

Anke van den Berg

List of Publications by Year in descending order

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Version: 2024-02-01

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	Actionability of on-target ALK Resistance Mutations in Patients With Non-Small Cell Lung Cancer: Local Experience and Review of the Literature. <i>Clinical Lung Cancer</i> , 2022, 23, e104-e115.	2.6	13
2	Cell-of-origin classification using the Hans and Lymph2Cx algorithms in primary cutaneous large B-cell lymphomas. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2022, 480, 667-675.	2.8	12
3	The Role of the MYC/miR-150/MYB/ZDHHC11 Network in Hodgkin Lymphoma and Diffuse Large B-Cell Lymphoma. <i>Genes</i> , 2022, 13, 227.	2.4	3
4	Understanding the role of long-acting muscarinic antagonists in asthma treatment. <i>Annals of Allergy, Asthma and Immunology</i> , 2022, 128, 352-360.	1.0	7
5	CD4+ T cells in classical Hodgkin lymphoma express exhaustion associated transcription factors TOX and TOX2. <i>Oncolmmunology</i> , 2022, 11, 2033433.	4.6	9
6	Killer Cell Immunoglobulin-Like Receptor Haplotype B Modulates Susceptibility to EBV-Associated Classic Hodgkin Lymphoma. <i>Frontiers in Immunology</i> , 2022, 13, 829943.	4.8	4
7	Alpine altitude climate treatment for severe and uncontrolled asthma: An EAACI position paper. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2022, 77, 1991-2024.	5.7	21
8	The Microbiome in Bronchial Biopsies from Smokers and Ex-Smokers with Stable COPD - A Metatranscriptomic Approach. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2022, 19, 81-87.	1.6	1
9	Low Mutational Burden of Extranodal Marginal Zone Lymphoma of Mucosa-Associated Lymphoid Tissue in Patients with Primary Sjogren's Syndrome. <i>Cancers</i> , 2022, 14, 1010.	3.7	5
10	The relation between age and airway epithelial barrier function. <i>Respiratory Research</i> , 2022, 23, 43.	3.6	13
11	Identification of the estrogen receptor beta as a possible new tamoxifen-sensitive target in diffuse large B-cell lymphoma. <i>Blood Cancer Journal</i> , 2022, 12, 36.	6.2	8
12	Reproducibility of Gene Expression Signatures in Diffuse Large B-Cell Lymphoma. <i>Cancers</i> , 2022, 14, 1346.	3.7	1
13	Biological and Clinical Implications of Gene-Expression Profiling in Diffuse Large B-Cell Lymphoma: A Proposal for a Targeted BLYM-777 Consortium Panel as Part of a Multilayered Analytical Approach. <i>Cancers</i> , 2022, 14, 1857.	3.7	6
14	Airway Wall Splice-QTL Analysis Reveals Novel Downstream Mechanisms for Well-Known GWAS Asthma-SNPs. , 2022, , .		0
15	Frequent 4EBP1 Amplification Induces Synthetic Dependence on FGFR Signaling in Cancer. <i>Cancers</i> , 2022, 14, 2397.	3.7	1
16	Prototype ORACLE Score Validation in NOVELTY: Predicted versus Observed Asthma Exacerbation Rates. , 2022, , .		0
17	Cytochrome P450 27C1 Level Dictates Lung Cancer Tumorigenicity and Sensitivity towards Multiple Anticancer Agents and Its Potential Interplay with the IGF-1R/Akt/p53 Signaling Pathway. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7853.	4.1	4
18	Interaction between ERAP Alleles and HLA Class I Types Support a Role of Antigen Presentation in Hodgkin Lymphoma Development. <i>Cancers</i> , 2021, 13, 414.	3.7	6

