Nicholas Cross

List of Publications by Year
in descending order

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286 papers	32,163 citations	4658 85 h-index	4645 170 g-index
287 all docs	287 docs citations	287 times ranked	21264 citing authors

#	Article	IF	Citations
1	Clonal myelopoiesis promotes adverse outcomes in chronic kidney disease. Leukemia, 2022, 36, 507-515.	7.2	49
2	Chronic Eosinophilic Leukaemia Associated with <i>JAK2</i> Exon 13 Insertion/Deletion Mutations. Acta Haematologica, 2022, 145, 201-206.	1.4	2
3	Superior Efficacy of Midostaurin Over Cladribine in Advanced Systemic Mastocytosis: A Registry-Based Analysis. Journal of Clinical Oncology, 2022, 40, 1783-1794.	1.6	24
4	Standardization of molecular monitoring of CML: results and recommendations from the European treatment and outcome study. Leukemia, 2022, 36, 1834-1842.	7.2	10
5	Genome-wide association study identifies novel susceptibility loci for KIT D816V positive mastocytosis. American Journal of Human Genetics, 2021, 108, 284-294.	6.2	12
6	Adverse Prognostic Impact of the KIT D816V Transcriptional Activity in Advanced Systemic Mastocytosis. International Journal of Molecular Sciences, 2021, 22, 2562.	4.1	9
7	Allogeneic haematopoietic cell transplantation for myelofibrosis: proposed definitions and management strategies for graft failure, poor graft function and relapse: best practice recommendations of the EBMT Chronic Malignancies Working Party. Leukemia, 2021, 35, 2445-2459.	7.2	36
8	The use of genetic tests to diagnose and manage patients with myeloproliferative and myeloproliferative/myelodysplastic neoplasms, and related disorders. British Journal of Haematology, 2021, 195, 338-351.	2.5	8
9	Realâ€world tyrosine kinase inhibitor treatment pathways, monitoring patterns and responses in patients with chronic myeloid leukaemia in the United Kingdom: the UK TARGET CML study. British Journal of Haematology, 2021, 192, 62-74.	2.5	18
10	HGNC nomenclature for fusion genes. Leukemia, 2021, 35, 3039-3039.	7.2	1
11	HUGO Gene Nomenclature Committee (HGNC) recommendations for the designation of gene fusions. Leukemia, 2021, 35, 3040-3043.	7.2	42
12	A British Society for Haematology Guideline on the diagnosis and management of chronic myeloid leukaemia. British Journal of Haematology, 2020, 191, 171-193.	2.5	38
13	Analysis of chronic myeloid leukaemia during deep molecular response by genomic PCR: a traffic light stratification model with impact on treatment-free remission. Leukemia, 2020, 34, 2113-2124.	7.2	22
14	Clonal myelopoiesis in the UK Biobank cohort: ASXL1 mutations are strongly associated with smoking. Leukemia, 2020, 34, 2660-2672.	7.2	96
15	Is cancer latency an outdated concept? Lessons from chronic myeloid leukemia. Leukemia, 2020, 34, 2279-2284.	7.2	7
16	Mutational mechanisms of EZH2 inactivation in myeloid neoplasms. Leukemia, 2020, 34, 3206-3214.	7.2	8
17	A Novel t(1;9)(p36;p24.1) JAK2 Translocation and Review of the Literature. Acta Haematologica, 2019, 142, 105-112.	1.4	5
18	Inhibitory effects of midostaurin and avapritinib on myeloid progenitors derived from patients with KIT D816V positive advanced systemic mastocytosis. Leukemia, 2019, 33, 1195-1205.	7.2	38

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19	Recurrent activating STAT5B N642H mutation in myeloid neoplasms with eosinophilia. Leukemia, 2019, 33, 415-425.	7.2	65
20	PRR14L mutations are associated with chromosome 22 acquired uniparental disomy, age-related clonal hematopoiesis and myeloid neoplasia. Leukemia, 2019, 33, 1184-1194.	7.2	11
21	KIT D816 mutated/CBF-negative acute myeloid leukemia: a poor-risk subtype associated with systemic mastocytosis. Leukemia, 2019, 33, 1124-1134.	7.2	29
22	Routine Screening for <i>KIT</i> M541L Is Not Warranted in the Diagnostic Work-Up of Patients with Hypereosinophilia. Acta Haematologica, 2018, 139, 71-73.	1.4	5
23	Consensus on <scp>BCR</scp>â€<scp>ABL</scp>1 reporting in chronic myeloid leukaemia in the <scp>UK</scp>. British Journal of Haematology, 2018, 182, 777-788.	2.5	11
24	Absence of <i>CALR</i> Mutations in Idiopathic Erythrocytosis Patients with Low Serum Erythropoietin Levels. Acta Haematologica, 2018, 139, 217-219.	1.4	3
25	Guideline for the investigation and management of eosinophilia. British Journal of Haematology, 2017, 176, 553-572.	2.5	110
26	Nilotinib first-line therapy in patients with Philadelphia chromosome-negative/BCR-ABL-positive chronic myeloid leukemia in chronic phase: ENEST1st sub-analysis. Journal of Cancer Research and Clinical Oncology, 2017, 143, 1225-1233.	2.5	9
27	The clinical and molecular diversity of mast cell leukemia with or without associated hematologic neoplasm. Haematologica, 2017, 102, 1035-1043.	3.5	84
28	Response and progression on midostaurin in advanced systemic mastocytosis: KIT D816V and other molecular markers. Blood, 2017, 130, 137-145.	1.4	97
29	The effect of initial molecular profile on response to recombinant interferonâ€Î± (rIFNÎ±) treatment in early myelofibrosis. Cancer, 2017, 123, 2680-2687.	4.1	48
30	Impact of age on efficacy and toxicity of nilotinib in patients with chronic myeloid leukemia in chronic phase: ENEST1st subanalysis. Journal of Cancer Research and Clinical Oncology, 2017, 143, 1585-1596.	2.5	29
31	Ruxolitinib, a potent JAK1/JAK2 inhibitor, induces temporary reductions in the allelic burden of concurrent CSF3R mutations in chronic neutrophilic leukemia. Haematologica, 2017, 102, e238-e240.	3.5	38
32	Measurement ofBCR-ABL1by RT-qPCR in chronic myeloid leukaemia: findings from an International EQA Programme. British Journal of Haematology, 2017, 177, 414-422.	2.5	16
33	A Novel PCM1-PDGFRB Fusion in a Patient with a Chronic Myeloproliferative Neoplasm and an ins(8;5). Acta Haematologica, 2017, 138, 198-200.	1.4	5
34	Genomics of Myeloproliferative Neoplasms. Journal of Clinical Oncology, 2017, 35, 947-954.	1.6	62
35	The U2AF1S34F mutation induces lineage-specific splicing alterations in myelodysplastic syndromes. Journal of Clinical Investigation, 2017, 127, 2206-2221.	8.2	69
36	Exon-centric regulation of ATM expression is population-dependent and amenable to antisense modification by pseudoexon targeting. Scientific Reports, 2016, 6, 18741.	3.3	5

