

Anke van den Berg

List of Publications by Year in descending order

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316
papers

13,489
citations

18482

62
h-index

31849

101
g-index

327
all docs

327
docs citations

327
times ranked

17948
citing authors

#	ARTICLE	IF	CITATIONS
1	BIC and miR-155 are highly expressed in Hodgkin, primary mediastinal and diffuse large B cell lymphomas. <i>Journal of Pathology</i> , 2005, 207, 243-249.	4.5	640
2	MHC class II transactivator CIITA is a recurrent gene fusion partner in lymphoid cancers. <i>Nature</i> , 2011, 471, 377-381.	27.8	551
3	High Expression of the CC Chemokine TARC in Reed-Sternberg Cells. <i>American Journal of Pathology</i> , 1999, 154, 1685-1691.	3.8	349
4	Somatic mutations of the von Hippel â€” Lindau disease tumour suppressor gene in non-familial clear cell renal carcinoma. <i>Human Molecular Genetics</i> , 1994, 3, 2169-2173.	2.9	341
5	Cigarette Smokeâ€”induced Emphysema. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2006, 173, 751-758.	5.6	279
6	MicroRNAs, macrocontrol: Regulation of miRNA processing. <i>Rna</i> , 2010, 16, 1087-1095.	3.5	229
7	High expression of Bâ€”cell receptor inducible gene <i>BIC</i> in all subtypes of Hodgkin lymphoma. <i>Genes Chromosomes and Cancer</i> , 2003, 37, 20-28.	2.8	224
8	Lack of <i>BIC</i> and microRNA miR-155 expression in primary cases of Burkitt lymphoma. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 147-153.	2.8	219
9	Cytoplasmic p21 expression levels determine cisplatin resistance in human testicular cancer. <i>Journal of Clinical Investigation</i> , 2010, 120, 3594-3605.	8.2	193
10	Circulating tumor cells in small-cell lung cancer: a predictive and prognostic factor. <i>Annals of Oncology</i> , 2012, 23, 2937-2942.	1.2	191
11	Expression of miR-21 and its targets (PTEN, PDCD4, TM1) in flat epithelial atypia of the breast in relation to ductal carcinoma in situ and invasive carcinoma. <i>BMC Cancer</i> , 2009, 9, 163.	2.6	190
12	Single-cell sequencing reveals karyotype heterogeneity in murine and human malignancies. <i>Genome Biology</i> , 2016, 17, 115.	8.8	178
13	A genome-wide association study of Hodgkin's lymphoma identifies new susceptibility loci at 2p16.1 (REL), 8q24.21 and 10p14 (GATA3). <i>Nature Genetics</i> , 2010, 42, 1126-1130.	21.4	177
14	Association with HLA class I in Epstein-Barr-virus-positive and with HLA class III in Epstein-Barr-virus-negative Hodgkin's lymphoma. <i>Lancet, The</i> , 2005, 365, 2216-2224.	13.7	155
15	Genome-Wide Association Study of Classical Hodgkin Lymphoma and Epsteinâ€”Barr Virus Statusâ€”Defined Subgroups. <i>Journal of the National Cancer Institute</i> , 2012, 104, 240-253.	6.3	141
16	Immunoâ€”miRs: critical regulators of Tâ€”cell development, function and ageing. <i>Immunology</i> , 2015, 144, 1-10.	4.4	141
17	Chemokines, cytokines and their receptors in Hodgkin's lymphoma cell lines and tissues. <i>Annals of Oncology</i> , 2002, 13, 52-56.	1.2	136
18	Follicular lymphoma grade 3B includes 3 cytogenetically defined subgroups with primary t(14;18), 3q27, or other translocations: t(14;18) and 3q27 are mutually exclusive. <i>Blood</i> , 2003, 101, 1149-1154.	1.4	136

