

Jude Fitzgibbon

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6735277/publications.pdf>

Version: 2024-02-01

155
papers

8,020
citations

41258

49
h-index

53109

85
g-index

164
all docs

164
docs citations

164
times ranked

11452
citing authors

#	ARTICLE	IF	CITATIONS
1	Acquired somatic variants in inherited myeloid malignancies. <i>Leukemia</i> , 2022, 36, 1377-1381.	3.3	8
2	KDM4C in GC lymphoma: a new piece of the epigenetic puzzle. <i>Haematologica</i> , 2022, , .	1.7	1
3	CKS1 inhibition depletes leukemic stem cells and protects healthy hematopoietic stem cells in acute myeloid leukemia. <i>Science Translational Medicine</i> , 2022, 14, .	5.8	8
4	Molecular Genetics in Indolent Lymphomas. <i>Hematologic Malignancies</i> , 2021, , 5-20.	0.2	0
5	Germline ETV6 variants: not ALL created equally. <i>Blood</i> , 2021, 137, 288-289.	0.6	0
6	Generation and Surgical Analysis of to Study NF- κ B-Driven Pathogenesis of Diffuse Large B Cell Lymphoma. <i>Methods in Molecular Biology</i> , 2021, 2366, 321-342.	0.4	0
7	KDM5 inhibition offers a novel therapeutic strategy for the treatment of <i>KMT2D</i> mutant lymphomas. <i>Blood</i> , 2021, 138, 370-381.	0.6	33
8	Drug ranking using machine learning systematically predicts the efficacy of anti-cancer drugs. <i>Nature Communications</i> , 2021, 12, 1850.	5.8	68
9	The RUNX1 database (RUNX1db): establishment of an expert curated RUNX1 registry and genomics database as a public resource for familial platelet disorder with myeloid malignancy. <i>Haematologica</i> , 2021, 106, 3004-3007.	1.7	29
10	Genome-wide association study identifies susceptibility loci for acute myeloid leukemia. <i>Nature Communications</i> , 2021, 12, 6233.	5.8	17
11	Characterising Tumour and Microenvironmental Responses to R-CHOP in Immunocompetent Mouse Models of DLBCL. <i>Blood</i> , 2021, 138, 2401-2401.	0.6	0
12	The Paradoxical Efficacy of KDM6 Inhibition in Germinal Centre B-Cell Lymphomas. <i>Blood</i> , 2021, 138, 3289-3289.	0.6	0
13	An EZH2 Gene Expression Signature Is Predictive of Differential Efficacy of Chemotherapy Irrespective of EZH2 Mutation Status in Patients with Follicular Lymphoma Treated within the Gallium Trial. <i>Blood</i> , 2021, 138, 39-39.	0.6	1
14	Distinct genetic changes reveal evolutionary history and heterogeneous molecular grade of DLBCL with MYC/BCL2 double-hit. <i>Leukemia</i> , 2020, 34, 1329-1341.	3.3	66
15	AML through the prism of molecular genetics. <i>British Journal of Haematology</i> , 2020, 188, 49-62.	1.2	17
16	Identification of Recurrent Mutations in the microRNA-Binding Sites of B-Cell Lymphoma-Associated Genes in Follicular Lymphoma. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8795.	1.8	1
17	The Biological Basis of Histologic Transformation. <i>Hematology/Oncology Clinics of North America</i> , 2020, 34, 771-784.	0.9	4
18	Recent Advancements in Hematology: Knowledge, Methods and Dissemination, Part 1. <i>Hemato</i> , 2020, 1, 10-22.	0.2	0

