Kirsten Grønbæk

List of Publications by Year in descending order

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153

all docs

148 5,805 40 papers citations h-index

153

docs citations

h-index g-index

153 9266
times ranked citing authors

70

#	Article	IF	CITATIONS
1	Epigenetic therapy in combination with a multi-epitope cancer vaccine targeting shared tumor antigens for high-risk myelodysplastic syndromeÂ-Âa phase I clinical trial. Cancer Immunology, Immunotherapy, 2022, 71, 433-444.	2.0	8
2	Expression patterns and prognostic potential of circular RNAs in mantle cell lymphoma: a study of younger patients from the MCL2 and MCL3 clinical trials. Leukemia, 2022, 36, 177-188.	3.3	11
3	Preâ€treatment healthâ€related quality of life parameters have prognostic impact in patients >65Âyears with newly diagnosed mantle cell lymphoma: The Nordic Lymphoma Group MCL4 (LENAâ€BERIT) experience. Hematological Oncology, 2022, 40, 23-31.	0.8	1
4	Humoral response to two doses of BNT162b2 vaccination in people with HIV. Journal of Internal Medicine, 2022, 291, 513-518.	2.7	33
5	Incidence of Positive Severe Acute Respiratory Syndrome Coronavirus Polymerase Chain Reaction After Coronavirus Disease 2019 Vaccination With up to 8 Months of Follow-up: Real-life Data From the Capital Region of Denmark. Clinical Infectious Diseases, 2022, 75, e675-e682.	2.9	7
6	Level of unique T cell clonotypes is associated with clonal hematopoiesis and survival in patients with lymphoma undergoing ASCT. Bone Marrow Transplantation, 2022, , .	1.3	1
7	Comprehensive and unbiased multiparameter high-throughput screening by compaRe finds effective and subtle drug responses in AML models. ELife, 2022, 11 , .	2.8	2
8	"Randomized phase II study of azacitidine ± lenalidomide in higher-risk myelodysplastic syndromes and acute myeloid leukemia with a karyotype including Del(5q)― Leukemia, 2022, 36, 1436-1439.	3.3	6
9	A predictive model for bone marrow disease in cytopenia based on noninvasive procedures. Blood Advances, 2022, 6, 3541-3550.	2.5	2
10	Inflammatory Cytokine Profiles Do Not Differ Between Patients With Idiopathic Cytopenias of Undetermined Significance and Myelodysplastic Syndromes. HemaSphere, 2022, 6, e0713.	1.2	3
11	Family caregiver ambassador support for caregivers of patients with newly diagnosed hematological cancer: a feasibility study. Supportive Care in Cancer, 2022, 30, 6923-6935.	1.0	8
12	The Thioredoxin-Interacting Protein TXNIP Is a Putative Tumour Suppressor in Cutaneous T-Cell Lymphoma. Dermatology, 2021, 237, 283-290.	0.9	8
13	Protein phosphatase, Mg 2+ /Mn 2+ â€dependent 1D (PPM1D) mutations in haematological cancer. British Journal of Haematology, 2021, 192, 697-705.	1.2	7
14	Genome-Wide Circular RNA Expression Patterns Reflect Resistance to Immunomodulatory Drugs in Multiple Myeloma Cells. Cancers, 2021, 13, 365.	1.7	19
15	Risk of new malignancies among patients with CLL treated with chemotherapy: results of a Danish populationâ€based study. British Journal of Haematology, 2021, 193, 339-345.	1.2	12
16	The Danish Myelodysplastic Syndromes Database: Patient Characteristics and Validity of Data Records. Clinical Epidemiology, 2021, Volume 13, 439-451.	1.5	5
17	Mutations known from B-cell lymphoid malignancies are not found in CD34 ⁺ stem cells from patients with lymphoma. Leukemia and Lymphoma, 2021, 62, 2808-2811.	0.6	1
18	Acute and persistent symptoms in non-hospitalized PCR-confirmed COVID-19 patients. Scientific Reports, 2021, 11, 13153.	1.6	147