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37	Development and evaluation of a secondary reference panel for BCR-ABL1 quantification on the International Scale. Leukemia, 2016, 30, 1844-1852.	7.2	51
38	Antisense Oligonucleotides Modulating Activation of a Nonsense-Mediated RNA Decay Switch Exon in the ATM Gene. Nucleic Acid Therapeutics, 2016, 26, 392-400.	3.6	9
39	Standardization of molecular monitoring for chronic myeloid leukemia in Latin America using locally produced secondary cellular calibrators. Leukemia, 2016, 30, 2258-2260.	7.2	12
40	Impact of centralized evaluation of bone marrow histology in systemic mastocytosis. European Journal of Clinical Investigation, 2016, 46, 392-397.	3.4	21
41	Splenomegaly, elevated alkaline phosphatase and mutations in the SRSF2/ASXL1/RUNX1 gene panel are strong adverse prognostic markers in patients with systemic mastocytosis. Leukemia, 2016, 30, 2342-2350.	7.2	73
42	Diagnostic challenges in the work up of hypereosinophilia: pitfalls in bone marrow core biopsy interpretation. Annals of Hematology, 2016, 95, 557-562.	1.8	27
43	Additional mutations in SRSF2, ASXL1 and/or RUNX1 identify a high-risk group of patients with KIT D816V+ advanced systemic mastocytosis. Leukemia, 2016, 30, 136-143.	7.2	185
44	Frontline nilotinib in patients with chronic myeloid leukemia in chronic phase: results from the European ENEST1st study. Leukemia, 2016, 30, 57-64.	7.2	91
45	An international consortium proposal of uniform response criteria for myelodysplastic/myeloproliferative neoplasms (MDS/MPN) in adults. Blood, 2015, 125, 1857-1865.	1.4	153
46	Neutrophilic leukemoid reaction in multiple myeloma. American Journal of Hematology, 2015, 90, 1090-1090.	4.1	3
47	<scp>KIT</scp><scp>D</scp>816<scp>V</scp> and <scp>JAK</scp>2<scp>V</scp>617<scp>F</scp> mutations are seen recurrently in hypereosinophilia of unknown significance. American Journal of Hematology, 2015, 90, 774-777.	4.1	50
48	Fusion of PDGFRB to MPRIP, CPSF6, and GOLGB1 in three patients with eosinophiliaâ€associated myeloproliferative neoplasms. Genes Chromosomes and Cancer, 2015, 54, 762-770.	2.8	27
49	Identification of U2AF(35)-dependent exons by RNA-Seq reveals a link between 3â€² splice-site organization and activity of U2AF-related proteins. Nucleic Acids Research, 2015, 43, 3747-3763.	14.5	35
50	Limited duration of complete remission on ruxolitinib in myeloid neoplasms with PCM1-JAK2 and BCR-JAK2 fusion genes. Annals of Hematology, 2015, 94, 233-238.	1.8	74
51	Laboratory recommendations for scoring deep molecular responses following treatment for chronic myeloid leukemia. Leukemia, 2015, 29, 999-1003.	7.2	280
52	Combining gene mutation with gene expression data improves outcome prediction in myelodysplastic syndromes. Nature Communications, 2015, 6, 5901.	12.8	196
53	Detection of leukemia-associated mutations in peripheral blood DNA of hematologically normal elderly individuals. Leukemia, 2015, 29, 1600-1602.	7.2	16
54	Molecular profiling of myeloid progenitor cells in multi-mutated advanced systemic mastocytosis identifies KIT D816V as a distinct and late event. Leukemia, 2015, 29, 1115-1122.	7.2	134

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55	Low frequency mutations independently predict poor treatment-free survival in early stage chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. Haematologica, 2015, 100, e237-e239.	3.5	21
56	KIT mutation analysis in mast cell neoplasms: recommendations of the European Competence Network on Mastocytosis. Leukemia, 2015, 29, 1223-1232.	7.2	229
57	An International MDS/MPN Working Group's perspective and recommendations on molecular pathogenesis, diagnosis and clinical characterization of myelodysplastic/myeloproliferative neoplasms. Haematologica, 2015, 100, 1117-1130.	3.5	97
58	Profound parental bias associated with chromosome 14 acquired uniparental disomy indicates targeting of an imprinted locus. Leukemia, 2015, 29, 2069-2074.	7.2	13
59	Genetic variation at MECOM, TERT, JAK2 and HBS1L-MYB predisposes to myeloproliferative neoplasms. Nature Communications, 2015, 6, 6691.	12.8	145
60	Evaluation of methods to detect CALR mutations in myeloproliferative neoplasms. Leukemia Research, 2015, 39, 82-87.	0.8	55
61	A certified plasmid reference material for the standardisation of BCRâ€“ABL1 mRNA quantification by real-time quantitative PCR. Leukemia, 2015, 29, 369-376.	7.2	72
62	Molecular pathogenesis of atypical CML, CMML and MDS/MPN-unclassifiable. International Journal of Hematology, 2015, 101, 229-242.	1.6	56
63	A multi-centre phase 2 study of azacitidine in chronic myelomonocytic leukaemia. Leukemia, 2014, 28, 1570-1572.	7.2	34
64	Identification and functional characterization of imatinibâ€sensitive DTD1â€PDGFRB and CCDC88Câ€PDGFRB fusion genes in eosinophiliaâ€associated myeloid/lymphoid neoplasms. Genes Chromosomes and Cancer, 2014, 53, 411-421.	2.8	23
65	Patients with myeloid malignancies bearing PDGFRB fusion genes achieve durable long-term remissions with imatinib. Blood, 2014, 123, 3574-3577.	1.4	118
66	Mosaic loss of chromosome Y in peripheral blood is associated with shorter survival and higher risk of cancer. Nature Genetics, 2014, 46, 624-628.	21.4	320
67	The number of prognostically detrimental mutations and prognosis in primary myelofibrosis: an international study of 797 patients. Leukemia, 2014, 28, 1804-1810.	7.2	263
68	The KIT D816V expressed allele burden for diagnosis and disease monitoring of systemic mastocytosis. Annals of Hematology, 2014, 93, 81-88.	1.8	142
69	Modification of British Committee for Standards in Haematology diagnostic criteria for essential thrombocythaemia. British Journal of Haematology, 2014, 167, 421-423.	2.5	40
70	The future of JAK inhibition in myelofibrosis and beyond. Blood Reviews, 2014, 28, 189-196.	5.7	25
71	Bioinformatic analyses of CALR mutations in myeloproliferative neoplasms support a role in signaling. Leukemia, 2014, 28, 2106-2109.	7.2	20
72	Megalencephaly Syndromes: Exome Pipeline Strategies for Detecting Low-Level Mosaic Mutations. PLoS ONE, 2014, 9, e86940.	2.5	20