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19	SETD2: an epigenetic modifier with tumor suppressor functionality. <i>Oncotarget</i> , 2016, 7, 50719-50734.	1.8	136
20	Isolation of the Human Semaphorin III/F Gene (SEMA3F) at Chromosome 3p21, a Region Deleted in Lung Cancer. <i>Genomics</i> , 1996, 32, 39-48.	2.9	134
21	Hodgkin Lymphoma Cell Lines Are Characterized by a Specific miRNA Expression Profile. <i>Neoplasia</i> , 2009, 11, 167-179.	5.3	133
22	HLA-A*02 is associated with a reduced risk and HLA-A*01 with an increased risk of developing EBV+ Hodgkin lymphoma. <i>Blood</i> , 2007, 110, 3310-3315.	1.4	131
23	Regulation of pri-microRNA BIC transcription and processing in Burkitt lymphoma. <i>Oncogene</i> , 2007, 26, 3769-3776.	5.9	131
24	Rapid Generation of MicroRNA Sponges for MicroRNA Inhibition. <i>PLoS ONE</i> , 2012, 7, e29275.	2.5	125
25	Gene expression profiling of microdissected Hodgkin Reed-Sternberg cells correlates with treatment outcome in classical Hodgkin lymphoma. <i>Blood</i> , 2012, 120, 3530-3540.	1.4	122
26	Plasma vesicle miRNAs for therapy response monitoring in Hodgkin lymphoma patients. <i>JCI Insight</i> , 2016, 1, e89631.	5.0	121
27	HLA Class II Expression by Hodgkin Reed-Sternberg Cells Is an Independent Prognostic Factor in Classical Hodgkin's Lymphoma. <i>Journal of Clinical Oncology</i> , 2007, 25, 3101-3108.	1.6	118
28	Aging disturbs the balance between effector and regulatory CD4+ T cells. <i>Experimental Gerontology</i> , 2014, 60, 190-196.	2.8	115
29	Proteomics analysis of Hodgkin lymphoma: identification of new players involved in the cross-talk between HRS cells and infiltrating lymphocytes. <i>Blood</i> , 2008, 111, 2339-2346.	1.4	114
30	Comprehensive analysis of miRNA expression in T-cell subsets of rheumatoid arthritis patients reveals defined signatures of naive and memory Tregs. <i>Genes and Immunity</i> , 2014, 15, 115-125.	4.1	111
31	Serum chemokine levels in Hodgkin lymphoma patients: highly increased levels of CCL17 and CCL22. <i>British Journal of Haematology</i> , 2008, 140, 527-536.	2.5	110
32	miRNA analysis in B-cell chronic lymphocytic leukaemia: proliferation centres characterized by low miR-150 and high miR-155 expression. <i>Journal of Pathology</i> , 2008, 215, 13-20.	4.5	109
33	A high throughput experimental approach to identify miRNA targets in human cells. <i>Nucleic Acids Research</i> , 2009, 37, e137-e137.	14.5	105
34	Randomized, Placebo-Controlled Phase III Study of Docetaxel Plus Carboplatin With Celecoxib and Cyclooxygenase-2 Expression As a Biomarker for Patients With Advanced Non-Small-Cell Lung Cancer: The NVALT-4 Study. <i>Journal of Clinical Oncology</i> , 2011, 29, 4320-4326.	1.6	105
35	miRNA profiling of B-cell subsets: specific miRNA profile for germinal center B cells with variation between centroblasts and centrocytes. <i>Laboratory Investigation</i> , 2009, 89, 708-716.	3.7	103
36	The microenvironment in classical Hodgkin lymphoma: An actively shaped and essential tumor component. <i>Seminars in Cancer Biology</i> , 2014, 24, 15-22.	9.6	102