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19	The Impact of Sedentary Lifestyle, High-fat Diet, Tobacco Smoke, and Alcohol Intake on the Hematopoietic Stem Cell Niches. HemaSphere, 2021, 5, e615.	1.2	5
20	Serum proteome modulations upon treatment provides biological insight on response to treatment in relapsed mantle cell lymphoma. Cancer Reports, 2021, , e1524.	0.6	1
21	The role of vitamin C in epigenetic cancer therapy. Free Radical Biology and Medicine, 2021, 170, 179-193.	1.3	23
22	A user's guide to multicolor flow cytometry panels for comprehensive immune profiling. Analytical Biochemistry, 2021, 627, 114210.	1.1	12
23	Allogeneic Hematopoietic Stem Cell Transplantation for Chronic Myelomonocytic Leukemia: Clinical and Molecular Genetic Prognostic Factors in a Nordic Population. Transplantation and Cellular Therapy, 2021, 27, 991.e1-991.e9.	0.6	6
24	Detailed Long-Term Follow-Up of Patients Who Relapsed After the Nordic Mantle Cell Lymphoma Trials: MCL2 and MCL3. HemaSphere, 2021, 5, e510.	1.2	18
25	TET2 mutations are associated with hypermethylation at key regulatory enhancers in normal and malignant hematopoiesis. Nature Communications, 2021, 12, 6061.	5.8	47
26	Structural aberrations are associated with poor survival in patients with clonal cytopenia of undetermined significance. Haematologica, 2021, 106, 1762-1766.	1.7	6
27	Level of Unique T-Cell Clonotypes Are Associated with Clonal Hematopoiesis and Survival in Patients with Lymphoma Intended for Autologous Stem Cell Transplant. Blood, 2021, 138, 3942-3942.	0.6	0
28	<i>KMT2D</i> mutations and <i>TP53</i> disruptions are poor prognostic biomarkers in mantle cell lymphoma receiving high-dose therapy: a FIL study. Haematologica, 2020, 105, 1604-1612.	1.7	96
29	The diagnostic and prognostic role of flow cytometry in idiopathic and clonal cytopenia of undetermined significance (ICUS/CCUS): A singleâ€center analysis of 79 patients. Cytometry Part B - Clinical Cytometry, 2020, 98, 250-258.	0.7	10
30	Cellâ€ofâ€origin determined by both gene expression profiling and immunohistochemistry is the strongest predictor of survival in patients with diffuse large Bâ€cell lymphoma. American Journal of Hematology, 2020, 95, 57-67.	2.0	27
31	Human endogenous retroviruses form a reservoir of T cell targets in hematological cancers. Nature Communications, 2020, 11, 5660.	5.8	55
32	p53 is associated with highâ€risk and pinpoints <i>TP53</i> missense mutations in mantle cell lymphoma. British Journal of Haematology, 2020, 191, 796-805.	1.2	31
33	Identification of unique and shared mitochondrial DNA mutations in neurodegeneration and cancer by single-cell mitochondrial DNA structural variation sequencing (MitoSV-seq). EBioMedicine, 2020, 57, 102868.	2.7	11
34	Activation of a Subset of Evolutionarily Young Transposable Elements and Innate Immunity Are Linked to Clinical Responses to 5-Azacytidine. Cancer Research, 2020, 80, 2441-2450.	0.4	33
35	Clonal hematopoiesis evolves from pretreatment clones and stabilizes after end of chemotherapy in patients with MCL. Blood, 2020, 135, 2000-2004.	0.6	26
36	The value of circulating microRNAs for early diagnosis of B-cell lymphoma: A case-control study on historical samples. Scientific Reports, 2020, 10, 9637.	1.6	10