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73	Mutations in <scp>SETBP</scp>1 are recurrent in myelodysplastic syndromes and often coexist with cytogenetic markers associated with disease progression. British Journal of Haematology, 2013, 163, 235-239.	2.5	37
74	Distribution of genomic breakpoints in chronic myeloid leukemia: analysis of 308 patients. Leukemia, 2013, 27, 2105-2107.	7.2	23
75	Establishing optimal quantitative-polymerase chain reaction assays for routine diagnosis and tracking of minimal residual disease in JAK2-V617F-associated myeloproliferative neoplasms: a joint European LeukemiaNet/MPN&MPNr-EuroNet (COST action BM0902) study. Leukemia, 2013, 27, 2032-2039.	7.2	96
76	SomaticCALRMutations in Myeloproliferative Neoplasms with NonmutatedJAK2. New England Journal of Medicine, 2013, 369, 2391-2405.	27.0	1,556
77	JAK2V617F allele burden, JAK2 46/1 haplotype and clinical features of Chinese with myeloproliferative neoplasms. Leukemia, 2013, 27, 1763-1767.	7.2	24
78	Clinical and biological implications of driver mutations in myelodysplastic syndromes. Blood, 2013, 122, 3616-3627.	1.4	1,562
79	The clinical significance of NOTCH1 and SF3B1 mutations in the UK LRF CLL4 trial. Blood, 2013, 121, 468-475.	1.4	190
80	Recurrent SETBP1 mutations in atypical chronic myeloid leukemia. Nature Genetics, 2013, 45, 18-24.	21.4	359
81	Recurrent CEP85Lâ€“PDGFRB fusion in patient with t(5;6) and imatinib-responsive myeloproliferative neoplasm with eosinophilia. Leukemia and Lymphoma, 2013, 54, 1527-1531.	1.3	10
82	Prognostic Score Including Gene Mutations in Chronic Myelomonocytic Leukemia. Journal of Clinical Oncology, 2013, 31, 2428-2436.	1.6	462
83	Comprehensive mutational profiling in advanced systemic mastocytosis. Blood, 2013, 122, 2460-2466.	1.4	222
84	Molecular diagnosis of the myeloproliferative neoplasms: <scp>UK</scp> guidelines for the detection of <scp>JAK</scp>2 <scp>V</scp>617<scp>F</scp> and other relevant mutations. British Journal of Haematology, 2013, 160, 25-34.	2.5	87
85	Establishment and Validation of Analytical Reference Panels for the Standardization of Quantitative BCR-ABL1 Measurements on the International Scale. Clinical Chemistry, 2013, 59, 938-948.	3.2	46
86	Why do we see JAK2 exon 12 mutations in myeloproliferative neoplasms?. Leukemia, 2013, 27, 1930-1932.	7.2	4
87	Mutations and prognosis in primary myelofibrosis. Leukemia, 2013, 27, 1861-1869.	7.2	653
88	Ponatinib as targeted therapy for FGFR1 fusions associated with the 8p11 myeloproliferative syndrome. Haematologica, 2013, 98, 103-106.	3.5	70
89	Long-term follow-up of treatment with imatinib in eosinophilia-associated myeloid/lymphoid neoplasms with PDGFR rearrangements in blast phase. Leukemia, 2013, 27, 2254-2256.	7.2	61
90	Ruxolitinib as potential targeted therapy for patients with JAK2 rearrangements. Haematologica, 2013, 98, 404-408.	3.5	35

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91	Molecular similarity between myelodysplastic form of chronic myelomonocytic leukemia and refractory anemia with ring sideroblasts. Haematologica, 2013, 98, 576-583.	3.5	9
92	Impact of isolated germline JAK2V617I mutation on human hematopoiesis. Blood, 2013, 121, 4156-4165.	1.4	42
93	Aberrant DNA methylation profile of chronic and transformed classic Philadelphia-negative myeloproliferative neoplasms. Haematologica, 2013, 98, 1414-1420.	3.5	46
94	Limited clinical activity of nilotinib and sorafenib in FIP1L1-PDGFRA positive chronic eosinophilic leukemia with imatinib-resistant T674I mutation. Leukemia, 2012, 26, 162-164.	7.2	55
95	Philadelphia Chromosome-Negative Myeloproliferative Neoplasm With a Novel Platelet-Derived Growth Factor Receptor-Î² Rearrangement Responsive to Imatinib. Journal of Clinical Oncology, 2012, 30, e109-e111.	1.6	4
96	Decrease in JAK2V617F allele burden is not a prerequisite to clinical response in patients with polycythemia vera. Haematologica, 2012, 97, 538-542.	3.5	31
97	Inactivation of polycomb repressive complex 2 components in myeloproliferative and myelodysplastic/myeloproliferative neoplasms. Blood, 2012, 119, 1208-1213.	1.4	162
98	Activating CBL mutations are associated with a distinct MDS/MPN phenotype. Annals of Hematology, 2012, 91, 1713-1720.	1.8	29
99	TET2 Mutations Are Associated with Specific 5-Methylcytosine and 5-Hydroxymethylcytosine Profiles in Patients with Chronic Myelomonocytic Leukemia. PLoS ONE, 2012, 7, e31605.	2.5	70
100	Acquired Uniparental Disomy in Myeloproliferative Neoplasms. Hematology/Oncology Clinics of North America, 2012, 26, 981-991.	2.2	14
101	Standardized definitions of molecular response in chronic myeloid leukemia. Leukemia, 2012, 26, 2172-2175.	7.2	339
102	Mutations affecting mRNA splicing define distinct clinical phenotypes and correlate with patient outcome in myelodysplastic syndromes. Blood, 2012, 119, 3211-3218.	1.4	220
103	Favorable outcome of allogeneic hematopoietic cell transplantation for 8p11 myeloproliferative syndrome associated with BCRâ€FGFR1 gene fusion. Pediatric Blood and Cancer, 2012, 59, 194-196.	1.5	25
104	Molecular profiling of chronic myelomonocytic leukemia reveals diverse mutations in >80% of patients with TET2 and EZH2 being of high prognostic relevance. Leukemia, 2011, 25, 877-879.	7.2	131
105	BCR-ABL kinase domain mutation analysis in chronic myeloid leukemia patients treated with tyrosine kinase inhibitors: recommendations from an expert panel on behalf of European LeukemiaNet. Blood, 2011, 118, 1208-1215.	1.4	486
106	Novel imatinib-sensitive PDGFRA-activating point mutations in hypereosinophilic syndrome induce growth factor independence and leukemia-like disease. Blood, 2011, 117, 2935-2943.	1.4	76
107	Response of ETV6-FLT3â€“positive myeloid/lymphoid neoplasm with eosinophilia to inhibitors of FMS-like tyrosine kinase 3. Blood, 2011, 118, 2239-2242.	1.4	75
108	EZH2 mutational status predicts poor survival in myelofibrosis. Blood, 2011, 118, 5227-5234.	1.4	242