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37	Generation of miRNA sponge constructs. <i>Methods</i> , 2012, 58, 113-117.	3.8	95
38	Strongly enhanced IL-10 production using stable galectin-1 homodimers. <i>Molecular Immunology</i> , 2007, 44, 506-513.	2.2	93
39	The role of microRNAs in normal hematopoiesis and hematopoietic malignancies. <i>Leukemia</i> , 2006, 20, 1931-1936.	7.2	92
40	Resistance mechanisms after tyrosine kinase inhibitors afatinib and crizotinib in non-small cell lung cancer, a review of the literature. <i>Critical Reviews in Oncology/Hematology</i> , 2016, 100, 107-116.	4.4	92
41	A gene in the chromosomal region 3p21 with greatly reduced expression in lung cancer is similar to the gene for ubiquitin-activating enzyme.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 6071-6075.	7.1	91
42	Dimeric galectin-1 induces IL-10 production in T-lymphocytes: an important tool in the regulation of the immune response. <i>Journal of Pathology</i> , 2004, 204, 511-518.	4.5	87
43	Interaction between host T cells and Reedâ€“Sternberg cells in Hodgkin lymphomas. <i>Seminars in Cancer Biology</i> , 2000, 10, 345-350.	9.6	84
44	HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?. <i>Oncimmunology</i> , 2017, 6, e1295202.	4.6	84
45	Latent Epstein-Barr Virus Infection of Tumor Cells in Classical Hodgkin's Lymphoma Predicts Adverse Outcome in Older Adult Patients. <i>Journal of Clinical Oncology</i> , 2009, 27, 3815-3821.	1.6	83
46	Clonal relation in a case of CLL, ALCL, and Hodgkin composite lymphoma. <i>Blood</i> , 2002, 100, 1425-1429.	1.4	79
47	Specific expression of miR-17-5p and miR-127 in testicular and central nervous system diffuse large B-cell lymphoma. <i>Modern Pathology</i> , 2009, 22, 547-555.	5.5	78
48	A meta-analysis of Hodgkin lymphoma reveals 19p13.3 TCF3 as a novel susceptibility locus. <i>Nature Communications</i> , 2014, 5, 3856.	12.8	78
49	HLAâ€“G protein expression as a potential immune escape mechanism in classical Hodgkinâ€“TMs lymphoma. <i>Tissue Antigens</i> , 2008, 71, 219-226.	1.0	76
50	Rheumatoid Arthritis, Immunosenescence and the Hallmarks of Aging. <i>Current Aging Science</i> , 2015, 8, 131-146.	1.2	76
51	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. <i>Leukemia</i> , 2014, 28, 2248-2251.	7.2	74
52	An Alternative Route for Multistep Tumorigenesis in a Novel Case of Hereditary Renal Cell Cancer and a t(2;3)(q35;q21) Chromosome Translocation. <i>American Journal of Human Genetics</i> , 1998, 62, 1475-1483.	6.2	73
53	MiRNA profiling in B nonâ€“Hodgkin lymphoma: a <i>c-MYC</i> -related miRNA profile characterizes Burkitt lymphoma. <i>British Journal of Haematology</i> , 2010, 149, 896-899.	2.5	71
54	Involvement of multiple loci on chromosome 3 in renal cell cancer development. , 1997, 19, 59-76.		69