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37	Clinical impact of clonal hematopoiesis in patients with lymphoma undergoing ASCT: a national population-based cohort study. Leukemia, 2020, 34, 3256-3268.	3.3	46
38	Angiotensinogen promoter methylation predicts bevacizumab treatment response of patients with recurrent glioblastoma. Molecular Oncology, 2020, 14, 964-973.	2.1	2
39	Improved Outcomes after Allogenic Hematopoietic Stem Cell Transplantation with Fludarabine/Treosulfan for Patients with Myelodysplastic Syndromes. Biology of Blood and Marrow Transplantation, 2020, 26, 1091-1098.	2.0	7
40	Clonal hematopoiesis in elderly twins: concordance, discordance, and mortality. Blood, 2020, 135, 261-268.	0.6	47
41	Profiling of ribose methylations in ribosomal RNA from diffuse large B-cell lymphoma patients for evaluation of ribosomes as drug targets. NAR Cancer, 2020, 2, zcaa035.	1.6	29
42	Pre-Treatment Health-Related Quality of Life Parameters May Have Prognostic Impact in Elderly Patients with Mantle Cell Lymphoma. the Nordic Lymphoma Group MCL4 (LENA-BERIT) Experience. Blood, 2020, 136, 8-9.	0.6	0
43	Therapeutic Cancer Vaccination Targeting Shared Tumor Associated Antigens in Combination with Azacitidine for High Risk Myelodysplastic Syndrome - a Phase I Clinical Trial. Blood, 2020, 136, 23-24.	0.6	2
44	Oral vitamin C supplementation to patients with myeloid cancer on azacitidine treatment: Normalization of plasma vitamin C induces epigenetic changes. Clinical Epigenetics, 2019, 11, 143.	1.8	55
45	Increased neonatal level of arginase 2 in cases of childhood acute lymphoblastic leukemia implicates immunosuppression in the etiology. Haematologica, 2019, 104, e514-e516.	1.7	8
46	Epigenetic therapy in hematological cancers. Apmis, 2019, 127, 316-328.	0.9	16
47	Nordic Guidelines for Germline Predisposition to Myeloid Neoplasms in Adults: Recommendations for Genetic Diagnosis, Clinical Management and Follow-up. HemaSphere, 2019, 3, e321.	1.2	51
48	Expression of CRBN, IKZF1, and IKZF3 does not predict lenalidomide sensitivity and mutations in the cereblon pathway are infrequent in multiple myeloma. Leukemia and Lymphoma, 2019, 60, 180-188.	0.6	26
49	A Circular RNA Molecule, circRAB11FIP1, Is Associated with TP53 Mutations and Is of Potential Prognostic and Functional Significance in Mantle Cell Lymphoma: Data from the Nordic MCL2 and MCL3 Studies. Blood, 2019, 134, 1495-1495.	0.6	1
50	Diagnostic 2-Gene Classifier in Early-Stage Mycosis Fungoides: A Retrospective Multicenter Study. Blood, 2019, 134, 2772-2772.	0.6	0
51	Ibrutinib, lenalidomide, and rituximab in relapsed or refractory mantle cell lymphoma (PHILEMON): a multicentre, open-label, single-arm, phase 2 trial. Lancet Haematology,the, 2018, 5, e109-e116.	2.2	117
52	Vitamin C $\hat{a} \in A$ new player in regulation of the cancer epigenome. Seminars in Cancer Biology, 2018, 51, 59-67.	4.3	73
53	Dual inhibition of DNMTs and EZH2 can overcome both intrinsic and acquired resistance of myeloma cells to IMiDs in a cereblonâ€independent manner. Molecular Oncology, 2018, 12, 180-195.	2.1	62
54	Mother–child transmission of epigenetic information by tunable polymorphic imprinting. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E11970-E11977.	3.3	33