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109	The European LeukemiaNet: achievements and perspectives. Haematologica, 2011, 96, 156-162.	3.5	15
110	Identification of FOXP1 and SNX2 as novel ABL1 fusion partners in acute lymphoblastic leukaemia. British Journal of Haematology, 2011, 153, 43-46.	2.5	40
111	Guidelines for the measurement of BCRâ€ABL1 transcripts in chronic myeloid leukaemia. British Journal of Haematology, 2011, 153, 179-190.	2.5	94
112	IDH2 somatic mutations in chronic myeloid leukemia patients in blast crisis. Leukemia, 2011, 25, 178-181.	7.2	27
113	The t(4;9)(q11;q33) fuses CEP110 to KIT in a case of acute myeloid leukemia. Leukemia, 2011, 25, 1049-1050.	7.2	2
114	The myeloproliferative neoplasm-associated JAK2 46/1 haplotype is not overrepresented in chronic myelogenous leukemia. Annals of Hematology, 2011, 90, 365-366.	1.8	6
115	JAK2V617F allele burden in polycythemia vera correlates with grade of myelofibrosis, but is not substantially affected by therapy. Leukemia Research, 2011, 35, 177-182.	0.8	56
116	Aberrations of EZH2 in Cancer. Clinical Cancer Research, 2011, 17, 2613-2618.	7.0	503
117	Impact ofBCR-ABLmutations on patients with chronic myeloid leukemia. Cell Cycle, 2011, 10, 250-260.	2.6	64
118	TFG, a target of chromosome translocations in lymphoma and soft tissue tumors, fuses to GPR128 in healthy individuals. Haematologica, 2010, 95, 20-26.	3.5	63
119	Transcription factor mutations in myelodysplastic/myeloproliferative neoplasms. Haematologica, 2010, 95, 1473-1480.	3.5	67
120	A polymorphism associated with STAT3 expression and response of chronic myeloid leukemia to interferon Â. Haematologica, 2010, 95, 148-152.	3.5	29
121	Screening for diverse PDGFRA or PDGFRB fusion genes is facilitated by generic quantitative reverse transcriptase polymerase chain reaction analysis. Haematologica, 2010, 95, 738-744.	3.5	72
122	The t(14;20) is a poor prognostic factor in myeloma but is associated with long-term stable disease in monoclonal gammopathies of undetermined significance. Haematologica, 2010, 95, 1221-1225.	3.5	84
123	In search of the original leukemic clone in chronic myeloid leukemia patients in complete molecular remission after stem cell transplantation or imatinib. Blood, 2010, 116, 1329-1335.	1.4	78
124	Atypical mRNA fusions in PMLâ€RARA positive, RARAâ€PML negative acute promyelocytic leukemia. Genes Chromosomes and Cancer, 2010, 49, 471-479.	2.8	26
125	Analysis of genomic breakpoints in p190 and p210 BCRâ€“ABL indicate distinct mechanisms of formation. Leukemia, 2010, 24, 1742-1750.	7.2	43
126	Inactivating mutations of the histone methyltransferase gene EZH2 in myeloid disorders. Nature Genetics, 2010, 42, 722-726.	21.4	1,034

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127	Interlaboratory Diagnostic Validation of Conformation-Sensitive Capillary Electrophoresis for Mutation Scanning. Clinical Chemistry, 2010, 56, 593-602.	3.2	11
128	Establishment of the first World Health Organization International Genetic Reference Panel for quantitation of BCR-ABL mRNA. Blood, 2010, 116, e111-e117.	1.4	141
129	No association between myeloproliferative neoplasms and the Crohn's disease-associated STAT3 predisposition SNP rs744166. Haematologica, 2010, 95, 1226-1227.	3.5	3
130	The JAK2 46/1 haplotype predisposes to MPL-mutated myeloproliferative neoplasms. Blood, 2010, 115, 4517-4523.	1.4	93
131	Loss of 1p and rearrangement of MYC are associated with progression of smouldering myeloma to myeloma: sequential analysis of a single case. Haematologica, 2009, 94, 1024-1028.	3.5	47
132	Molecular basis of myelodysplastic/myeloproliferative neoplasms. Haematologica, 2009, 94, 1634-1638.	3.5	41
133	Frequent upregulation of MYC in plasma cell leukemia. Genes Chromosomes and Cancer, 2009, 48, 624-636.	2.8	65
134	No evidence for increased prevalence of JAK2 V617F in women with a history of recurrent miscarriage. British Journal of Haematology, 2009, 144, 802-803.	2.5	12
135	Clonal diversity in the myeloproliferative neoplasms: independent origins of genetically distinct clones. British Journal of Haematology, 2009, 144, 904-908.	2.5	75
136	Acute myeloid leukaemia with associated eosinophilia: justification for FIP1L1â€PDGFRA screening in cases lacking the CBFBâ€MYH11 fusion gene. British Journal of Haematology, 2009, 146, 225-227.	2.5	11
137	Imatinib sensitivity as a consequence of a CSF1R-Y571D mutation and CSF1/CSF1R signaling abnormalities in the cell line GDM1. Leukemia, 2009, 23, 358-364.	7.2	13
138	Detection and molecular monitoring of FIP1L1-PDGFRA-positive disease by analysis of patient-specific genomic DNA fusion junctions. Leukemia, 2009, 23, 332-339.	7.2	35
139	The molecular anatomy of the FIP1L1-PDGFRA fusion gene. Leukemia, 2009, 23, 271-278.	7.2	23
140	Harmonization of molecular monitoring of CML therapy in Europe. Leukemia, 2009, 23, 1957-1963.	7.2	196
141	Clinical evidence for a graft-versus-tumour effect following allogeneic HSCT for t(8;13) atypical myeloproliferative disorder. Bone Marrow Transplantation, 2009, 44, 197-199.	2.4	4
142	JAK2 haplotype is a major risk factor for the development of myeloproliferative neoplasms. Nature Genetics, 2009, 41, 446-449.	21.4	365
143	Standardisation of molecular monitoring for chronic myeloid leukaemia. Best Practice and Research in Clinical Haematology, 2009, 22, 355-365.	1.7	85
144	Timing of acquisition of deletion 13 in plasma cell dyscrasias is dependent on genetic context. Haematologica, 2009, 94, 1708-1713.	3.5	68

