

Anke van den Berg

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

294
papers

10,942
citations

56
h-index

93
g-index

327
ext. papers

12,411
ext. citations

5.7
avg, IF

5.81
L-index

#	Paper	IF	Citations
294	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	5.1	3
293	Understanding the role of long-acting muscarinic antagonists in asthma treatment.. <i>Annals of Allergy, Asthma and Immunology</i> , 2022 ,	3.2	1
292	CD4+ T cells in classical Hodgkin lymphoma express exhaustion associated transcription factors TOX and TOX2: Characterizing CD4+ T cells in Hodgkin lymphoma.. <i>Oncolmmunology</i> , 2022 , 11, 2033433	7.2	0
291	Killer Cell Immunoglobulin-Like Receptor Haplotype B Modulates Susceptibility to EBV-Associated Classic Hodgkin Lymphoma.. <i>Frontiers in Immunology</i> , 2022 , 13, 829943	8.4	1
290	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 , 1-7	2	0
289	The relation between age and airway epithelial barrier function.. <i>Respiratory Research</i> , 2022 , 23, 43	7.3	1
288	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	7	2
287	Frequent 4EBP1 Amplification Induces Synthetic Dependence on FGFR Signaling in Cancer. <i>Cancers</i> , 2022 , 14, 2397	6.6	1
286	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	2.2	
285	Responsivity and Reproducibility of Sputum Inflammatory Biomarkers During COPD Exacerbation and Stable Phases - A Pilot Study. <i>International Journal of COPD</i> , 2021 , 16, 3055-3064	3	
284	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	4.7	6
283	Interaction between ERAP Alleles and HLA Class I Types Support a Role of Antigen Presentation in Hodgkin Lymphoma Development. <i>Cancers</i> , 2021 , 13,	6.6	2
282	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	4.5	1
281	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	5	2
280	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> , 2021 ,	4.9	2
279	Frequent mutated B2M, EZH2, IRF8, and TNFRSF14 in primary bone diffuse large B-cell lymphoma reflect a GCB phenotype. <i>Blood Advances</i> , 2021 , 5, 3760-3775	7.8	3
278	Molecular imaging in lymphoma beyond F-FDG-PET: understanding the biology and its implications for diagnostics and therapy. <i>Lancet Haematology</i> , 2020 , 7, e479-e489	14.6	5

277	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	5.9	18
276	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020 , 12,	6.6	10
275	Non-Coding RNAs in Cancer Radiosensitivity: MicroRNAs and lncRNAs as Regulators of Radiation-Induced Signaling Pathways. <i>Cancers</i> , 2020 , 12,	6.6	21
274	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, 2040620719898373	5.7	5
273	Interim thymus and activation regulated chemokine versus interim F-fluorodeoxyglucose positron-emission tomography in classical Hodgkin lymphoma response evaluation. <i>British Journal of Haematology</i> , 2020 , 190, 40-44	4.5	9
272	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	6.6	10
271	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	3.6	18
270	Microenvironment, Cross-Talk, and Immune Escape Mechanisms. <i>Hematologic Malignancies</i> , 2020 , 69-86		0
269	Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma. <i>Cancer Treatment Reviews</i> , 2020 , 82, 101931	14.4	20
268	Analysis of Released Circulating Tumor Cells During Surgery for Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2020 , 26, 1656-1666	12.9	12
267	An All-In-One Transcriptome-Based Assay to Identify Therapy-Guiding Genomic Aberrations in Nonsmall Cell Lung Cancer Patients. <i>Cancers</i> , 2020 , 12,	6.6	2
266	Rosetting T cells in Hodgkin lymphoma are activated by immunological synapse components HLA class II and CD58. <i>Blood</i> , 2020 , 136, 2437-2441	2.2	10
265	MiR-378a-3p Is Critical for Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020 , 12,	6.6	1
264	Targeted sequencing of circulating cell-free DNA in stage II-III resectable oesophageal squamous cell carcinoma patients. <i>BMC Cancer</i> , 2019 , 19, 818	4.8	10
263	Increased miR-142-3p Expression Might Explain Reduced Regulatory T Cell Function in Granulomatosis With Polyangiitis. <i>Frontiers in Immunology</i> , 2019 , 10, 2170	8.4	10
262	Tumour necrosis as assessed with F-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	8	3
261	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	5.1	13
260	Age-related gene and miRNA expression changes in airways of healthy individuals. <i>Scientific Reports</i> , 2019 , 9, 3765	4.9	17