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55	Mitoxantrone resistance in a small cell lung cancer cell line is associated with ABCA2 upregulation. <i>British Journal of Cancer</i> , 2004, 90, 2411-2417.	6.4	69
56	Treatment of the bronchial tree from beginning to end: targeting small airway inflammation in asthma. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2013, 68, 16-26.	5.7	69
57	Long noncoding RNAs as a novel component of the Myc transcriptional network. <i>FASEB Journal</i> , 2015, 29, 2338-2346.	0.5	67
58	The European Hematology Association Roadmap for European Hematology Research: a consensus document. <i>Haematologica</i> , 2016, 101, 115-208.	3.5	67
59	Common and differential chemokine expression patterns in rs cells of NLP, EBV positive and negative classical hodgkin lymphomas. <i>International Journal of Cancer</i> , 2002, 99, 665-672.	5.1	66
60	Development of Lymphoma in Autoimmune Lymphoproliferative Syndrome (ALPS) and its Relationship to Fas Gene Mutations. <i>Leukemia and Lymphoma</i> , 2004, 45, 423-431.	1.3	66
61	A homozygous deletion in a small cell lung cancer cell line involving a 3p21 region with a marked instability in yeast artificial chromosomes. <i>Cancer Research</i> , 1994, 54, 4183-7.	0.9	66
62	Molecular, cytogenetic, and immunophenotypic characterization of follicular lymphoma grade 3B; a separate entity or part of the spectrum of diffuse large B-cell lymphoma or follicular lymphoma?. <i>Human Pathology</i> , 2006, 37, 528-533.	2.0	65
63	Protease activity of plasma hemopexin. <i>Kidney International</i> , 2005, 68, 603-610.	5.2	63
64	Differential expression and distribution of epithelial adhesion molecules in non-small cell lung cancer and normal bronchus. <i>Journal of Clinical Pathology</i> , 2007, 60, 608-614.	2.0	63
65	Current smoking-specific gene expression signature in normal bronchial epithelium is enhanced in squamous cell lung cancer. <i>Journal of Pathology</i> , 2009, 218, 182-191.	4.5	63
66	The CD4+CD26 ^{hi} T-cell population in classical Hodgkin's lymphoma displays a distinctive regulatory T-cell profile. <i>Laboratory Investigation</i> , 2008, 88, 482-490.	3.7	62
67	Dichotomous ALK-IHC Is a Better Predictor for ALK Inhibition Outcome than Traditional ALK-FISH in Advanced Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 4251-4258.	7.0	62
68	Dual Role of miR-21 in CD4+ T-Cells: Activation-Induced miR-21 Supports Survival of Memory T-Cells and Regulates CCR7 Expression in Naive T-Cells. <i>PLoS ONE</i> , 2013, 8, e76217.	2.5	61
69	A cosmid and cDNA fine physical map of a human chromosome 13q14 region frequently lost in B-cell chronic lymphocytic leukemia and identification of a new putative tumor suppressor gene, Leu5. <i>FEBS Letters</i> , 1998, 426, 266-270.	2.8	59
70	Genomic aberrations in squamous cell lung carcinoma related to lymph node or distant metastasis. <i>Lung Cancer</i> , 2009, 66, 372-378.	2.0	57
71	Expression of the T-Cell Transcription Factors, GATA-3 and T-bet, in the Neoplastic Cells of Hodgkin Lymphomas. <i>American Journal of Pathology</i> , 2005, 166, 127-134.	3.8	56
72	Plasma thymus and activation-regulated chemokine as an early response marker in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012, 97, 410-415.	3.5	56

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73	TP53 gene mutations in Hodgkin lymphoma are infrequent and not associated with absence of Epstein-Barr virus. <i>International Journal of Cancer</i> , 2001, 94, 60-66.	5.1	54
74	Low Frequency of FAS Mutations in Reed-Sternberg Cells of Hodgkin's Lymphoma. <i>American Journal of Pathology</i> , 2003, 162, 29-35.	3.8	54
75	Toll-like receptors in the pathogenesis of human B cell malignancies. <i>Journal of Hematology and Oncology</i> , 2014, 7, 57.	17.0	54
76	Major role for a 3p21 region and lack of involvement of the t(3;8) breakpoint region in the development of renal cell carcinoma suggested by loss of heterozygosity analysis. , 1996, 15, 64-72.		52
77	HLA Associations in Classical Hodgkin Lymphoma: EBV Status Matters. <i>PLoS ONE</i> , 2012, 7, e39986.	2.5	52
78	Genetic Associations in Classical Hodgkin Lymphoma: A Systematic Review and Insights into Susceptibility Mechanisms. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2737-2747.	2.5	52
79	Extensive mutation scanning of RET in sporadic medullary thyroid carcinoma and of RET and VHL in sporadic pheochromocytoma reveals involvement of these genes in only a minority of cases. <i>Journal of Clinical Endocrinology and Metabolism</i> , 1996, 81, 2881-2884.	3.6	49
80	The role of female sex hormones in the development and severity of allergic and non-allergic asthma. <i>Clinical and Experimental Allergy</i> , 2009, 39, 1477-1481.	2.9	48
81	Prognostic Model to Predict Post-Autologous Stem-Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. <i>Journal of Clinical Oncology</i> , 2017, 35, 3722-3733.	1.6	48
82	Combining genomewide association study and lung eQTL analysis provides evidence for novel genes associated with asthma. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2016, 71, 1712-1720.	5.7	47
83	Building bridges for innovation in ageing: Synergies between action groups of the EIP on AHA. <i>Journal of Nutrition, Health and Aging</i> , 2017, 21, 92-104.	3.3	47
84	miR-24-3p Is Overexpressed in Hodgkin Lymphoma and Protects Hodgkin and Reed-Sternberg Cells from Apoptosis. <i>American Journal of Pathology</i> , 2017, 187, 1343-1355.	3.8	46
85	A comparison of genomic structures and expression patterns of two closely related flanking genes in a critical lung cancer region at 3p21.3. <i>European Journal of Human Genetics</i> , 1999, 7, 478-486.	2.8	45
86	Intricate crosstalk between MYC and non-coding RNA's regulates hallmarks of cancer. <i>Molecular Oncology</i> , 2019, 13, 26-45.	4.6	45
87	Immune reactions in classical Hodgkin's lymphoma. <i>Seminars in Hematology</i> , 1999, 36, 253-9.	3.4	45
88	Genetic susceptibility to Hodgkin's lymphoma associated with the human leukocyte antigen region. <i>European Journal of Haematology</i> , 2005, 75, 34-41.	2.2	44
89	Analysis of serum immune markers in seropositive and seronegative rheumatoid arthritis and in high-risk seropositive arthralgia patients. <i>Scientific Reports</i> , 2016, 6, 26021.	3.3	44
90	Biomarkers for evaluation of treatment response in classical Hodgkin lymphoma: comparison of sGClectin-1, sCD163 and sCD30 with TARC. <i>British Journal of Haematology</i> , 2016, 175, 868-875.	2.5	44