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19	Soluble PD-L1 is a promising disease biomarker but does not reflect tissue expression in classic Hodgkin lymphoma. <i>British Journal of Haematology</i> , 2021, 193, 506-514.	2.5	9
20	Clinical Value of EGFR Copy Number Gain Determined by Amplicon-Based Targeted Next Generation Sequencing in Patients with EGFR-Mutated NSCLC. <i>Targeted Oncology</i> , 2021, 16, 215-226.	3.6	7
21	Ultra-low-dose CT combined with noise reduction techniques for quantification of emphysema in COPD patients: An intra-individual comparison study with standard-dose CT. <i>European Journal of Radiology</i> , 2021, 138, 109646.	2.6	19
22	Frequent mutated <i>B2M</i> , <i>EZH2</i> , <i>IRF8</i> , and <i>TNFRSF14</i> in primary bone diffuse large B-cell lymphoma reflect a GCB phenotype. <i>Blood Advances</i> , 2021, 5, 3760-3775.	5.2	11
23	HLA Expression in Relation to HLA Type in Classic Hodgkin Lymphoma Patients. <i>Cancers</i> , 2021, 13, 5833.	3.7	1
24	Low Mutational Burden of Extra Nodal Marginal Zone Lymphoma of Mucosa-Associated Lymphoid Tissue in Patients with Primary Sjogren's Syndrome. <i>Blood</i> , 2021, 138, 1331-1331.	1.4	1
25	Responsivity and Reproducibility of Sputum Inflammatory Biomarkers During COPD Exacerbation and Stable Phases – A Pilot Study. <i>International Journal of COPD</i> , 2021, Volume 16, 3055-3064.	2.3	1
26	Genetic Modification Approaches for Parasporins <i>Bacillus thuringiensis</i> Proteins with Anticancer Activity. <i>Molecules</i> , 2021, 26, 7476.	3.8	1
27	Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma. <i>Cancer Treatment Reviews</i> , 2020, 82, 101931.	7.7	33
28	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
29	An All-In-One Transcriptome-Based Assay to Identify Therapy-Guiding Genomic Aberrations in Nonsmall Cell Lung Cancer Patients. <i>Cancers</i> , 2020, 12, 2843.	3.7	6
30	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
31	MiR-378a-3p Is Critical for Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020, 12, 3546.	3.7	12
32	MicroRNA Profiling in Bronchial Biopsies of Asthma Patients. , 2020, , .		0
33	Molecular imaging in lymphoma beyond 18F-FDG-PET: understanding the biology and its implications for diagnostics and therapy. <i>Lancet Haematology</i> , 2020, 7, e479-e489.	4.6	14
34	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
35	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020, 12, 1464.	3.7	19
36	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

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37	WEE1 inhibition synergizes with CHOP chemotherapy and radiation therapy through induction of premature mitotic entry and DNA damage in diffuse large B-cell lymphoma. <i>Therapeutic Advances in Hematology</i> , 2020, 11, 204062071989837.	2.5	12
38	Interim thymus and activation regulated chemokine versus interim 18 Fâ€fluorodeoxyglucose positronâ€emission tomography in classical Hodgkin lymphoma response evaluation. <i>British Journal of Haematology</i> , 2020, 190, 40-44.	2.5	15
39	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
40	Relevance and Effectiveness of Molecular Tumor Board Recommendations for Patients With Nonâ€Small-Cell Lung Cancer With Rare or Complex Mutational Profiles. <i>JCO Precision Oncology</i> , 2020, 4, 393-410.	3.0	32
41	Cell-free DNA as biomarker in Hodgkin lymphoma patients. <i>Klinische Padiatrie</i> , 2020, 232, .	0.6	0
42	Microenvironment, Cross-Talk, and Immune Escape Mechanisms. <i>Hematologic Malignancies</i> , 2020, , 69-86.	0.2	1
43	Cellular Senescence in Lung Fibroblasts from COPD Patients Is Associated with Altered Extracellular Matrix Regulation. , 2019, , .		0
44	Long Non-coding RNAs in the Development and Maintenance of Lymphoid Malignancies. , 2019, , 127-149.		0
45	A super-SILAC based proteomics analysis of diffuse large B-cell lymphoma-NOS patient samples to identify new proteins that discriminate GCB and non-GCB lymphomas. <i>PLoS ONE</i> , 2019, 14, e0223260.	2.5	4
46	WEE1 Inhibition Enhances Anti-Apoptotic Dependency as a Result of Premature Mitotic Entry and DNA Damage. <i>Cancers</i> , 2019, 11, 1743.	3.7	12
47	Current Smoking is Associated with Decreased Expression of miR-335-5p in Parenchymal Lung Fibroblasts. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5176.	4.1	15
48	Targeted sequencing of circulating cell-free DNA in stage II-III resectable oesophageal squamous cell carcinoma patients. <i>BMC Cancer</i> , 2019, 19, 818.	2.6	16
49	Increased miR-142-3p Expression Might Explain Reduced Regulatory T Cell Function in Granulomatosis With Polyangiitis. <i>Frontiers in Immunology</i> , 2019, 10, 2170.	4.8	18
50	Tumour necrosis as assessed with 18F-FDG PET is a potential prognostic marker in diffuse large B cell lymphoma independent of MYC rearrangements. <i>European Radiology</i> , 2019, 29, 6018-6028.	4.5	6
51	Recommendations for the clinical interpretation and reporting of copy number gains using gene panel NGS analysis in routine diagnostics. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2019, 474, 673-680.	2.8	24
52	Age-related gene and miRNA expression changes in airways of healthy individuals. <i>Scientific Reports</i> , 2019, 9, 3765.	3.3	34
53	CX Chemokine Receptor 7 Contributes to Survival of KRAS-Mutant Non-Small Cell Lung Cancer upon Loss of Epidermal Growth Factor Receptor. <i>Cancers</i> , 2019, 11, 455.	3.7	18
54	NGS-Based High-Throughput Screen to Identify MicroRNAs Regulating Growth of B-Cell Lymphoma. <i>Methods in Molecular Biology</i> , 2019, 1956, 269-282.	0.9	3