259	CX Chemokine Receptor 7 Contributes to Survival of -Mutant Non-Small Cell Lung Cancer upon Loss of Epidermal Growth Factor Receptor. <i>Cancers</i> , 2019 , 11,	6.6	13
258	NGS-Based High-Throughput Screen to Identify MicroRNAs Regulating Growth of B-Cell Lymphoma. <i>Methods in Molecular Biology</i> , 2019 , 1956, 269-282	1.4	1
257	Long Non-coding RNAs in the Development and Maintenance of Lymphoid Malignancies 2019 , 127-149		
256	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	3.7	3
255	WEE1 Inhibition Enhances Anti-Apoptotic Dependency as a Result of Premature Mitotic Entry and DNA Damage. <i>Cancers</i> , 2019 , 11,	6.6	6
254	Current Smoking is Associated with Decreased Expression of miR-335-5p in Parenchymal Lung Fibroblasts. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	8
253	Marked TGF- β -Regulated miRNA expression changes in both COPD and control lung fibroblasts. <i>Scientific Reports</i> , 2019 , 9, 18214	4.9	9
252	Heterogeneous Pattern of Dependence on Anti-Apoptotic BCL-2 Family Proteins upon CHOP Treatment in Diffuse Large B-Cell Lymphoma. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	6
251	Intricate crosstalk between MYC and non-coding RNAs regulates hallmarks of cancer. <i>Molecular Oncology</i> , 2019 , 13, 26-45	7.9	31
250	Differential miRNA Expression Profiles in Cumulus and Mural Granulosa Cells from Human Pre-ovulatory Follicles. <i>MicroRNA (Sharjah, United Arab Emirates)</i> , 2019 , 8, 61-67	2.9	13
249	Argonaute 2 RNA Immunoprecipitation Reveals Distinct miRNA Targetomes of Primary Burkitt Lymphoma Tumors and Normal B Cells. <i>American Journal of Pathology</i> , 2018 , 188, 1289-1299	5.8	4
248	Effects of fluticasone propionate and budesonide on the expression of immune defense genes in bronchial epithelial cells. <i>Pulmonary Pharmacology and Therapeutics</i> , 2018 , 50, 47-56	3.5	15
247	Combined PD-1 and JAK1/2 inhibition in refractory primary mediastinal B-cell lymphoma. <i>Annals of Hematology</i> , 2018 , 97, 905-907	3	2
246	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	3.3	30
245	Targeting the Microenvironment in Hodgkin Lymphoma: Opportunities and Challenges. <i>Molecular Pathology Library</i> , 2018 , 59-90		
244	Identification of Two Protein-Signaling States Delineating Transcriptionally Heterogeneous Human Medulloblastoma. <i>Cell Reports</i> , 2018 , 22, 3206-3216	10.6	10
243	Impact of acute exposure to cigarette smoke on airway gene expression. <i>Physiological Genomics</i> , 2018 , 50, 705-713	3.6	16
242	Involvement of MicroRNAs in the Aging-Related Decline of CD28 Expression by Human T Cells. <i>Frontiers in Immunology</i> , 2018 , 9, 1400	8.4	8