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91	Non-Coding RNAs in Cancer Radiosensitivity: MicroRNAs and lncRNAs as Regulators of Radiation-Induced Signaling Pathways. <i>Cancers</i> , 2020, 12, 1662.	3.7	44
92	Involvement of multiple loci on chromosome 3 in renal cell cancer development. <i>Genes Chromosomes and Cancer</i> , 1997, 19, 59-76.	2.8	44
93	MicroRNA profiling of human primary macrophages exposed to dengue virus identifies miRNA-3614-5p as antiviral and regulator of ADAR1 expression. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005981.	3.0	43
94	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. <i>Oncotarget</i> , 2016, 7, 2391-2400.	1.8	43
95	A High Throughput Experimental Approach to Identify miRNA Target Genes in Hodgkin Lymphoma.. <i>Blood</i> , 2008, 112, 1461-1461.	1.4	42
96	Analysis of multiple renal cell adenomas and carcinomas suggests allelic loss at 3p21 to be a prerequisite for malignant development. , 1997, 19, 228-232.		41
97	Splenic Marginal Zone Lymphomas Presenting with Splenomegaly and Typical Immunophenotype Are Characterized by Allelic Loss in 7q31-32. <i>Modern Pathology</i> , 2003, 16, 1210-1217.	5.5	41
98	Cytokine gene expression profile distinguishes CD4+/CD57+ T cells of the nodular lymphocyte predominance type of Hodgkin's lymphoma from their tonsillar counterparts. <i>Journal of Pathology</i> , 2006, 208, 423-430.	4.5	41
99	Rapid BRAF mutation tests in patients with advanced melanoma: comparison of immunohistochemistry, Droplet Digital PCR, and the Idylla Mutation Platform. <i>Melanoma Research</i> , 2018, 28, 96-104.	1.2	41
100	The Microenvironment in Epstein-Barr Virus-Associated Malignancies. <i>Pathogens</i> , 2018, 7, 40.	2.8	40
101	CD58 mutations are common in Hodgkin lymphoma cell lines and loss of CD58 expression in tumor cells occurs in Hodgkin lymphoma patients who relapse. <i>Genes and Immunity</i> , 2016, 17, 363-366.	4.1	39
102	ZDHHC11 and ZDHHC11B are critical novel components of the oncogenic MYC-miR-150-MYB network in Burkitt lymphoma. <i>Leukemia</i> , 2017, 31, 1470-1473.	7.2	39
103	Mutations in EMT-Related Genes in ALK Positive Crizotinib Resistant Non-Small Cell Lung Cancers. <i>Cancers</i> , 2018, 10, 10.	3.7	39
104	Partial 3q duplication syndrome and assignment of D3S5 to 3q25-3q28. <i>Human Genetics</i> , 1991, 87, 151-154.	3.8	38
105	NormalFHIT transcripts in renal cell cancer- and lung cancer-derived cell lines, including a cell line with a homozygous deletion in the FRA3B region. , 1997, 19, 220-227.		38
106	An 80 Kb P1 clone from chromosome 3p21.3 suppresses tumor growth in vivo. <i>Oncogene</i> , 1996, 13, 2387-96.	5.9	38
107	Induction of glomerular alkaline phosphatase after challenge with lipopolysaccharide. <i>International Journal of Experimental Pathology</i> , 2003, 84, 135-144.	1.3	37
108	Emerging roles for long noncoding RNAs in B-cell development and malignancy. <i>Critical Reviews in Oncology/Hematology</i> , 2017, 120, 77-85.	4.4	37