#	ARTICLE	IF	CITATIONS
19	Recent Advancements in Hematology: Knowledge, Methods and Dissemination. Hemato, 2020, 1, 5-6.	0.2	0
20	A frameshift variant in specificity protein 1 triggers superactivation of Sp1-mediated transcription in familial bone marrow failure. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 17151-17155.	3.3	2
21	The complex genetic landscape of familial MDS and AML reveals pathogenic germline variants. Nature Communications, 2020, 11, 1044.	5.8	81
22	Genetic heterogeneity highlighted by differential FDG-PET response in diffuse large B-cell lymphoma. Haematologica, 2020, 105, 318-321.	1.7	5
23	Mesenchymal niche remodeling impairs hematopoiesis via stanniocalcin 1 in acute myeloid leukemia. Journal of Clinical Investigation, 2020, 130, 3038-3050.	3.9	48
24	Systematic Evaluation of Somatic <i>Cis</i> -Regulatory Mutations in Follicular Lymphoma. Blood, 2020, 136, 26-27.	0.6	0
25	Transmission of diffuse large B-cell lymphoma by an allogeneic stem-cell transplant. Haematologica, 2019, 104, e174-e177.	1.7	5
26	Oncogenic Rag GTPase signalling enhances B cell activation and drives follicular lymphoma sensitive to pharmacological inhibition of mTOR. Nature Metabolism, 2019, 1, 775-789.	5.1	40
27	Mutant CEBPA directly drives the expression of the targetable tumor-promoting factor CD73 in AML. Science Advances, 2019, 5, eaaw4304.	4.7	28
28	Follicular lymphoma. Nature Reviews Disease Primers, 2019, 5, 83.	18.1	148
29	Evaluation of the m7-FLIPI in Patients with Follicular Lymphoma Treated within the Gallium Trial: EZH2 mutation Status May be a Predictive Marker for Differential Efficacy of Chemotherapy. Blood, 2019, 134, 122-122.	0.6	41
30	Longitudinal Analyses of Diagnostic-Relapse Biopsies of Diffuse Large B Cell Lymphoma Reveal a Poor Risk Subset of ABC Patients Based on the Expression of a 30 Gene Panel. Blood, 2019, 134, 2769-2769.	0.6	0
31	Predicting early relapse in follicular lymphoma: have we turned a corner?. Lancet Oncology, The, 2018, 19, 441-442.	5.1	1
32	Proteomic and genomic integration identifies kinase and differentiation determinants of kinase inhibitor sensitivity in leukemia cells. Leukemia, 2018, 32, 1818-1822.	3.3	36
33	Genomic profiling reveals spatial intra-tumor heterogeneity in follicular lymphoma. Leukemia, 2018, 32, 1261-1265.	3.3	87
34	Rituximab as a first step in tackling transformation. Lancet Haematology, the, 2018, 5, e326-e327.	2.2	0
35	Precision medicine and lymphoma. Current Opinion in Hematology, 2018, 25, 329-334.	1.2	8
36	GATA2 monoallelic expression underlies reduced penetrance in inherited GATA2-mutated MDS/AML. Leukemia, 2018, 32, 2502-2507.	3.3	48

#	ARTICLE	IF	CITATIONS
37	Homozygous OB-fold variants in telomere protein TPP1 are associated with dyskeratosis congenita-like phenotypes. <i>Blood</i> , 2018, 132, 1349-1353.	0.6	16
38	Follicular lymphoma, a B cell malignancy addicted to epigenetic mutations. <i>Epigenetics</i> , 2017, 12, 370-377.	1.3	31
39	Familial CEBPA -mutated acute myeloid leukemia. <i>Seminars in Hematology</i> , 2017, 54, 87-93.	1.8	54
40	Recurrent somatic JAK-STAT pathway variants within a RUNX1-mutated pedigree. <i>European Journal of Human Genetics</i> , 2017, 25, 1020-1024.	1.4	13
41	Follicular lymphoma: State-of-the-art ICML workshop in Lugano 2015. <i>Hematological Oncology</i> , 2017, 35, 397-407.	0.8	11
42	Early loss of Crebbp confers malignant stem cell properties on lymphoid progenitors. <i>Nature Cell Biology</i> , 2017, 19, 1093-1104.	4.6	58
43	Proteomic and genomic integration identifies kinase and differentiation determinants of kinase inhibitor sensitivity in leukemia cells. <i>Leukemia</i> , 2017, , .	3.3	0
44	The molecular pathogenesis of the NUP98-HOXA9 fusion protein in acute myeloid leukemia. <i>Leukemia</i> , 2017, 31, 2000-2005.	3.3	28
45	The routes for transformation of follicular lymphoma. <i>Current Opinion in Hematology</i> , 2016, 23, 385-391.	1.2	16
46	Frequent NFKBIE deletions are associated with poor outcome in primary mediastinal B-cell lymphoma. <i>Blood</i> , 2016, 128, 2666-2670.	0.6	82
47	Germline heterozygous DDX41 variants in a subset of familial myelodysplasia and acute myeloid leukemia. <i>Leukemia</i> , 2016, 30, 2083-2086.	3.3	62
48	TNFRSF14 aberrations in follicular lymphoma increase clinically significant allogeneic T-cell responses. <i>Blood</i> , 2016, 128, 72-81.	0.6	29
49	Inherited DDX41 mutations: 11 genes and counting. <i>Blood</i> , 2016, 127, 960-961.	0.6	21
50	Pediatric-type FL: simply different. <i>Blood</i> , 2016, 128, 1030-1031.	0.6	2
51	Advances in the molecular diagnosis of diffuse large B-cell lymphoma in the era of precision medicine. <i>Expert Review of Molecular Diagnostics</i> , 2016, 16, 1093-1102.	1.5	3
52	The International Human Epigenome Consortium: A Blueprint for Scientific Collaboration and Discovery. <i>Cell</i> , 2016, 167, 1145-1149.	13.5	404
53	Clinical utility gene card for: Familial platelet disorder with associated myeloid malignancies. <i>European Journal of Human Genetics</i> , 2016, 24, 3-4.	1.4	5
54	5. Follicular lymphoma. , 2016, , 75-100.		0