241	Mutations in EMT-Related Genes in ALK Positive Crizotinib Resistant Non-Small Cell Lung Cancers. <i>Cancers</i> , 2018 , 10,	6.6	25
240	The Microenvironment in Epstein-Barr Virus-Associated Malignancies. <i>Pathogens</i> , 2018 , 7,	4.5	29
239	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	3.7	14
238	Combined loss of HLA I and HLA II expression is more common in the non-GCB type of diffuse large B cell lymphoma. <i>Histopathology</i> , 2018 , 72, 886-888	7.3	3
237	Circulating miRNAs in patients with Barrett's esophagus, high-grade dysplasia and esophageal adenocarcinoma. <i>Journal of Gastrointestinal Oncology</i> , 2018 , 9, 1150-1156	2.8	8
236	KRAS Mutation as a Resistance Mechanism to BRAF/MEK Inhibition in NSCLC. <i>Journal of Thoracic Oncology</i> , 2018 , 13, e249-e251	8.9	10
235	Mutational Evolution in Relapsed Diffuse Large B-Cell Lymphoma. <i>Cancers</i> , 2018 , 10,	6.6	9
234	Tuberous sclerosis complex is required for tumor maintenance in MYC-driven Burkitt's lymphoma. <i>EMBO Journal</i> , 2018 , 37,	13	6
233	MicroRNA High Throughput Loss-of-Function Screening Reveals an Oncogenic Role for miR-21-5p in Hodgkin Lymphoma. <i>Cellular Physiology and Biochemistry</i> , 2018 , 49, 144-159	3.9	17
232	Serum periostin does not reflect type 2-driven inflammation in COPD. <i>Respiratory Research</i> , 2018 , 19, 112	7.3	6
231	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	4.1	11
230	MBRS-36. IDENTIFICATION OF TWO PROTEIN-SIGNALING STATES DELINEATING TRANSCRIPTIONALLY HETEROGENEOUS HUMAN MEDULLOBLASTOMA. <i>Neuro-Oncology</i> , 2018 , 20, i136-1136	7.8	78
229	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	5.7	6
228	Latrophilin receptors: novel bronchodilator targets in asthma. <i>Thorax</i> , 2017 , 72, 74-82	7.3	8
227	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	12.9	48
226	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	5.8	32
225	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	10.3	14
224	HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?. <i>Onc Immunology</i> , 2017 , 6, e1295202	7.2	60

223	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	10.7	24
222	Emerging roles for long noncoding RNAs in B-cell development and malignancy. <i>Critical Reviews in Oncology/Hematology</i> , 2017 , 120, 77-85	7	25
221	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	4.8	27
220	Prognostic Model to Predict Post-Autologous Stem-Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. <i>Journal of Clinical Oncology</i> , 2017 , 35, 3722-3733	2.2	29
219	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	5.2	36
218	PTTG1IP and MAML3, 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	9.3	12
217	Lymphadenopathy driven by TCR-V8V1 T-cell expansion in FAS-related autoimmune lymphoproliferative syndrome. <i>Blood Advances</i> , 2017 , 1, 1101-1106	7.8	2
216	HLA expression and HLA type associations in relation to EBV status in Hispanic Hodgkin lymphoma patients. <i>PLoS ONE</i> , 2017 , 12, e0174457	3.7	6
215	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	3.7	15
214	Identification of transforming growth factor-beta-regulated microRNAs and the microRNA-targetomes in primary lung fibroblasts. <i>PLoS ONE</i> , 2017 , 12, e0183815	3.7	28
213	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, 144-151	4.6	21
212	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	9.3	36
211	Single-cell sequencing reveals karyotype heterogeneity in murine and human malignancies. <i>Genome Biology</i> , 2016 , 17, 115	18.3	99
210	Functional Studies on Primary Tubular Epithelial Cells Indicate a Tumor Suppressor Role of SETD2 in Clear Cell Renal Cell Carcinoma. <i>Neoplasia</i> , 2016 , 18, 339-46	6.4	16
209	Budesonide and fluticasone propionate differentially affect the airway epithelial barrier. <i>Respiratory Research</i> , 2016 , 17, 2	7.3	26
208	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, 107-16	7	76
207	The European Hematology Association Roadmap for European Hematology Research: a consensus document. <i>Haematologica</i> , 2016 , 101, 115-208	6.6	46
206	Plasma vesicle miRNAs for therapy response monitoring in Hodgkin lymphoma patients. <i>JCI Insight</i> , 2016 , 1, e89631	9.9	93