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55	Epidemiology of chronic redâ€cell transfusion recipients in Sweden and Denmark–a 10 year followâ€up study. Vox Sanguinis, 2018, 113, 770-778.	0.7	0
56	Human adult HSCs can be discriminated from lineage-committed HPCs by the expression of endomucin. Blood Advances, 2018, 2, 1628-1632.	2.5	10
57	Lenalidomide plus bendamustine-rituximab does not overcome the adverse impact of <i>TP53</i> mutations in mantle cell lymphoma. Haematologica, 2018, 103, e541-e543.	1.7	19
58	DNA Methylation Levels of the ELMO Gene Promoter CpG Islands in Human Glioblastomas. International Journal of Molecular Sciences, 2018, 19, 679.	1.8	19
59	Longâ€ŧerm clinical outcomes of patients with hematologically unexplained cytopenia after routine assessment: A single center study. European Journal of Haematology, 2018, 101, 595-603.	1.1	4
60	Long Non-Coding RNAs Guide the Fine-Tuning of Gene Regulation in B-Cell Development and Malignancy. International Journal of Molecular Sciences, 2018, 19, 2475.	1.8	33
61	Enzyme-free digital counting of endogenous circular RNA molecules in B-cell malignancies. Laboratory Investigation, 2018, 98, 1657-1669.	1.7	93
62	Phase I Results of a Multicenter Clinical Trial Combining Guadecitabine, a DNA Methyltransferase Inhibitor, with Atezolizumab, an Immune Checkpoint Inhibitor, in Patients with Relapsed or Refractory Myelodysplastic Syndrome or Chronic Myelomonocytic Leukemia. Blood, 2018, 132, 1811-1811.	0.6	7
63	Nucleosome Positioning and NDR Structure at RNA Polymerase III Promoters. Scientific Reports, 2017, 7, 41947.	1.6	29
64	Anemia is present years before myelodysplastic syndrome diagnosis: Results from the preâ€diagnostic period. American Journal of Hematology, 2017, 92, E130-E132.	2.0	5
65	Molecular Monitoring after Autologous Stem Cell Transplantation and Preemptive Rituximab Treatment of Molecular Relapse; Results from the Nordic Mantle Cell Lymphoma Studies (MCL2 and) Tj ETQq1 1 428-435.	0.784314	rgBT /Overlo
66	TP53 mutations identify younger mantle cell lymphoma patients who do not benefit from intensive chemoimmunotherapy. Blood, 2017, 130, 1903-1910.	0.6	296
67	DNA Methyltransferase Inhibitors in Myeloid Cancer. Cancer Journal (Sudbury, Mass), 2017, 23, 277-285.	1.0	6
68	Epigenetic changes in myelofibrosis: Distinct methylation changes in the myeloid compartments and in cases with ASXL1 mutations. Scientific Reports, 2017, 7, 6774.	1.6	16
69	Mature lymphoid malignancies: origin, stem cells, and chronicity. Blood Advances, 2017, 1, 2444-2455.	2.5	13
70	Immune Mechanisms in Myelodysplastic Syndrome. International Journal of Molecular Sciences, 2016, 17, 944.	1.8	48
71	Lenalidomide-bendamustine-rituximab in patients older than 65 years with untreated mantle cell lymphoma. Blood, 2016, 128, 1814-1820.	0.6	75
72	Frequent NFKBIE deletions are associated with poor outcome in primary mediastinal B-cell lymphoma. Blood, 2016, 128, 2666-2670.	0.6	82

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73	15â€year followâ€up of the Second Nordic Mantle Cell Lymphoma trial (<scp>MCL</scp> 2): prolonged remissions without survival plateau. British Journal of Haematology, 2016, 175, 410-418.	1.2	170
74	Vitamin C increases viral mimicry induced by 5-aza-2′-deoxycytidine. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10238-10244.	3.3	171
75	Mutations in idiopathic cytopenia of undetermined significance assist diagnostics and correlate to dysplastic changes. American Journal of Hematology, 2016, 91, 1234-1238.	2.0	32
76	TP53 hotspot mutations are predictive of survival in primary central nervous system lymphoma patients treated with combination chemotherapy. Acta Neuropathologica Communications, 2016, 4, 40.	2.4	19
77	Assessment of Quantitative and Allelic <i>MGMT</i> Methylation Patterns as a Prognostic Marker in Glioblastoma. Journal of Neuropathology and Experimental Neurology, 2016, 75, 246-255.	0.9	33
78	Molecular Monitoring and Tailored Strategy with Pre-Emptive Rituximab Treatment for Molecular Relapse; Results from the Nordic Mantle Cell Lymphoma Studies (MCL2 and MCL3) with Median Follow-up of 8.5 Years. Blood, 2016, 128, 146-146.	0.6	4
79	Ibrutinib-Lenalidomide-Rituximab in Patients with Relapsed/Refractory Mantle Cell Lymphoma: First Results from the Nordic Lymphoma Group MCL6 (PHILEMON) Phase II Trial. Blood, 2016, 128, 148-148.	0.6	12
80	Tumor suppressor microRNAs are downregulated in myelodysplastic syndrome with spliceosome mutations. Oncotarget, 2016, 7, 9951-9963.	0.8	27
81	ÎFÎŝÎΙΕ Deletions: A Novel Marker of Clinical Aggressiveness in Primary Mediastinal B-Cell Lymphoma. Blood, 2016, 128, 609-609.	0.6	0
82	Cereblon Is Downregulated By Promoter Nucleosome Occupancy in Acquired IMiD Resistance: The Potential of IMiD Resensitization By Epigenetic Therapy. Blood, 2016, 128, 3258-3258.	0.6	0
83	Increased Risk of Second Hematological and Non-Hematological Malignancies in CLL Patients Treated with Chemotherapy As Compared to Untreated Patients and Matched Controls - Results from a Danish Population Based Study. Blood, 2016, 128, 3219-3219.	0.6	0
84	Loss of PRDM11 promotes MYC-driven lymphomagenesis. Blood, 2015, 125, 1272-1281.	0.6	18
85	miR-18b overexpression identifies mantle cell lymphoma patients with poor outcome and improves the MIPI-B prognosticator. Blood, 2015, 125, 2669-2677.	0.6	44
86	Whole-exome sequencing and genome-wide methylation analyses identify novel disease associated mutations and methylation patterns in idiopathic hypereosinophilic syndrome. Oncotarget, 2015, 6, 40588-40597.	0.8	14
87	Hypermethylation of the VTRNA1-3 Promoter is Associated with Poor Outcome in Lower Risk Myelodysplastic Syndrome Patients. Genes, 2015, 6, 977-990.	1.0	19
88	Development and Blind Clinical Validation of a MicroRNA Based Predictor of Response to Treatment with R-CHO(E)P in DLBCL. PLoS ONE, 2015, 10, e0115538.	1.1	19
89	DNA Methylation and Hydroxymethylation in Cancer. , 2015, , 9-30.		4
90	Differential Expression of miR-155 and miR-21 in Tumor and Stroma Cells in Diffuse Large B-Cell Lymphoma. Applied Immunohistochemistry and Molecular Morphology, 2015, 23, 188-195.	0.6	10