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109	Combined osimertinib, dabrafenib and trametinib treatment for advanced non-small-cell lung cancer patients with an osimertinib-induced BRAF V600E mutation. <i>Lung Cancer</i> , 2020, 146, 358-361.	2.0	37
110	T-cell Activation Induces Dynamic Changes in miRNA Expression Patterns in CD4 and CD8 T-cell Subsets. <i>MicroRNA (Sharjah, United Arab Emirates)</i> , 2015, 4, 117-122.	1.2	37
111	The Human Leukocyte Antigen Class I Region Is Associated with EBV-Positive Hodgkin's Lymphoma: HLA-A and HLA Complex Group 9 Are Putative Candidate Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 2280-2284.	2.5	36
112	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cell-Specific Long Noncoding RNAs. <i>American Journal of Pathology</i> , 2016, 186, 2462-2472.	3.8	36
113	High expression of calcium-binding proteins, S100A10, S100A11 and CALM2 in anaplastic large cell lymphoma. <i>British Journal of Haematology</i> , 2005, 131, 596-608.	2.5	35
114	Brain Death Induces Inflammation in the Donor Intestine. <i>Transplantation</i> , 2008, 86, 148-154.	1.0	35
115	MiR-17/106b seed family regulates p21 in Hodgkin's lymphoma. <i>Journal of Pathology</i> , 2011, 225, 609-617.	4.5	35
116	A gene from human chromosome region 3p21 with reduced expression in small cell lung cancer. <i>Cancer Research</i> , 1992, 52, 1536-41.	0.9	35
117	Germline FAS gene mutation in a case of ALPS and NLP Hodgkin lymphoma. <i>Blood</i> , 2002, 99, 1492-1494.	1.4	34
118	P53 Mutation Analysis of Colorectal Liver Metastases: Relation to Actual Survival, Angiogenic Status, and p53 Overexpression. <i>Clinical Cancer Research</i> , 2005, 11, 4067-4073.	7.0	34
119	Age-related gene and miRNA expression changes in airways of healthy individuals. <i>Scientific Reports</i> , 2019, 9, 3765.	3.3	34
120	Identification of transforming growth factor-beta-regulated microRNAs and the microRNA-targetomes in primary lung fibroblasts. <i>PLoS ONE</i> , 2017, 12, e0183815.	2.5	34
121	Direct molecular analysis of a deletion of 3p in tumors from patients with sporadic renal cell carcinoma. <i>Cancer Genetics and Cytogenetics</i> , 1988, 32, 281-285.	1.0	33
122	Maternal smoking during pregnancy decreases Wnt signalling in neonatal mice. <i>Thorax</i> , 2010, 65, 553-554.	5.6	33
123	Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma. <i>Cancer Treatment Reviews</i> , 2020, 82, 101931.	7.7	33
124	Analysis of Released Circulating Tumor Cells During Surgery for Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 1656-1666.	7.0	33
125	Analysis of chromosomal copy number changes and oncoprotein expression in primary central nervous system lymphomas: frequent loss of chromosome arm 6q. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2003, 443, 164-169.	2.8	32
126	Relevance and Effectiveness of Molecular Tumor Board Recommendations for Patients With Non-Small-Cell Lung Cancer With Rare or Complex Mutational Profiles. <i>JCO Precision Oncology</i> , 2020, 4, 393-410.	3.0	32