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55	Small Airways Disease Parameters Defined by PExA Particle Mass in Asthma, Asthma Remission and Healthy Controls. , 2019, , .		0
56	Marked TGF- β -regulated miRNA expression changes in both COPD and control lung fibroblasts. Scientific Reports, 2019, 9, 18214.	3.3	16
57	Heterogeneous Pattern of Dependence on Anti-Apoptotic BCL-2 Family Proteins upon CHOP Treatment in Diffuse Large B-Cell Lymphoma. International Journal of Molecular Sciences, 2019, 20, 6036.	4.1	13
58	Intricate crosstalk between MYC and non-coding RNA regulates hallmarks of cancer. Molecular Oncology, 2019, 13, 26-45.	4.6	45
59	Argonaute 2 RNA Immunoprecipitation Reveals Distinct miRNA Targetomes of Primary Burkitt Lymphoma Tumors and Normal B Cells. American Journal of Pathology, 2018, 188, 1289-1299.	3.8	7
60	Effects of fluticasone propionate and budesonide on the expression of immune defense genes in bronchial epithelial cells. Pulmonary Pharmacology and Therapeutics, 2018, 50, 47-56.	2.6	19
61	Combined PD-1 and JAK1/2 inhibition in refractory primary mediastinal B-cell lymphoma. Annals of Hematology, 2018, 97, 905-907.	1.8	3
62	Rapid BRAF mutation tests in patients with advanced melanoma: comparison of immunohistochemistry, Droplet Digital PCR, and the Idylla Mutation Platform. Melanoma Research, 2018, 28, 96-104.	1.2	41
63	Targeting the Microenvironment in Hodgkin Lymphoma: Opportunities and Challenges. Molecular Pathology Library, 2018, , 59-90.	0.1	0
64	Identification of Two Protein-Signaling States Delineating Transcriptionally Heterogeneous Human Medulloblastoma. Cell Reports, 2018, 22, 3206-3216.	6.4	19
65	Combined loss of HLA I and HLA II expression is more common in the non-GCB type of diffuse large B cell lymphoma. Histopathology, 2018, 72, 886-888.	2.9	4
66	Circulating miRNAs in patients with Barrett's esophagus, high-grade dysplasia and esophageal adenocarcinoma. Journal of Gastrointestinal Oncology, 2018, 9, 1150-1156.	1.4	11
67	KRAS Mutation as a Resistance Mechanism to BRAF/MEK Inhibition in NSCLC. Journal of Thoracic Oncology, 2018, 13, e249-e251.	1.1	28
68	Differential miRNA Expression Profiles in Cumulus and Mural Granulosa Cells from Human Pre-ovulatory Follicles. MicroRNA (Sharjah, United Arab Emirates), 2018, 8, 61-67.	1.2	23
69	Mutational Evolution in Relapsed Diffuse Large B-Cell Lymphoma. Cancers, 2018, 10, 459.	3.7	16
70	Tuberous sclerosis complex is required for tumor maintenance in MYC-driven Burkitt's lymphoma. EMBO Journal, 2018, 37, .	7.8	10
71	MicroRNA High Throughput Loss-of-Function Screening Reveals an Oncogenic Role for miR-21-5p in Hodgkin Lymphoma. Cellular Physiology and Biochemistry, 2018, 49, 144-159.	1.6	20
72	Serum periostin does not reflect type 2-driven inflammation in COPD. Respiratory Research, 2018, 19, 112.	3.6	8