#	ARTICLE	IF	CITATIONS
55	Reduced Expression of Histone Methyltransferases KMT2C and KMT2D Correlates with Improved Outcome in Pancreatic Ductal Adenocarcinoma. <i>Cancer Research</i> , 2016, 76, 4861-4871.	0.4	72
56	DNAJC21 Mutations Link a Cancer-Prone Bone Marrow Failure Syndrome to Corruption in 60S Ribosome Subunit Maturation. <i>American Journal of Human Genetics</i> , 2016, 99, 115-124.	2.6	85
57	Epigenetic dysregulation in follicular lymphoma. <i>Epigenomics</i> , 2016, 8, 77-84.	1.0	20
58	Recurrent mTORC1-activating RRAGC mutations in follicular lymphoma. <i>Nature Genetics</i> , 2016, 48, 183-188.	9.4	160
59	Early Loss of CREBBP Confers Malignant Stem Cell Properties on Lymphoid Progenitors. <i>Blood</i> , 2016, 128, 460-460.	0.6	1
60	IRF1 Deletions: A Novel Marker of Clinical Aggressiveness in Primary Mediastinal B-Cell Lymphoma. <i>Blood</i> , 2016, 128, 609-609.	0.6	0
61	Disease evolution and outcomes in familial AML with germline CEBPA mutations. <i>Blood</i> , 2015, 126, 1214-1223.	0.6	157
62	High Throughput Sequencing Analysis of the Immunoglobulin Heavy Chain Gene from Flow-Sorted B Cell Sub-Populations Define the Dynamics of Follicular Lymphoma Clonal Evolution. <i>PLoS ONE</i> , 2015, 10, e0134833.	1.1	28
63	A multi-gene signature predicts outcome in patients with pancreatic ductal adenocarcinoma. <i>Genome Medicine</i> , 2014, 6, 105.	3.6	106
64	Myeloblastic therapy with autologous haematopoietic stem cell support as consolidation of first remission in acute myeloid leukaemia – “very long follow-up”. <i>British Journal of Haematology</i> , 2014, 167, 724-726.	1.2	4
65	Integrated genomic analysis identifies recurrent mutations and evolution patterns driving the initiation and progression of follicular lymphoma. <i>Nature Genetics</i> , 2014, 46, 176-181.	9.4	624
66	Investigational epigenetically targeted drugs in early phase trials for the treatment of haematological malignancies. <i>Expert Opinion on Investigational Drugs</i> , 2014, 23, 1321-1332.	1.9	7
67	Genome-wide copy-number analyses reveal genomic abnormalities involved in transformation of follicular lymphoma. <i>Blood</i> , 2014, 123, 1681-1690.	0.6	110
68	GATA2 mutations in sporadic and familial acute myeloid leukaemia patients with CEBPA mutations. <i>British Journal of Haematology</i> , 2013, 161, 701-705.	1.2	47
69	Highly variable clinical manifestations in a large family with a novel GATA2 mutation. <i>Leukemia</i> , 2013, 27, 2247-2248.	3.3	27
70	EZH2 mutations are frequent and represent an early event in follicular lymphoma. <i>Blood</i> , 2013, 122, 3165-3168.	0.6	274
71	The co-receptor BTLA negatively regulates human V α 9V β 2 T-cell proliferation: a potential way of immune escape for lymphoma cells. <i>Blood</i> , 2013, 122, 922-931.	0.6	87
72	Functional Analysis of a Breast Cancer-Associated FGFR2 Single Nucleotide Polymorphism Using Zinc Finger Mediated Genome Editing. <i>PLoS ONE</i> , 2013, 8, e78839.	1.1	14