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91	Allele-Specific DNA Methylation Detection by Pyrosequencing $\hat{A}^{@}$. Methods in Molecular Biology, 2015, 1315, 271-289.	0.4	5
92	Hypomethylation and up-regulation of <i> PD-1 < /i > in T cells by azacytidine in MDS/AML patients: A rationale for combined targeting of PD-1 and DNA methylation. Oncotarget, 2015, 6, 9612-9626.</i>	0.8	166
93	A novel del(8)(q23.2q24.11) contributing to disease progression in a case of JAK2/TET2 double mutated chronic myelomonocytic leukemia. Leukemia Research Reports, 2014, 3, 94-97.	0.2	1
94	Validation of a diagnostic microRNA classifier in cutaneous T-cell lymphomas. Leukemia and Lymphoma, 2014, 55, 957-958.	0.6	28
95	SOX11 and TP53 add prognostic information to MIPI in a homogenously treated cohort of mantle cell lymphoma $\hat{a} \in \mathbb{C}$ a Nordic Lymphoma Group study. British Journal of Haematology, 2014, 166, 98-108.	1.2	110
96	A Dual Program for Translation Regulation in Cellular Proliferation and Differentiation. Cell, 2014, 158, 1281-1292.	13.5	414
97	Circulating YKL-40 in patients with essential thrombocythemia and polycythemia vera treated with the novel histone deacetylase inhibitor vorinostat. Leukemia Research, 2014, 38, 816-821.	0.4	12
98	Predicting response to epigenetic therapy. Journal of Clinical Investigation, 2014, 124, 47-55.	3.9	78
99	Nordic MCL3 study: 90Y-ibritumomab-tiuxetan added to BEAM/C in non-CR patients before transplant in mantle cell lymphoma. Blood, 2014, 123, 2953-2959.	0.6	90
100	MicroRNA expression in early mycosis fungoides is distinctly different from atopic dermatitis and advanced cutaneous T-cell lymphoma. Anticancer Research, 2014, 34, 7207-17.	0.5	55
101	MicroRNAs in mantle cell lymphoma. Leukemia and Lymphoma, 2013, 54, 1867-1875.	0.6	11
102	Somatic mutations of the CREBBP and EP300 genes affect response to histone deacetylase inhibition in malignant DLBCL clones. Leukemia Research Reports, 2013, 2, 1-3.	0.2	30
103	Kinetics of del(7q) driven leukemogenesis in a patient with JAK2 V617F and TET2 mutated chronic myeloproliferative neoplasm. Leukemia Research Reports, 2013, 2, 51-53.	0.2	1
104	Downregulation but lack of promoter hypermethylation or somatic mutations of the potential tumor suppressor <i><scp>CXXC</scp>5</i> in <scp>MDS</scp> and <scp>AML</scp> with deletion 5q. European Journal of Haematology, 2013, 90, 259-260.	1.1	17
105	Analysis of Epigenetic Modifications of DNA in Human Cells. Current Protocols in Human Genetics, 2013, 77, Unit20.2.	3.5	9
106	Aberrant micro <scp>RNA</scp> expression in multiple myeloma. European Journal of Haematology, 2013, 91, 95-105.	1.1	40
107	Clonal expansion of renal cell carcinoma-infiltrating T lymphocytes. Oncolmmunology, 2013, 2, e26014.	2.1	20
108	Tubulointerstitial Nephritis in a Patient With Probable Autoimmune Lymphoproliferative Syndrome. Journal of Pediatric Hematology/Oncology, 2013, 35, e187-e189.	0.3	4