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73	Novel genes and insights in complete asthma remission: A genome-wide association study on clinical and complete asthma remission. <i>Clinical and Experimental Allergy</i> , 2018, 48, 1286-1296.	2.9	17
74	MBRS-36. IDENTIFICATION OF TWO PROTEIN-SIGNALING STATES DELINEATING TRANSCRIPTIONALLY HETEROGENEOUS HUMAN MEDULLOBLASTOMA. <i>Neuro-Oncology</i> , 2018, 20, i136-i136.	1.2	0
75	Impact of acute exposure to cigarette smoke on airway gene expression. <i>Physiological Genomics</i> , 2018, 50, 705-713.	2.3	24
76	Involvement of MicroRNAs in the Aging-Related Decline of CD28 Expression by Human T Cells. <i>Frontiers in Immunology</i> , 2018, 9, 1400.	4.8	13
77	Mutations in EMT-Related Genes in ALK Positive Crizotinib Resistant Non-Small Cell Lung Cancers. <i>Cancers</i> , 2018, 10, 10.	3.7	39
78	The Microenvironment in Epstein-Barr Virus-Associated Malignancies. <i>Pathogens</i> , 2018, 7, 40.	2.8	40
79	Identification of relevant drugable targets in diffuse large B-cell lymphoma using a genome-wide unbiased CD20 guilt-by association approach. <i>PLoS ONE</i> , 2018, 13, e0193098.	2.5	20
80	OPO316...Increased expression of microrna-142-3p is associated with the functional defect of regulatory t cells in anti-neutrophil cytoplasmic antibody associated vasculitis. , 2018, , .		0
81	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
82	Argonaute 2 immunoprecipitation revealed large tumor suppressor kinase 1 as a novel proapoptotic target of miR-21 in T cells. <i>FEBS Journal</i> , 2017, 284, 555-567.	4.7	7
83	Latrophilin receptors: novel bronchodilator targets in asthma. <i>Thorax</i> , 2017, 72, 74-82.	5.6	12
84	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
85	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
86	Copy number alterations assessed at the single-cell level revealed mono- and polyclonal seeding patterns of distant metastasis in a small-cell lung cancer patient. <i>Annals of Oncology</i> , 2017, 28, 1668-1670.	1.2	19
87	HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?. <i>Oncolmmunology</i> , 2017, 6, e1295202.	4.6	84
88	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
89	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
90	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

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91	<i>PTTG1IP</i> and <i>MAML3</i> , novel genomewide association study genes for severity of hyperresponsiveness in adult asthma. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2017, 72, 792-801.	5.7	12
92	All-in-one RNA-based assay to detect therapeutic biomarkers in lung cancer. <i>Annals of Oncology</i> , 2017, 28, vii10.	1.2	0
93	Lymphadenopathy driven by TCR-V β 1 T-cell expansion in FAS-related autoimmune lymphoproliferative syndrome. <i>Blood Advances</i> , 2017, 1, 1101-1106.	5.2	3
94	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
95	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
96	HLA expression and HLA type associations in relation to EBV status in Hispanic Hodgkin lymphoma patients. <i>PLoS ONE</i> , 2017, 12, e0174457.	2.5	7
97	Overall survival in EGFR mutated non-small-cell lung cancer patients treated with afatinib after EGFR TKI and resistant mechanisms upon disease progression. <i>PLoS ONE</i> , 2017, 12, e0182885.	2.5	21
98	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
99	Abstract 754: Treatment decision-making of rare ERBB2 (HER2) mutations in lung cancer; a role for multidisciplinary molecular tumor boards. , 2017, , .		0
100	Abstract 2718: Molecular Tumor Board treatment predictions on rare EGFR exon 20 mutations. , 2017, , .		1
101	Abstract 2785: A comprehensive RNA-based assay for treatment prediction in non-small cell lung cancer patients. , 2017, , .		0
102	Target gene identification of TGF- β -induced miR-455-3p and miR-21-3p in lung fibroblasts. , 2017, , .		0
103	Characterization of the Microenvironment of Nodular Lymphocyte Predominant Hodgkin Lymphoma. <i>International Journal of Molecular Sciences</i> , 2016, 17, 2127.	4.1	23
104	Chronic Obstructive Pulmonary Disease Is Not Associated with KRAS Mutations in Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2016, 11, e0152317.	2.5	10
105	Small RNA sequencing reveals a comprehensive miRNA signature of BRCA1-associated high-grade serous ovarian cancer. <i>Journal of Clinical Pathology</i> , 2016, 69, 979-985.	2.0	11
106	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
107	Meta-analysis of genome-wide association studies reveals genetic overlap between Hodgkin lymphoma and multiple sclerosis. <i>International Journal of Epidemiology</i> , 2016, 45, 728-740.	1.9	20
108	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