#	ARTICLE	IF	CITATIONS
73	Investigating The Role Of MLL2 (Mll4) In B Cell Development. Blood, 2013, 122, 343-343.	0.6	0
74	Familial AML With Germline CEBPA Mutations: Extended Clinical Outcomes and Analysis Of Secondary Mutations Using Whole Exome Sequencing. Blood, 2013, 122, 740-740.	0.6	0
75	Germ-line GATA2 p.THR354MET mutation in familial myelodysplastic syndrome with acquired monosomy 7 and ASXL1 mutation demonstrating rapid onset and poor survival. Haematologica, 2012, 97, 890-894.	1.7	85
76	Promoter methylation of argininosuccinate synthetase-1 sensitises lymphomas to arginine deiminase treatment, autophagy and caspase-dependent apoptosis. Cell Death and Disease, 2012, 3, e342-e342.	2.7	107
77	Overexpression of wild-type or mutants forms of CEBPA alter normal human hematopoiesis. Leukemia, 2012, 26, 1537-1546.	3.3	23
78	BLUEPRINT to decode the epigenetic signature written in blood. Nature Biotechnology, 2012, 30, 224-226.	9.4	323
79	High Incidence of EZH2 Mutations with Variable Mutation Load in Follicular Lymphoma and Its Consequences for EZH2 Targeted Therapy. Blood, 2012, 120, 545-545.	0.6	0
80	Whole Genome Sequencing in Sequential Biopsies Reveals the Genetic Evolution of Follicular Lymphoma to Transformed Follicular Lymphoma. Blood, 2012, 120, 145-145.	0.6	0
81	SNP rs6457327 in the HLA region on chromosome 6p is predictive of the transformation of follicular lymphoma. Blood, 2011, 117, 3147-3150.	0.6	17
82	MicroRNA profiles of t(14;18)â€“negative follicular lymphoma support a late germinal center B-cell phenotype. Blood, 2011, 118, 5550-5558.	0.6	77
83	It's a targeted world in non-Hodgkin's lymphoma. British Journal of Nursing, 2011, 20, S28-S28.	0.3	1
84	Serum selenium concentration at diagnosis and outcome in patients with haematological malignancies. British Journal of Haematology, 2011, 154, 448-456.	1.2	20
85	EZH2 Y641 mutations in follicular lymphoma. Leukemia, 2011, 25, 726-729.	3.3	132
86	Transformation of Indolent B-Cell Lymphomas. Journal of Clinical Oncology, 2011, 29, 1827-1834.	0.8	164
87	The clinical relevance of <i>WT1</i> gene mutations in acute leukaemia. Hematological Oncology, 2010, 28, 13-19.	0.8	38
88	Molecular signatures in the diagnosis and management of follicular lymphoma. Current Opinion in Hematology, 2010, 17, 333-340.	1.2	8
89	A novel <i>RUNX1</i> mutation in a kindred with familial platelet disorder with propensity to acute myeloid leukaemia: male predominance of affected individuals. European Journal of Haematology, 2010, 85, 552-553.	1.1	13
90	AZD1152 Rapidly and Negatively Affects the Growth and Survival of Human Acute Myeloid Leukemia Cells <i>In vitro</i> and <i>In vivo</i> . Cancer Research, 2009, 69, 4150-4158.	0.4	72