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109	Investigation of MGMT and DAPK1 methylation patterns in diffuse large B-cell lymphoma using allelic MSP-pyrosequencing. Scientific Reports, 2013, 3, 2789.	1.6	30
110	MicroRNA Profiling in Ocular Adnexal Lymphoma: A Role for MYC and NFKB1 Mediated Dysregulation of MicroRNA Expression in Aggressive Disease. , 2013, 54, 5169.		27
111	Genome-wide profiling identifies a DNA methylation signature that associates with TET2 mutations in diffuse large B-cell lymphoma. Haematologica, 2013, 98, 1912-1920.	1.7	116
112	Lenalidomide, Bendamustine, and Rituximab As First-Line Therapy For Patients >65 Years With Mantle Cell Lymphoma: Preliminary Results From The Nordic Lymphoma Group MCL4 (LENA-BERIT) Phase I-II Trial. Blood, 2013, 122, 4377-4377.	0.6	11
113	Diffuse Large B-Cell Lymphoma With Combined TP53 mutation and MIR34A methylation: Another "double hit―Lymphoma With Very Poor Outcome?. Blood, 2013, 122, 83-83.	0.6	9
114	A critical appraisal of tools available for monitoring epigenetic changes in clinical samples from patients with myeloid malignancies. Haematologica, 2012, 97, 1380-1388.	1.7	20
115	Allelic methylation levels of the noncoding VTRNA2-1 located on chromosome 5q31.1 predict outcome in AML. Blood, 2012, 119, 206-216.	0.6	97
116	Lack of somatic mutations in the catalytic domains of CREBBP and EP300 genes implies a role for histone deacetylase inhibition in myeloproliferative neoplasms. Leukemia Research, 2012, 36, 485-487.	0.4	6
117	miR34s in Normal and Malignant B-Cells: miR34A Plays a Dominant Role in Normal B-Cells, and aggressive Diffuse Large B-Cell Lymphoma Carry Combined Lesions of TP53, MIR34A, and MIR34B/C. Blood, 2012, 120, 296-296.	0.6	1
118	Nordic MCL3 Study: Zevalin Combined with High-Dose Chemotherapy Followed by Autologous Stem Cell Support As Late Intensification for Mantle Cell Lymphoma (MCL) Patients < 66 Years Not in CR After Induction Chemoimmunotherapy: No Benefit of Zevalin. Blood, 2012, 120, 747-747.	0.6	3
119	miRNA Profiling Predicts Survival and Identifies a Novel Putative Oncomir in Diffuse Large B-Cell Lymphoma Treated with Immunochemotherapy. Blood, 2012, 120, 1548-1548.	0.6	0
120	MicroRNA Profiling in Low-Grade Ocular MALT and Diffuse Large B-Cell Lymphoma: A Role for MYC and NFKB1 Mediated Dysregulation of MicroRNA Expression in Aggressive Disease. Blood, 2012, 120, 1600-1600.	0.6	0
121	Allelic Methylation Levels of VTRNA2-1 Predict Outcome in Higher Risk MDS Patients Not Treated by Azacytidine Blood, 2012, 120, 2394-2394.	0.6	0
122	Advances in DNA methylation: 5-hydroxymethylcytosine revisited. Clinica Chimica Acta, 2011, 412, 831-836.	0.5	93
123	MicroRNA-130a–mediated down-regulation of Smad4 contributes to reduced sensitivity to TGF-β1 stimulation in granulocytic precursors. Blood, 2011, 118, 6649-6659.	0.6	53
124	Clinical effect of increasing doses of lenalidomide in high-risk myelodysplastic syndrome and acute myeloid leukemia with chromosome 5 abnormalities. Haematologica, 2011, 96, 963-971.	1.7	52
125	miR-449 inhibits cell proliferation and is down-regulated in gastric cancer. Molecular Cancer, 2011, 10, 29.	7.9	206
126	Diagnostic microRNA profiling in cutaneous T-cell lymphoma (CTCL). Blood, 2011, 118, 5891-5900.	0.6	237