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109	Biomarkers for evaluation of treatment response in classical Hodgkin lymphoma: comparison of <sc>sG</sc> alectinâ€1, <sc>sCD</sc>163 and <sc>sCD</sc>30 with TARC. British Journal of Haematology, 2016, 175, 868-875.	2.5	44
110	CD58 mutations are common in Hodgkin lymphoma cell lines and loss of CD58 expression in tumor cells occurs in Hodgkin lymphoma patients who relapse. Genes and Immunity, 2016, 17, 363-366.	4.1	39
111	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cellâ€Specific Long Noncoding RNAs. American Journal of Pathology, 2016, 186, 2462-2472.	3.8	36
112	Combining genomewide association study and lung <sc>eQTL</sc> analysis provides evidence for novel genes associated with asthma. Allergy: European Journal of Allergy and Clinical Immunology, 2016, 71, 1712-1720.	5.7	47
113	Single-cell sequencing reveals karyotype heterogeneity in murine and human malignancies. Genome Biology, 2016, 17, 115.	8.8	178
114	Functional Studies on Primary Tubular Epithelial Cells Indicate a Tumor Suppressor Role of SETD2 in Clear Cell Renal Cell Carcinoma. Neoplasia, 2016, 18, 339-346.	5.3	23
115	Budesonide and fluticasone propionate differentially affect the airway epithelial barrier. Respiratory Research, 2016, 17, 2.	3.6	30
116	Resistance mechanisms after tyrosine kinase inhibitors afatinib and crizotinib in non-small cell lung cancer, a review of the literature. Critical Reviews in Oncology/Hematology, 2016, 100, 107-116.	4.4	92
117	The European Hematology Association Roadmap for European Hematology Research: a consensus document. Haematologica, 2016, 101, 115-208.	3.5	67
118	Plasma vesicle miRNAs for therapy response monitoring in Hodgkin lymphoma patients. JCI Insight, 2016, 1, e89631.	5.0	121
119	A Novel Prognostic Model Based on Tumor Microenvironment Biology in Relapse Biopsies Predicts Post-Autologous Stem Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. Blood, 2016, 128, 1093-1093.	1.4	12
120	Proteomics Based Identification of Proteins with Deregulated Expression in B Cell Lymphomas. PLoS ONE, 2016, 11, e0146624.	2.5	8
121	Genomic Aberrations in Crizotinib Resistant Lung Adenocarcinoma Samples Identified by Transcriptome Sequencing. PLoS ONE, 2016, 11, e0153065.	2.5	18
122	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. Oncotarget, 2016, 7, 2391-2400.	1.8	43
123	SETD2: an epigenetic modifier with tumor suppressor functionality. Oncotarget, 2016, 7, 50719-50734.	1.8	136
124	CD57+ T-cells are a subpopulation of T-follicular helper cells in nodular lymphocyte predominant Hodgkin lymphoma. Experimental Hematology and Oncology, 2015, 4, 27.	5.0	13
125	Rheumatoid Arthritis, Immunosenescence and the Hallmarks of Aging. Current Aging Science, 2015, 8, 131-146.	1.2	76
126	Microenvironment, Crosstalk, and Immune Escape Mechanisms. Hematologic Malignancies, 2015, , 65-78.	0.2	0