#	ARTICLE	IF	CITATIONS
91	Array-based DNA methylation profiling in follicular lymphoma. <i>Leukemia</i> , 2009, 23, 1858-1866.	3.3	65
92	Follicular lymphomas with and without translocation t(14;18) differ in gene expression profiles and genetic alterations. <i>Blood</i> , 2009, 114, 826-834.	0.6	177
93	Clinical relevance of MDM2 SNP 309 and TP53 Arg72Pro in follicular lymphoma. <i>Haematologica</i> , 2009, 94, 148-150.	1.7	18
94	Transformation of follicular lymphoma to diffuse large B-cell lymphoma may occur by divergent evolution from a common progenitor cell or by direct evolution from the follicular lymphoma clone. <i>Blood</i> , 2009, 113, 3553-3557.	0.6	129
95	Regions of acquired uniparental disomy at diagnosis of follicular lymphoma are associated with both overall survival and risk of transformation. <i>Blood</i> , 2009, 113, 2298-2301.	0.6	75
96	Familial myelodysplasia and acute myeloid leukaemia – a review. <i>British Journal of Haematology</i> , 2008, 140, 123-132.	1.2	159
97	Mutation of the Wilms' Tumor 1 Gene Is a Poor Prognostic Factor Associated With Chemotherapy Resistance in Normal Karyotype Acute Myeloid Leukemia: The United Kingdom Medical Research Council Adult Leukaemia Working Party. <i>Journal of Clinical Oncology</i> , 2008, 26, 5429-5435.	0.8	185
98	Segmental uniparental disomy is a commonly acquired genetic event in relapsed acute myeloid leukemia. <i>Blood</i> , 2008, 112, 814-821.	0.6	97
99	The presence of TP53 mutation at diagnosis of follicular lymphoma identifies a high-risk group of patients with shortened time to disease progression and poorer overall survival. <i>Blood</i> , 2008, 112, 3126-3129.	0.6	112
100	Five new pedigrees with inherited RUNX1 mutations causing familial platelet disorder with propensity to myeloid malignancy. <i>Blood</i> , 2008, 112, 4639-4645.	0.6	222
101	WTX is rarely mutated in acute myeloid leukemia. <i>Haematologica</i> , 2008, 93, 947-948.	1.7	10
102	Five New Pedigrees with Inherited RUNX1 Mutations Causing Familial Platelet Disorder with Propensity to Myeloid Malignancy (FPD/AML). <i>Blood</i> , 2008, 112, 5067-5067.	0.6	0
103	Recurrent Chromosomal Intermingling Interactions at the BCL2 Locus in T(14;18) +Ve and -Ve Cell Lines. <i>Blood</i> , 2008, 112, 2048-2048.	0.6	0
104	Genetic Abnormalities Involved in the Development and Progression of Follicular Lymphoma. <i>Blood</i> , 2008, 112, 2049-2049.	0.6	0
105	Role of Different C/EBP β Mutations in AML Transformation. <i>Blood</i> , 2008, 112, 1343-1343.	0.6	0
106	Chemosensitization of B-Cell Lymphomas by Methylseleninic Acid Involves Nuclear Factor- κ B Inhibition and the Rapid Generation of Other Selenium Species. <i>Cancer Research</i> , 2007, 67, 10984-10992.	0.4	41
107	Wilms' tumour 1 mutations are associated with FLT3-ITD and failure of standard induction chemotherapy in patients with normal karyotype AML. <i>Leukemia</i> , 2007, 21, 550-551.	3.3	88
108	Genome-wide detection of recurring sites of uniparental disomy in follicular and transformed follicular lymphoma. <i>Leukemia</i> , 2007, 21, 1514-1520.	3.3	78