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145	Frequent CBL mutations associated with 11q acquired uniparental disomy in myeloproliferative neoplasms. Blood, 2009, 113, 6182-6192.	1.4	349
146	Evolutional change of karyotype with t(8;9)(p22;p24) and HLA-DR immunophenotype in relapsed acute myeloid leukemia. International Journal of Hematology, 2008, 88, 197-201.	1.6	11
147	The t(1;9)(p34;q34) and t(8;12)(p11;q15) fuse preâ€mRNA processing proteins SFPQ (PSF) and CPSF6 to ABL and FGFR1. Genes Chromosomes and Cancer, 2008, 47, 379-385.	2.8	55
148	International standardisation of quantitative real-time RT-PCR for BCR-ABL. Leukemia Research, 2008, 32, 505-506.	0.8	30
149	Comparison of mutated ABL1 and JAK2 as oncogenes and drug targets in myeloproliferative disorders. Leukemia, 2008, 22, 1320-1334.	7.2	24
150	BCR-ABL1-positive CML and BCR-ABL1-negative chronic myeloproliferative disorders: some common and contrasting features. Leukemia, 2008, 22, 1975-1989.	7.2	21
151	JAK2 Mutations are present in all cases of polycythemia vera. Leukemia, 2008, 22, 1289-1289.	7.2	53
152	Safety and efficacy of imatinib in chronic eosinophilic leukaemia and hypereosinophilic syndrome â€“ a phaseâ€ study. British Journal of Haematology, 2008, 143, 707-715.	2.5	128
153	Fibroblast Growth Factor Receptor and Platelet-Derived Growth Factor Receptor Abnormalities in Eosinophilic Myeloproliferative Disorders. Acta Haematologica, 2008, 119, 199-206.	1.4	59
154	Rapid identification of JAK2 exon 12 mutations using high resolution melting analysis. Haematologica, 2008, 93, 1560-1564.	3.5	49
155	Desirable performance characteristics for BCR-ABL measurement on an international reporting scale to allow consistent interpretation of individual patient response and comparison of response rates between clinical trials. Blood, 2008, 112, 3330-3338.	1.4	350
156	Chronic Myeloproliferative Disorders: The Role of Tyrosine Kinases in Pathogenesis, Diagnosis and Therapy. Pathobiology, 2007, 74, 81-88.	3.8	29
157	Durable responses to imatinib in patients with PDGFRB fusion geneâ€“positive and BCR-ABLâ€“negative chronic myeloproliferative disorders. Blood, 2007, 109, 61-64.	1.4	156
158	Activity of TKI258 against primary cells and cell lines with FGFR1 fusion genes associated with the 8p11 myeloproliferative syndrome. Blood, 2007, 110, 3729-3734.	1.4	91
159	Low-dose imatinib mesylate leads to rapid induction of major molecular responses and achievement of complete molecular remission in FIP1L1-PDGFRAâ€“positive chronic eosinophilic leukemia. Blood, 2007, 109, 4635-4640.	1.4	195
160	Heterogeneous prognostic impact of derivative chromosome 9 deletions in chronic myelogenous leukemia. Blood, 2007, 110, 1283-1290.	1.4	72
161	Gene mapping and expression analysis of 16q loss of heterozygosity identifies WWOX and CYLD as being important in determining clinical outcome in multiple myeloma. Blood, 2007, 110, 3291-3300.	1.4	133
162	Tyrosine Kinases as Therapeutic Targets in BCR-ABL Negative Chronic Myeloproliferative Disorders. Current Drug Targets, 2007, 8, 205-216.	2.1	30

#	Article	IF	Citations
163	Amplification Refractory Mutation System, a Highly Sensitive and Simple Polymerase Chain Reaction Assay, for the Detection of JAK2 V617F Mutation in Chronic Myeloproliferative Disorders. Journal of Molecular Diagnostics, 2007, 9, 272-276.	2.8	60
164	Methylation-Sensitive High-Resolution Melting-Curve Analysis of the SNRPN Gene as a Diagnostic Screen for Prader-Willi and Angelman Syndromes. Clinical Chemistry, 2007, 53, 1960-1962.	3.2	87
165	Characterization of three new imatinib-responsive fusion genes in chronic myeloproliferative disorders generated by disruption of the platelet-derived growth factor receptor Â gene. Haematologica, 2007, 92, 163-169.	3.5	84
166	Evaluation of JAK2^{V617F} in B and T Cell Neoplasms: Identification of JAK2^{V617F} Mutation of Undetermined Significance (JMUS) in the Bone Marrow of Three Individuals. Acta Haematologica, 2007, 118, 209-214.	1.4	17
167	FIP1L1-PDGFRA in chronic eosinophilic leukemia and BCR-ABL1 in chronic myeloid leukemia affect different leukemic cells. Leukemia, 2007, 21, 397-402.	7.2	21
168	Recurrent finding of the FIP1L1-PDGFRA fusion gene in eosinophilia-associated acute myeloid leukemia and lymphoblastic T-cell lymphoma. Leukemia, 2007, 21, 1183-1188.	7.2	170
169	A novel ETV6-PDGFRB fusion transcript missed by standard screening in a patient with an imatinib responsive chronic myeloproliferative disease. Leukemia, 2007, 21, 1839-1841.	7.2	33
170	Chronic myeloid leukemia â€“ some topical issues. Leukemia, 2007, 21, 1347-1352.	7.2	12
171	The severity of FIP1L1â€“PDGFRA-positive chronic eosinophilic leukaemia is associated with polymorphic variation at the IL5RA locus. Leukemia, 2007, 21, 2428-2432.	7.2	26
172	Two novel imatinibâ€responsive PDGFRA fusion genes in chronic eosinophilic leukaemia. British Journal of Haematology, 2007, 138, 77-81.	2.5	79
173	A constitutively active SPTBN1-FLT3 fusion in atypical chronic myeloid leukemia is sensitive to tyrosine kinase inhibitors and immunotherapy. Experimental Hematology, 2007, 35, 1723-1727.	0.4	44
174	Simultaneous MLPA-based multiplex point mutation and deletion analysis of the Dystrophin gene. Molecular Biotechnology, 2007, 35, 135-140.	2.4	24
175	Diagnostic and therapeutic management of eosinophilia-associated chronic myeloproliferative disorders. Haematologica, 2007, 92, 1153-1158.	3.5	14
176	Monitoring CML patients responding to treatment with tyrosine kinase inhibitors: review and recommendations for harmonizing current methodology for detecting BCR-ABL transcripts and kinase domain mutations and for expressing results. Blood, 2006, 108, 28-37.	1.4	1,117
177	Eosinophilic disorders: Molecular pathogenesis, new classification, and modern therapy. Best Practice and Research in Clinical Haematology, 2006, 19, 535-569.	1.7	60
178	Signal transduction therapy in haematological malignancies: identification and targeting of tyrosine kinases. Clinical Science, 2006, 111, 233-249.	4.3	10
179	Serial measurement of BCR-ABL transcripts in the peripheral blood after allogeneic stem cell transplantation for chronic myeloid leukemia: an attempt to define patients who may not require further therapy. Blood, 2006, 107, 4171-4176.	1.4	119
180	Minimal molecular response in polycythemia vera patients treated with imatinib or interferon alpha. Blood, 2006, 107, 3339-3341.	1.4	113