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127	Lenalidomide, Bendamustine, and Rituximab As First-Line Therapy for Patients > 65 Years with Mantle Cell Lymphoma: Results From the Phase I Portion of the Nordic Lymphoma Group MCL4 (LENA-BERIT) Trial. Blood, 2011, 118, 2700-2700.	0.6	12
128	Mir-130a-Mediated Downregulation of SMAD4 Contributes to Reduced Sensitivity to $TGF\hat{l}^2$ Stimulation in Promyelocytic Cells,. Blood, 2011, 118, 3383-3383.	0.6	0
129	Allelic Methylation Levels of the Non-Coding RNA Gene VTRNA2-1 Located on Chromosome 5q31.1 Predict Outcome in AML,. Blood, 2011, 118, 3450-3450.	0.6	O
130	TET2 mutations in Diffuse Large B-Cell Lymphoma: The Role of TET2 in the Regulation of Methylation Patterns at TET2 Target Genes. Blood, 2011, 118, 1364-1364.	0.6	0
131	Deficient SOCS3 and SHP-1 Expression in Psoriatic T Cells. Journal of Investigative Dermatology, 2010, 130, 1590-1597.	0.3	40
132	Cancer Risk in Long-term Users of Valproate: A Population-Based Case-Control Study. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 1714-1719.	1.1	28
133	Splenectomy in two children with autoimmune lymphoproliferative syndrome and massive splenomegaly. Pediatric Blood and Cancer, 2009, 53, 1124-1126.	0.8	1
134	Engaging the lysosomal compartment to combat B cell malignancies. Journal of Clinical Investigation, 2009, 119, 2133-6.	3.9	5
135	A novel splice mutation in theTP53 gene associated with Leydig cell tumor and primitive neuroectodermal tumor. Pediatric Blood and Cancer, 2008, 50, 701-703.	0.8	6
136	Epigenetic Changes in Cancer as Potential Targets for Prophylaxis and Maintenance Therapy. Basic and Clinical Pharmacology and Toxicology, 2008, 103, 389-396.	1.2	26
137	Frequent hypermethylation of DBC1 in malignant lymphoproliferative neoplasms. Modern Pathology, 2008, 21, 632-638.	2.9	22
138	Epigenetic changes in cancer. Apmis, 2007, 115, 1039-1059.	0.9	320
139	MicroRNAs and cancer. Apmis, 2007, 115, 1090-1106.	0.9	162
140	Genetic and epigenetic alterations of the reduced folate carrier in untreated diffuse large B-cell lymphoma. European Journal of Haematology, 2007, 80, 071119183417001-???.	1.1	13
141	Microarray-based classification of diffuse large B-cell lymphoma. European Journal of Haematology, 2005, 74, 453-465.	1.1	42
142	Genetic and epigenetic alterations of the APC gene in malignant melanoma. Oncogene, 2004, 23, 5215-5226.	2.6	105
143	Aberrations of the Chk2 tumour suppressor in advanced urinary bladder cancer. Oncogene, 2004, 23, 8545-8551.	2.6	42
144	Infrequent somatic Fas mutations but no evidence of Bcl 10 mutations or t(11 ; 18) in primary cutaneous MALT-type lymphoma. Journal of Pathology, 2003, 201, 134-140.	2.1	30

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145	ATM mutations are associated with inactivation of the ARF-TP53 tumor suppressor pathway in diffuse large B-cell lymphoma. Blood, 2002, 100, 1430-1437.	0.6	78
146	Profiling DNA methylation by melting analysis. Methods, 2002, 27, 121-127.	1.9	46
147	Mutational analysis of the tumour suppressor gene MMAC1/PTEN in malignant myeloid disorders. European Journal of Haematology, 2000, 65, 109-113.	1.1	64
148	Detection of mutations in GC-rich DNA by bisulphite denaturing gradient gel electrophoresis. Nucleic Acids Research, 1998, 26, 1548-1549.	6. 5	27