#	ARTICLE	IF	CITATIONS
109	Transformation of follicular lymphoma to diffuse large B-cell lymphoma proceeds by distinct oncogenic mechanisms. <i>British Journal of Haematology</i> , 2007, 136, 286-293.	1.2	142
110	Should Patients with Myelodysplasia and/or Acute Myeloid Leukemia (MDS/AML) Be Screened for Inherited RUNX1 Mutations Prior to Sibling Donor Allografting?.. <i>Blood</i> , 2007, 110, 5027-5027.	0.6	0
111	Development of a quantitative real-time polymerase chain reaction method for monitoring CEBPA mutations in normal karyotype acute myeloid leukaemia. <i>British Journal of Haematology</i> , 2006, 133, 060203080756008.	1.2	16
112	A novel K509I mutation of KIT identified in familial mastocytosisâ€”in vitro and in vivo responsiveness to imatinib therapy. <i>Leukemia Research</i> , 2006, 30, 373-378.	0.4	129
113	The activity of methylated and non-methylated selenium species in lymphoma cell lines and primary tumours. <i>Annals of Oncology</i> , 2006, 17, 773-779.	0.6	35
114	Wilmâ€™s Tumour 1 (WT1) Mutations Are Associated with FLT3-ITD Mutation and Poor Prognosis in Normal Karyotype AML.. <i>Blood</i> , 2006, 108, 2302-2302.	0.6	0
115	aUPD in the Clonal Evolution of Follicular Lymphoma.. <i>Blood</i> , 2006, 108, 2065-2065.	0.6	0
116	Development of a human acute myeloid leukaemia screening panel and consequent identification of novel gene mutation in FLT3 and CCND3. <i>British Journal of Haematology</i> , 2005, 128, 318-323.	1.2	25
117	A limited role for TP53 mutation in the transformation of follicular lymphoma to diffuse large B-cell lymphoma. <i>Leukemia</i> , 2005, 19, 1459-1465.	3.3	58
118	Mutations of the PU.1 Ets domain are specifically associated with murine radiation-induced, but not human therapy-related, acute myeloid leukaemia. <i>Oncogene</i> , 2005, 24, 3678-3683.	2.6	58
119	Association between Acquired Uniparental Disomy and Homozygous Gene Mutation in Acute Myeloid Leukemias. <i>Cancer Research</i> , 2005, 65, 9152-9154.	0.4	221
120	The Activity of the Novel Aurora Kinase B Inhibitor AZD1152 in Acute Myeloid Leukaemia Cells.. <i>Blood</i> , 2005, 106, 3374-3374.	0.6	12
121	Methylseleninic Acid Results in Rapid Changes in Intracellular Selenium Species and Sensitises Human Lymphoma Cells to Doxorubicin.. <i>Blood</i> , 2005, 106, 1768-1768.	0.6	0
122	Reliable detection of clonal IgH/Bcl2 MBR rearrangement in follicular lymphoma: methodology and clinical significance. <i>British Journal of Haematology</i> , 2004, 124, 325-328.	1.2	16
123	Mutation ofCEBPAin Familial Acute Myeloid Leukemia. <i>New England Journal of Medicine</i> , 2004, 351, 2403-2407.	13.9	295
124	Tissue Microarray Is a Useful Tool in the Evaluation of Genes Implicated in Transformation of Follicular Lymphoma.. <i>Blood</i> , 2004, 104, 2267-2267.	0.6	1
125	Sequential Time-Point Mutational Analysis of TP53 in Follicular Lymphoma Undergoing Transformation to Large B-Cell Lymphoma.. <i>Blood</i> , 2004, 104, 1361-1361.	0.6	12
126	The Induction of Apoptosis in Lymphoma Cells by Methylseleninic Acid and Selenodiglutathione.. <i>Blood</i> , 2004, 104, 2502-2502.	0.6	0