#	Article	IF	Citations
181	International Working Group (IWG) consensus criteria for treatment response in myelofibrosis with myeloid metaplasia, for the IWG for Myelofibrosis Research and Treatment (IWG-MRT). Blood, 2006, 108, 1497-1503.	1.4	317
182	JAK2 V617F mutation in classic chronic myeloproliferative diseases: a report on a series of 349 patients. Leukemia, 2006, 20, 534-535.	7.2	41
183	Identification of a novel imatinib responsive KIF5B-PDGFRA fusion gene following screening for PDGFRA overexpression in patients with hypereosinophilia. Leukemia, 2006, 20, 827-832.	7.2	122
184	Deletion of chromosome 13 detected by conventional cytogenetics is a critical prognostic factor in myeloma. Leukemia, 2006, 20, 1610-1617.	7.2	141
185	JAK2V617F mutational frequency in polycythemia vera: 100%, >90%, less?. Leukemia, 2006, 20, 2067-2067.	7.2	45
186	Rationale for the recommendations for harmonizing current methodology for detecting BCR-ABL transcripts in patients with chronic myeloid leukaemia. Leukemia, 2006, 20, 1925-1930.	7.2	184
187	A novel K509I mutation of KIT identified in familial mastocytosisâ€"in vitro and in vivo responsiveness to imatinib therapy. Leukemia Research, 2006, 30, 373-378.	0.8	129
188	Transient response to imatinib in a chronic eosinophilic leukemia associated with ins(9;4)(q33;q12q25) and a CDK5RAP2â€PDGFRA fusion gene. Genes Chromosomes and Cancer, 2006, 45, 950-956.	2.8	72
189	Cough and hypereosinophilia due to FIP1L1-PDGFRA fusion gene with tyrosine kinase activity. European Respiratory Journal, 2006, 27, 230-232.	6.7	37
190	Idiopathic Hypereosinophilic Syndrome in Children. Journal of Pediatric Hematology/Oncology, 2005, 27, 663-665.	0.6	38
191	NUP98-LEDGF fusion and t(9;11) in transformed chronic myeloid leukemia. Leukemia Research, 2005, 29, 1469-1472.	0.8	28
192	Der(6)t(1;6)(q21-23;p21.3): a specific cytogenetic abnormality in myelofibrosis with myeloid metaplasia. British Journal of Haematology, 2005, 130, 229-232.	2.5	47
193	Disruption and aberrant expression of HMGA2 as a consequence of diverse chromosomal translocations in myeloid malignancies. Leukemia, 2005, 19, 245-252.	7.2	69
194	Broad molecular screening of an unclassifiable myeloproliferative disorder reveals an unexpected ETV6/ABL1 fusion transcript. Leukemia, 2005, 19, 1096-1099.	7.2	20
195	The t(8;17)(p11;q23) in the 8p11 myeloproliferative syndrome fuses MYO18A to FGFR1. Leukemia, 2005, 19, 1005-1009.	7.2	85
196	Age has a profound effect on the incidence and significance of chromosome abnormalities in myeloma. Leukemia, 2005, 19, 1634-1642.	7.2	73
197	Persistence of bone marrow micrometastases in patients receiving adjuvant therapy for breast cancer: Results at 4 years. International Journal of Cancer, 2005, 114, 94-100.	5.1	63
198	DNA Topoisomerase II in Therapy-Related Acute Promyelocytic Leukemia. New England Journal of Medicine, 2005, 352, 1529-1538.	27.0	262

#	Article	IF	Citations
199	The t(8;9)(p22;p24) Is a Recurrent Abnormality in Chronic and Acute Leukemia that Fuses PCM1 to JAK2. Cancer Research, 2005, 65, 2662-2667.	0.9	269
200	Accurate Detection and Quantitation of Heteroplasmic Mitochondrial Point Mutations by Pyrosequencing. Genetic Testing and Molecular Biomarkers, 2005, 9, 190-199.	1.7	94
201	Widespread occurrence of the JAK2 V617F mutation in chronic myeloproliferative disorders. Blood, 2005, 106, 2162-2168.	1.4	798
202	Hydroxyurea Compared with Anagrelide in High-Risk Essential Thrombocythemia. New England Journal of Medicine, 2005, 353, 33-45.	27.0	838
203	p53-Binding Protein 1 Is Fused to the Platelet-Derived Growth Factor Receptor Î² in a Patient with a t(5;15)(q33;q22) and an Imatinib-Responsive Eosinophilic Myeloproliferative Disorder. Cancer Research, 2004, 64, 7216-7219.	0.9	74
204	NIN, a Gene Encoding a CEP110-Like Centrosomal Protein, Is Fused to PDGFRB in a Patient with a t(5;14)(q33;q24) and an Imatinib-Responsive Myeloproliferative Disorder 1. Cancer Research, 2004, 64, 2673-2676.	0.9	67
205	Critical Role of STAT5 Activation in Transformation Mediated by ZNF198-FGFR1. Journal of Biological Chemistry, 2004, 279, 6666-6673.	3.4	43
206	Low expression of the putative tumour suppressor gene gravin in chronic myeloid leukaemia, myelodysplastic syndromes and acute myeloid leukaemia. British Journal of Haematology, 2004, 126, 508-511.	2.5	12
207	Targeting FGFR3 in multiple myeloma: inhibition of t(4;14)-positive cells by SU5402 and PD173074. Leukemia, 2004, 18, 962-966.	7.2	137
208	Dosage analysis of cancer predisposition genes by multiplex ligation-dependent probe amplification. British Journal of Cancer, 2004, 91, 1155-1159.	6.4	161
209	Distinct stem cell myeloproliferative/T lymphoma syndromes induced by ZNF198-FGFR1 and BCR-FGFR1 fusion genes from 8p11 translocations. Cancer Cell, 2004, 5, 287-298.	16.8	145
210	Oncogenic protein tyrosine kinases. Cellular and Molecular Life Sciences, 2004, 61, 2912-2923.	5.4	126
211	Identification of a novel gene, FGFR1OP2, fused to FGFR1 in 8p11 myeloproliferative syndrome. Genes Chromosomes and Cancer, 2004, 40, 78-83.	2.8	89
212	Clinical variability of patients with the t(6;8)(q27;p12) and FGFR1OP-FGFR1 fusion: two further cases. The Hematology Journal, 2004, 5, 534-537.	1.4	49
213	Paucimorphic Alleles versus Polymorphic Alleles and Rare Mutations in Disease Causation: Theory, Observation and Detection. Current Genomics, 2004, 5, 431-438.	1.6	7
214	MOZ-TIF2-induced acute myeloid leukemia requires the MOZ nucleosome binding motif and TIF2-mediated recruitment of CBP. Cancer Cell, 2003, 3, 259-271.	16.8	192
215	Genomic anatomy of the specific reciprocal translocation t(15;17) in acute promyelocytic leukemia. Genes Chromosomes and Cancer, 2003, 36, 175-188.	2.8	58
216	Novel translocations that disrupt the plateletâ€derived growth factor receptor Î² (PDGFRB) gene in BCRâ€“ABLâ€negative chronic myeloproliferative disorders. British Journal of Haematology, 2003, 120, 251-256.	2.5	87

