of transcripts from the <i>Dictyostelium discoideum</i> retrotransposable genetic element DRE. <i>Die Pharmazie</i> , 1994, 49, 923-5.	0.5	0
240	Structure of the promoter region of the gene encoding cytochrome c oxidase subunit V in <i>Dictyostelium</i> . <i>FEBS Journal</i> , 1993, 211, 411-414.	0.2	3
241	Different organization of the tRNA-gene-associated repetitive element, DRE, in NC4-derived strains and in other wild-type <i>Dictyostelium discoideum</i> strains. <i>FEBS Journal</i> , 1993, 217, 627-631.	0.2	11
242	Nucleotide Sequence of a <i>Dictyostelium discoideum</i> Gene Encoding a Protein Homologous to the Yeast Ribosomal Protein S31. <i>Biochemical and Biophysical Research Communications</i> , 1993, 190, 134-139.	2.1	8
243	The <i>Dictyostelium discoideum</i> 5S rDNA Is Organized in the Same Transcriptional Orientation as the Other rDNAs. <i>Biochemical and Biophysical Research Communications</i> , 1993, 191, 558-564.	2.1	13
244	Two distinct subforms of the retrotransposable DRE element in NC4 strains of <i>Dictyostelium discoideum</i> . <i>Nucleic Acids Research</i> , 1992, 20, 6247-6252.	14.5	14
245	Structure of DRE, a Retrotransposable Element Which Integrates with Position Specificity Upstream of <i>Dictyostelium discoideum</i> tRNA Genes. <i>Molecular and Cellular Biology</i> , 1992, 12, 229-239.	2.3	25
246	Establishment of a System for Conditional Gene Expression Using an Inducible tRNA Suppressor Gene. <i>Molecular and Cellular Biology</i> , 1992, 12, 4038-4045.	2.3	20
247	RNA polymerase III catalysed transcription can be regulated in <i>Saccharomyces cerevisiae</i> by the bacterial tetracycline repressor-operator system. <i>EMBO Journal</i> , 1992, 11, 1487-92.	7.8	35
248	Glycosylation sites identified by detection of glycosylated amino acids released from Edman degradation: The identification of Xaa-Pro-Xaa-Xaa as a motif for Thr-O-glycosylation. <i>Biochemical and Biophysical Research Communications</i> , 1991, 178, 1194-1201.	2.1	101
249	Transfer RNA genes from <i>Dictyostelium discoideum</i> are frequently associated with repetitive elements and contain consensus boxes in their 5' and 3'-flanking regions. <i>Journal of Molecular Biology</i> , 1991, 222, 537-552.	4.2	45
250	Nuclear factors which bind to <i>Dictyostelium discoideum</i> transfer RNA genes. <i>Current Genetics</i> , 1991, 20, 129-135.	1.7	4
251	Genomic organization of the transposable element Tdd-3 from <i>Dictyostelium discoideum</i> . <i>Nucleic Acids Research</i> , 1990, 18, 5751-5757.	14.5	15
252	Temperature sensitive synthesis of transfer RNAs in vivo in <i>Saccharomyces cerevisiae</i> . <i>EMBO Journal</i> , 1990, 9, 1253-8.	7.8	3

#	ARTICLE	IF	CITATIONS
253	Transfer RNA Genes: Landmarks for Integration of Mobile Genetic Elements in <i>Dictyostelium discoideum</i> . <i>Science</i> , 1989, 244, 1493-1496.	12.6	67
254	CMER, an RNA encoded by human cytomegalovirus is most likely transcribed by RNA polymerase III. <i>Nucleic Acids Research</i> , 1989, 17, 631-643.	14.5	9
255	tRNA ^{Glu} (GAA) Genes from the Cellular Slime Mold <i>Dictyostelium discoideum</i> . <i>DNA and Cell Biology</i> , 1989, 8, 193-204.	5.2	18
256	A family of non-allelic tRNAGUUVal genes from the cellular slime mold <i>Dictyostelium discoideum</i> . <i>Gene</i> , 1988, 73, 373-384.	2.2	21
257	Identification of a protein factor binding to the 5'-flanking region of a tRNA gene and being involved in modulation of tRNA gene transcription in <i>Saccharomyces cerevisiae</i> . <i>Nucleic Acids Research</i> , 1988, 16, 6737-6752.	14.5	36
258	Novel Diagnostic and Therapeutic Options for KMT2A-Rearranged Acute Leukemias. <i>Frontiers in Pharmacology</i> , 0, 13, .	3.5	6