#	ARTICLE	IF	CITATIONS
127	Mutation of CEBPA in Familial Acute Myeloid Leukaemia.. Blood, 2004, 104, 2012-2012.	0.6	0
128	Development of a Human Acute Myeloid Leukaemia Screening Panel and Identification of Novel Gene Mutations.. Blood, 2004, 104, 2991-2991.	0.6	0
129	Flow Cytometric FLT3 Expression in Acute Leukaemias Is of Diagnostic Value but Does Not Correlate with ITD / D835Y Mutation Status.. Blood, 2004, 104, 3014-3014.	0.6	0
130	Mutations of CEBPA in acute myeloid leukemia FAB types M1 and M2. Genes Chromosomes and Cancer, 2003, 37, 72-78.	1.5	98
131	Mutation of BRAF is uncommon in AML FAB type M1 and M2. Leukemia, 2003, 17, 274-275.	3.3	23
132	Presentation Serum Selenium Predicts for Overall Survival, Dose Delivery, and First Treatment Response in Aggressive Non-Hodgkin's Lymphoma. Journal of Clinical Oncology, 2003, 21, 2335-2341.	0.8	56
133	Mutations in the runt homology domain of CBF β 2 in myeloid malignancies with acquired trisomy 21. Cancer Genetics and Cytogenetics, 2002, 136, 151-152.	1.0	6
134	JH probe real-time quantitative polymerase chain reaction assay for Bcl-2/IgH rearrangements. British Journal of Haematology, 2002, 118, 550-558.	1.2	12
135	The relative role of peripheral blood and bone marrow for monitoring molecular evidence of disease in follicular lymphoma by quantitative real-time polymerase chain reaction. British Journal of Haematology, 2002, 118, 563-566.	1.2	15
136	Frequency of the <i>Bcl-2/IgH</i> Rearrangement in Normal Individuals: Implications for the Monitoring of Disease in Patients With Follicular Lymphoma. Journal of Clinical Oncology, 2001, 19, 420-424.	0.8	137
137	Localisation of a novel region of recurrent amplification in follicular lymphoma to an ~6.8 Mb region of 13q32-33. Genes Chromosomes and Cancer, 2001, 32, 236-243.	1.5	40
138	Familial follicular lymphoma: A Case Report with Molecular Analysis. British Journal of Haematology, 2000, 110, 744-745.	1.2	2
139	Tumour necrosis factor polymorphisms and susceptibility to follicular lymphoma. British Journal of Haematology, 1999, 107, 388-391.	1.2	25
140	Isolation and Chromosomal Localization of Two Human CDP-diacylglycerol Synthase (CDS) Genes. Genomics, 1998, 54, 140-144.	1.3	44
141	Molecular Evolution of the Cottoid Fish Endemic to Lake Baikal Deduced from Nuclear DNA Evidence. Molecular Phylogenetics and Evolution, 1997, 8, 415-422.	1.2	39
142	Spectral tuning and molecular evolution of rod visual pigments in the species flock of cottoid fish in Lake Baikal. Vision Research, 1996, 36, 1217-1224.	0.7	129
143	Human guanylate kinase (GUK1): cDNA sequence, expression and chromosomal localisation. FEBS Letters, 1996, 385, 185-188.	1.3	17
144	Paralogy Mapping: Identification of a Region in the Human MHC Triplicated onto Human Chromosomes 1 and 9 Allows the Prediction and Isolation of Novel PBX and NOTCH Loci. Genomics, 1996, 35, 101-108.	1.3	161

#	ARTICLE	IF	CITATIONS
145	Localisation of the gene encoding diacylglycerol kinase 3 (DAGK3) to human chromosome 3q27-28 and mouse chromosome 16. <i>Current Eye Research</i> , 1995, 14, 1041-1043.	0.7	1
146	The rhodopsin-encoding gene of bony fish lacks introns. <i>Gene</i> , 1995, 164, 273-277.	1.0	77
147	Chromosomal localization of three mouse diacylglycerol kinase (DAGK) genes: genes sharing sequence homology to the <i>Drosophila</i> retinal degeneration A (rdgA) gene. <i>Genomics</i> , 1995, 26, 599-601.	1.3	12
148	Regional Localization of 64 Cosmid Contigs, Including 18 Genes and 14 Markers, to Intervals on Human Chromosome 9q34. <i>Genomics</i> , 1995, 29, 257-260.	1.3	5
149	<i>Drosophila</i> Visual Transduction, a Model System For Human Eye Disease?. , 1995, , 255-261.		1
150	Cosmid Contigs Spanning 9q34 Including the Candidate Region forTSCI. <i>European Journal of Human Genetics</i> , 1995, 3, 65-77.	1.4	9
151	Localization of the gene encoding human phosphatidylinositol transfer protein (PITPN) to 17p13.3: a gene showing homology to the <i>Drosophila</i> retinal degeneration B gene (rdgB). <i>Cytogenetic and Genome Research</i> , 1994, 67, 205-207.	0.6	8
152	Localisation of the human blue cone pigment gene to chromosome band 7q31.3-32. <i>Human Genetics</i> , 1994, 93, 79-80.	1.8	18
153	Localization of the retinoid X receptor alpha gene (RXRA) to chromosome 9q34. <i>Annals of Human Genetics</i> , 1993, 57, 195-201.	0.3	7
154	Mapping of RXRB to human chromosome 6p21. 3. <i>Annals of Human Genetics</i> , 1993, 57, 203-209.	0.3	15
155	A dual role for the RNA helicase DHX34 in NMD and pre-mRNA splicing and its function in hematopoietic differentiation. <i>Rna</i> , 0, , rna.079277.122.	1.6	4