#	Article	IF	Citations
217	P2X7 polymorphism and chronic lymphocytic leukaemia: lack of correlation with incidence, survival and abnormalities of chromosome 12. Leukemia, 2003, 17, 2097-2100.	7.2	42
218	Imatinib for systemic mast-cell disease. Lancet, The, 2003, 362, 535-537.	13.7	242
219	A Tyrosine Kinase Created by Fusion of thePDGFRAandFIP1L1Genes as a Therapeutic Target of Imatinib in Idiopathic Hypereosinophilic Syndrome. New England Journal of Medicine, 2003, 348, 1201-1214.	27.0	1,655
220	Imatinib therapy for hypereosinophilic syndrome and other eosinophilic disorders. Blood, 2003, 101, 3391-3397.	1.4	206
221	CHIC2 deletion, a surrogate for FIP1L1-PDGFRA fusion, occurs in systemic mastocytosis associated with eosinophilia and predicts response to imatinib mesylate therapy. Blood, 2003, 102, 3093-3096.	1.4	368
222	The t(4;22)(q12;q11) in atypical chronic myeloid leukaemia fuses BCR to PDGFRA. Human Molecular Genetics, 2002, 11, 1391-1397.	2.9	139
223	Response to Imatinib Mesylate in Patients with Chronic Myeloproliferative Diseases with Rearrangements of the Platelet-Derived Growth Factor Receptor Beta. New England Journal of Medicine, 2002, 347, 481-487.	27.0	623
224	The 8p11 Myeloproliferative Syndrome: A Distinct Clinical Entity Caused by Constitutive Activation of <i>FGFR1</i>. Acta Haematologica, 2002, 107, 101-107.	1.4	220
225	Chronic Eosinophilic Leukaemia Presenting with Erythroderma, Mild Eosinophilia and Hyper-IgE: Clinical, Immunological and Cytogenetic Features and Therapeutic Approach. Acta Haematologica, 2002, 107, 108-112.	1.4	27
226	Myeloproliferative Disorders with Translocations of Chromosome 5q31â€“35: Role of the Platelet-Derived Growth Factor Receptor Beta. Acta Haematologica, 2002, 107, 113-122.	1.4	170
227	Tyrosine kinase fusion genes in chronic myeloproliferative diseases. Leukemia, 2002, 16, 1207-1212.	7.2	124
228	Molecular and chromosomal mechanisms of resistance to imatinib (STI571) therapy. Leukemia, 2002, 16, 2190-2196.	7.2	839
229	Myeloproliferative disorders. Best Practice and Research in Clinical Haematology, 2001, 14, 531-551.	1.7	39
230	Cytogenetics of chronic myeloid leukaemia. Best Practice and Research in Clinical Haematology, 2001, 14, 553-571.	1.7	36
231	ABL-BCR expression does not correlate with deletions on the derivative chromosome 9 or survival in chronic myeloid leukemia. Blood, 2001, 98, 2879-2880.	1.4	38
232	Early detection of BCR-ABL transcripts by quantitative reverse transcriptaseâ€“polymerase chain reaction predicts outcome after allogeneic stem cell transplantation for chronic myeloid leukemia. Blood, 2001, 97, 1560-1565.	1.4	154
233	The influence of INK4 proteins on growth and self-renewal kinetics of hematopoietic progenitor cells. Blood, 2001, 97, 2604-2610.	1.4	46
234	The t(8;22) in chronic myeloid leukemia fuses BCR to FGFR1: transforming activity and specific inhibition of FGFR1 fusion proteins. Blood, 2001, 98, 3778-3783.	1.4	189

#	Article	IF	Citations
235	A novel gene, NSD1, is fused to NUP98 in the t(5;11)(q35;p15.5) in de novo childhood acute myeloid leukemia. Blood, 2001, 98, 1264-1267.	1.4	245
236	Molecular studies in patients with chronic myeloid leukaemia in remission 5 years after allogeneic stem cell transplant define the risk of subsequent relapse. British Journal of Haematology, 2001, 115, 569-574.	2.5	66
237	Identification of four new translocations involving FGFR1 in myeloid disorders. Genes Chromosomes and Cancer, 2001, 32, 155-163.	2.8	91
238	Mutations of the transcription factor AML1/CBA2 are uncommon in blastic transformation of chronic myeloid leukaemia. Leukemia, 2001, 15, 476-477.	7.2	8
239	Detection and quantification of residual disease in chronic myelogenous leukemia. Leukemia, 2000, 14, 998-1005.	7.2	105
240	Cloning of ZNF237, a novel member of the MYM gene family that maps to human chromosome 13q11â†'q12. Cytogenetic and Genome Research, 2000, 89, 24-28.	1.1	5
241	Non-random involvement of chromosome 13 in patients with persistent or relapsed disease after bone-marrow transplantation for chronic myeloid leukemia. Genes Chromosomes and Cancer, 2000, 27, 278.	2.8	1
242	Assignment<footref rid="foot01">¹</footref> of ZNF262 to human chromosome band 1p34â†'p32 by in situ hybridization. Cytogenetic and Genome Research, 1999, 85, 306-307.	1.1	7
243	A case of myelofibrosis with a t(4;13)(q25;q12): evidence for involvement of a second 13q12 locus in chronic myeloproliferative disorders. British Journal of Haematology, 1999, 105, 771-774.	2.5	7
244	Quantification of minimal residual disease in patients with BCR-ABL-positive acute lymphoblastic leukaemia using quantitative competitive polymerase chain reaction. British Journal of Haematology, 1999, 106, 634-643.	2.5	50
245	Two patients with novel BCR/ABL fusion transcripts (e8/a2 and e13/a2) resulting from translocation breakpoints within BCR exons. British Journal of Haematology, 1999, 105, 434-436.	2.5	28
246	Accurate and rapid analysis of residual disease in patients with CML using specific fluorescent hybridization probes for real time quantitative RT-PCR. Leukemia, 1999, 13, 1825-1832.	7.2	255
247	Consistent Fusion of MOZ and TIF2 in AML with inv(8)(p11q13). Cancer Genetics and Cytogenetics, 1999, 113, 70-72.	1.0	50
248	The Genomic Structure of ZNF198 and Location of Breakpoints in the t(8;13) Myeloproliferative Syndrome. Genomics, 1999, 55, 118-121.	2.9	24
249	Cloning and Characterization ofRNF6,a Novel RING Finger Gene Mapping to 13q12. Genomics, 1999, 58, 94-97.	2.9	20
250	Adoptive immunotherapy for relapse of chronic myeloid leukemia after allogeneic bone marrow transplant: equal efficacy of lymphocytes from sibling and matched unrelated donors. Bone Marrow Transplantation, 1998, 21, 1055-1061.	2.4	67
251	Aberrant transcripts of the FHIT gene are expressed in normal and leukaemic haemopoietic cells. British Journal of Cancer, 1998, 78, 601-605.	6.4	10
252	BCR-ABL-positive progenitors in chronic myeloid leukaemia patients in complete cytogenetic remission after treatment with interferon-Î±. British Journal of Haematology, 1998, 102, 1271-1278.	2.5	18