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127	A Novel Risk Locus at 6p21.3 for Epstein-Barr Virus-Positive Hodgkin Lymphoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1838-1843.	2.5	20
128	Pim1 kinase activity preserves airway epithelial integrity upon house dust mite exposure. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2015, 309, L1344-L1353.	2.9	10
129	GATA6 expression in Barrett's oesophagus and oesophageal adenocarcinoma. <i>Digestive and Liver Disease</i> , 2015, 47, 73-80.	0.9	13
130	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
131	Long noncoding RNAs as a novel component of the Myc transcriptional network. <i>FASEB Journal</i> , 2015, 29, 2338-2346.	0.5	67
132	Genomic aberrations guiding treatment of non-small cell lung cancer patients. <i>Cancer Treatment Communications</i> , 2015, 4, 23-33.	0.4	17
133	Immune miRs: critical regulators of T cell development, function and ageing. <i>Immunology</i> , 2015, 144, 1-10.	4.4	141
134	Mir-155 Enhances B-Cell Lymphoma Growth By Targeting TBRC1. <i>Blood</i> , 2015, 126, 4820-4820.	1.4	14
135	Age-Associated Differences in MiRNA Signatures Are Restricted to CD45RO Negative T Cells and Are Associated with Changes in the Cellular Composition, Activation and Cellular Ageing. <i>PLoS ONE</i> , 2015, 10, e0137556.	2.5	23
136	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
137	Abstract 4757: Whole exome sequencing reveals a distinct mutation pattern in metastatic small cell lung cancer compared to non-small cell lung cancer. , 2015, , .		1
138	Abstract 4245: Detection of fusion genes in lung cancer biopsies of crizotinib resistant patients. , 2015, , .		0
139	Mutations in CD58 and Other Immune System Related Genes in Hodgkin Lymphoma. <i>Blood</i> , 2015, 126, 1439-1439.	1.4	0
140	Impact of Radiotherapy and Rituximab on Outcome of Stage I(E) Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2015, 126, 1503-1503.	1.4	0
141	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. <i>Leukemia</i> , 2014, 28, 2248-2251.	7.2	74
142	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
143	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
144	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

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145	Aging disturbs the balance between effector and regulatory CD4+ T cells. <i>Experimental Gerontology</i> , 2014, 60, 190-196.	2.8	115
146	Toll-like receptors in the pathogenesis of human B cell malignancies. <i>Journal of Hematology and Oncology</i> , 2014, 7, 57.	17.0	54
147	Shifted T-cell polarisation after agricultural dust exposure in mice and men. <i>Thorax</i> , 2014, 69, 630-637.	5.6	24
148	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
149	Embryological signaling pathways in Barrett's metaplasia development and malignant transformation; mechanisms and therapeutic opportunities. <i>Critical Reviews in Oncology/Hematology</i> , 2014, 92, 25-37.	4.4	13
150	Insulin-Like Growth Factor 1 Receptor Is a Prognostic Factor in Classical Hodgkin Lymphoma. <i>PLoS ONE</i> , 2014, 9, e87474.	2.5	22
151	Correlation of MicroRNA-16, MicroRNA-21 and MicroRNA-101 Expression with Cyclooxygenase-2 Expression and Angiogenic Factors in Cirrhotic and Noncirrhotic Human Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2014, 9, e95826.	2.5	10
152	GATA6 expression in Barrett's metaplasia development and malignant progression. <i>Journal of Clinical Oncology</i> , 2014, 32, 26-26.	1.6	0
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307	The genomic structure of the human UBE1L gene. <i>Gene Expression</i> , 1995, 4, 163-75.	1.2	15
308	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
309	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
310	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
311	A PCR-Aided Transcript Titration Assay Revealing Very Low Expression of a Gene at Band 3p21 in 33 Cell Lines Derived from all Types of Lung Cancer. <i>European Journal of Human Genetics</i> , 1993, 1, 156-163.	2.8	6
312	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
313	A new RFLP identified at the D3S48 locus. <i>Nucleic Acids Research</i> , 1991, 19, 4797-4797.	14.5	4
314	Partial 3q duplication syndrome and assignment of D3S5 to 3q25â€”3q28. <i>Human Genetics</i> , 1991, 87, 151-154.	3.8	38
315	131 Are chromosome 3 deletions a consistent abnormality in mesotheliomas?. <i>Cancer Genetics and Cytogenetics</i> , 1989, 38, 203.	1.0	2
316	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