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127	Production of hemopexin by TNF- α stimulated human mesangial cells. <i>Kidney International</i> , 2003, 63, 1681-1686.	5.2	31
128	BCL6 alternative translocation breakpoint cluster region associated with follicular lymphoma grade 3B. <i>Genes Chromosomes and Cancer</i> , 2005, 44, 301-304.	2.8	31
129	Prolonged protection of the new inhaled corticosteroid fluticasone furoate against AMP hyperresponsiveness in patients with asthma. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2010, 65, 1531-1535.	5.7	31
130	Budesonide and fluticasone propionate differentially affect the airway epithelial barrier. <i>Respiratory Research</i> , 2016, 17, 2.	3.6	30
131	Treatment of patients with MYC rearrangement positive large B-cell lymphoma with R-CHOP plus lenalidomide: results of a multicenter HOVON phase II trial. <i>Haematologica</i> , 2020, 105, 2805-2812.	3.5	30
132	BCL6 alternative breakpoint region break and homozygous deletion of 17q24 in the nodular lymphocyte predominance type of Hodgkin's lymphoma-derived cell line DEV. <i>Human Pathology</i> , 2006, 37, 675-683.	2.0	29
133	Expression of the c-Met oncogene by tumor cells predicts a favorable outcome in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012, 97, 572-578.	3.5	29
134	SF Treg cells transcribing high levels of Bcl-2 and microRNA-21 demonstrate limited apoptosis in RA. <i>Rheumatology</i> , 2015, 54, 950-958.	1.9	29
135	Mutation patterns in small cell and non-small cell lung cancer patients suggest a different level of heterogeneity between primary and metastatic tumors. <i>Carcinogenesis</i> , 2017, 38, bgw128.	2.8	29
136	The microenvironment of classical Hodgkin lymphoma: heterogeneity by Epstein-Barr virus presence and location within the tumor. <i>Blood Cancer Journal</i> , 2016, 6, e417-e417.	6.2	29
137	Identification of chromosomal copy number changes associated with transformation of follicular lymphoma to diffuse large B-cell lymphoma. <i>Human Pathology</i> , 2003, 34, 915-923.	2.0	28
138	KRAS Mutation as a Resistance Mechanism to BRAF/MEK Inhibition in NSCLC. <i>Journal of Thoracic Oncology</i> , 2018, 13, e249-e251.	1.1	28
139	Rosetting T cells in Hodgkin lymphoma are activated by immunological synapse components HLA class II and CD58. <i>Blood</i> , 2020, 136, 2437-2441.	1.4	28
140	Epidemiology of Classical Hodgkin Lymphoma and Its Association with Epstein Barr Virus in Northern China. <i>PLoS ONE</i> , 2011, 6, e21152.	2.5	28
141	Gene expression analysis of dendritic/Langerhans cells and Langerhans cell histiocytosis. <i>Journal of Pathology</i> , 2006, 209, 474-483.	4.5	27
142	A chronic obstructive pulmonary disease related signature in squamous cell lung cancer. <i>Lung Cancer</i> , 2011, 72, 177-183.	2.0	26
143	The entire miR-200 seed family is strongly deregulated in clear cell renal cell cancer compared to the proximal tubular epithelial cells of the kidney. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 165-173.	2.8	26
144	HLA-A*02:07 Is a Protective Allele for EBV Negative and a Susceptibility Allele for EBV Positive Classical Hodgkin Lymphoma in China. <i>PLoS ONE</i> , 2012, 7, e31865.	2.5	25

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145	Brain death causes structural and inflammatory changes in donor intestine. <i>Transplantation Proceedings</i> , 2005, 37, 448-449.	0.6	24
146	Global correlation of genome and transcriptome changes in classical Hodgkin lymphoma. <i>Hematological Oncology</i> , 2007, 25, 21-29.	1.7	24
147	Expression of CD1d and presence of invariant NKT cells in classical Hodgkin lymphoma. <i>American Journal of Hematology</i> , 2010, 85, 539-541.	4.1	24
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