#	Article	IF	Citations
253	A novel BCR-ABL fusion gene (e2/1a) in a patient with Philadelphia-positive chronic myelogenous leukaemia and an aggressive clinical course. British Journal of Haematology, 1998, 103, 791-794.	2.5	22
254	Assignment of the Steroid Receptor Coactivator-1 (SRC-1) Gene to Human Chromosome Band 2p23. Genomics, 1998, 52, 242-244.	2.9	10
255	The commonly deleted region at 9p21-22 in lymphoblastic leukemias spans at least 400 kb and includes p16 but not p15 or the IFN gene cluster. Leukemia, 1997, 11, 233-238.	7.2	29
256	An extremely delayed cytogenetic response to interferon-Î± in a patient with chronic myeloid leukaemia. Leukemia, 1997, 11, 614-616.	7.2	2
257	Rarity of dominant-negative mutations of the G-CSF receptor in patients with blast crisis of chronic myeloid leukemia or de novo acute leukemia. Leukemia, 1997, 11, 1005-1008.	7.2	24
258	Molecular response of CML patients treated with interferonâ€Î± monitored by quantitative Southern blot analysis. British Journal of Haematology, 1997, 97, 86-93.	2.5	34
259	The rate and kinetics of molecular response to donor leucocyte transfusions in chronic myeloid leukaemia patients treated for relapse after allogeneic bone marrow transplantation. British Journal of Haematology, 1997, 99, 945-950.	2.5	72
260	Rarity of microsatellite alterations in acute myeloid leukaemia. British Journal of Cancer, 1996, 74, 255-257.	6.4	37
261	Mutational analysis of the p15 and p16 genes in acute leukaemias. British Journal of Haematology, 1996, 92, 681-683.	2.5	22
262	Quantification of PMLâ€RARÎ± transcripts in acute promyelocytic leukaemia: explanation for the lack of sensitivity of RTâ€PCR for the detection of minimal residual disease and induction of the leukaemiaâ€specific mRNA by alpha interferon. British Journal of Haematology, 1996, 95, 95-101.	2.5	40
263	TELâ€AML1 fusion in acute lymphoblastic leukaemia of adults. British Journal of Haematology, 1996, 95, 673-677.	2.5	86
264	Characterization of genomic BCR-ABL breakpoints in chronic myeloid leukaemia by PCR. British Journal of Haematology, 1995, 90, 138-146.	2.5	63
265	Variable numbers of BCR-ABL transcripts persist in CML patients who achieve complete cytogenetic remission with interferon-Î±. British Journal of Haematology, 1995, 91, 126-131.	2.5	75
266	Localization of the 8;13 translocation breakpoint associated with myeloproliferative disease to a 1.5 mbp region of chromosome 13. Genes Chromosomes and Cancer, 1995, 12, 283-287.	2.8	25
267	Correlation between the proportion of Philadelphia chromosome-positive metaphase cells and levels ofBCR-ABL mRNA in chronic myeloid leukaemia. Genes Chromosomes and Cancer, 1995, 13, 110-114.	2.8	22
268	Null alleles of the aldolase B gene in patients with hereditary fructose intolerance.. Journal of Medical Genetics, 1994, 31, 499-503.	3.2	18
269	A comparison of the sensitivity of blood and bone marrow for the detection of minimal residual disease in chronic myeloid leukaemia. British Journal of Haematology, 1994, 86, 683-685.	2.5	56
270	An atypical myeloproliferative disorder with t(8;13) (p11;q12): a third case. British Journal of Haematology, 1994, 86, 879-880.	2.5	48

#	Article	IF	Citations
271	APPROPRIATE CONTROLS FOR REVERSE TRANSCRIPTION POLYMERASE CHAIN REACTION (RTâ€PCR). British Journal of Haematology, 1994, 87, 218-218.	2.5	18
272	Aplasia after donor lymphocyte infusion (DLI) for CML in relapse after sex-mismatched BMT: recovery of donor-type haemopoiesis predicted by non-isotopic in situ hybridization (ISH). British Journal of Haematology, 1994, 88, 400-402.	2.5	14
273	Minimal residual disease after allogeneic bone marrow transplantation for chronic myeloid leukaemia in first chronic phase: correlations with acute graftâ€versusâ€host disease and relapse. British Journal of Haematology, 1993, 84, 67-74.	2.5	159
274	Isolation and characterization of the genes encoding mouse and human type-5 acid phosphatase. Gene, 1993, 130, 201-207.	2.2	46
275	BCR-ABL antisense purging in chronic myeloid leukaemia. Lancet, The, 1993, 342, 614.	13.7	35
276	Minimal Residual Disease after Bone Marrow Transplant for Chronic Myeloid Leukaemia Detected by the Polymerase Chain Reaction. Leukemia and Lymphoma, 1993, 11, 39-43.	1.3	57
277	Aldolase B mutations in Italian families affected by hereditary fructose intolerance.. Journal of Medical Genetics, 1991, 28, 241-243.	3.2	20
278	Type 5 acid phosphatase. Sequence, expression and chromosomal localization of a differentiation-associated protein of the human macrophage. FEBS Journal, 1990, 189, 287-293.	0.2	88
279	Hereditary fructose intolerance. International Journal of Biochemistry & Cell Biology, 1990, 22, 685-689.	0.5	12
280	A new aldolase B variant, N334K, is a common cause of hereditary fructose intolerance in Yugoslavia. Nucleic Acids Research, 1990, 18, 1925-1925.	14.5	29
281	Molecular analysis of aldolase B genes in hereditary fructose intolerance. Lancet, The, 1990, 335, 306-309.	13.7	127
282	Catalytic deficiency of human aldolase B in hereditary fructose intolerance caused by a common missense mutation. Cell, 1988, 53, 881-885.	28.9	98
283	A potent inhibitor ofTaqpolymerase copurifies with human genomic DNA. Nucleic Acids Research, 1988, 16, 10355-10355.	14.5	143
284	Polymerase chain reaction automated at low cost. Nucleic Acids Research, 1988, 16, 5687-5688.	14.5	22
285	Tsetse fly rDNA: an analysis of structure and sequence. Nucleic Acids Research, 1987, 15, 15-30.	14.5	40
286	A novel arrangement of sequence elements surrounding the rDNA promoter and its spacer duplications in tsetse species. Journal of Molecular Biology, 1987, 195, 63-74.	4.2	17