205	A Novel Prognostic Model Based on Tumor Microenvironment Biology in Relapse Biopsies Predicts Post-Autologous Stem Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. <i>Blood</i> , 2016 , 128, 1093-1093	2.2	1
204	Proteomics Based Identification of Proteins with Deregulated Expression in B Cell Lymphomas. <i>PLoS ONE</i> , 2016 , 11, e0146624	3.7	5
203	Genomic Aberrations in Crizotinib Resistant Lung Adenocarcinoma Samples Identified by Transcriptome Sequencing. <i>PLoS ONE</i> , 2016 , 11, e0153065	3.7	11
202	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. <i>Oncotarget</i> , 2016 , 7, 2391-400	3.3	34
201	SETD2: an epigenetic modifier with tumor suppressor functionality. <i>Oncotarget</i> , 2016 , 7, 50719-50734	3.3	89
200	Characterization of the Microenvironment of Nodular Lymphocyte Predominant Hodgkin Lymphoma. <i>International Journal of Molecular Sciences</i> , 2016 , 17,	6.3	9
199	Chronic Obstructive Pulmonary Disease Is Not Associated with KRAS Mutations in Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2016 , 11, e0152317	3.7	8
198	Small RNA sequencing reveals a comprehensive miRNA signature of BRCA1-associated high-grade serous ovarian cancer. <i>Journal of Clinical Pathology</i> , 2016 ,	3.9	8
197	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	4.9	30
196	Meta-analysis of genome-wide association studies reveals genetic overlap between Hodgkin lymphoma and multiple sclerosis. <i>International Journal of Epidemiology</i> , 2016 , 45, 728-40	7.8	16
195	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	7	19
194	Biomarkers for evaluation of treatment response in classical Hodgkin lymphoma: comparison of sGalectin-1, sCD163 and sCD30 with TARC. <i>British Journal of Haematology</i> , 2016 , 175, 868-875	4.5	27
193	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, 363-6	4.4	22
192	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, 2462-72	5.8	27
191	Genomic aberrations guiding treatment of non-small cell lung cancer patients. <i>Cancer Treatment Communications</i> , 2015 , 4, 23-33		14
190	Immuno-miRs: critical regulators of T-cell development, function and ageing. <i>Immunology</i> , 2015 , 144, 1-10	7.8	110
189	CD57+ T-cells are a subpopulation of T-follicular helper cells in nodular lymphocyte predominant Hodgkin lymphoma. <i>Experimental Hematology and Oncology</i> , 2015 , 4, 27	7.8	11
188	Rheumatoid Arthritis, Immunosenescence and the Hallmarks of Aging. <i>Current Aging Science</i> , 2015 , 8, 131-46	2.2	48

187	Microenvironment, Crosstalk, and Immune Escape Mechanisms. <i>Hematologic Malignancies</i> , 2015 , 65-78	o	
186	A Novel Risk Locus at 6p21.3 for Epstein-Barr Virus-Positive Hodgkin Lymphoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1838-43	4	16
185	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, L1344-53	5.8	7
184	GATA6 expression in Barrett's oesophagus and oesophageal adenocarcinoma. <i>Digestive and Liver Disease</i> , 2015 , 47, 73-80	3.3	9
183	SF Treg cells transcribing high levels of Bcl-2 and microRNA-21 demonstrate limited apoptosis in RA. <i>Rheumatology</i> , 2015 , 54, 950-8	3.9	24
182	Long noncoding RNAs as a novel component of the Myc transcriptional network. <i>FASEB Journal</i> , 2015 , 29, 2338-46	0.9	42
181	Mir-155 Enhances B-Cell Lymphoma Growth By Targeting TBRG1. <i>Blood</i> , 2015 , 126, 4820-4820	2.2	3
180	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	3.7	21
179	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, 117-22	2.9	26
178	Mutations in CD58 and Other Immune System Related Genes in Hodgkin Lymphoma. <i>Blood</i> , 2015 , 126, 1439-1439	2.2	
177	Impact of Radiotherapy and Rituximab on Outcome of Stage I(E) Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2015 , 126, 1503-1503	2.2	
176	The microenvironment in classical Hodgkin lymphoma: an actively shaped and essential tumor component. <i>Seminars in Cancer Biology</i> , 2014 , 24, 15-22	12.7	79
175	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, 115-25	4.4	100
174	Aging disturbs the balance between effector and regulatory CD4+ T cells. <i>Experimental Gerontology</i> , 2014 , 60, 190-6	4.5	81
173	Toll-like receptors in the pathogenesis of human B cell malignancies. <i>Journal of Hematology and Oncology</i> , 2014 , 7, 57	22.4	38
172	Shifted T-cell polarisation after agricultural dust exposure in mice and men. <i>Thorax</i> , 2014 , 69, 630-7	7.3	24
171	A meta-analysis of Hodgkin lymphoma reveals 19p13.3 TCF3 as a novel susceptibility locus. <i>Nature Communications</i> , 2014 , 5, 3856	17.4	55
170	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	7	11

169	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. <i>Leukemia</i> , 2014 , 28, 2248-51	10.7	54
168	Genetic associations in classical hodgkin lymphoma: a systematic review and insights into susceptibility mechanisms. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 2737-47	4	30
167	Insulin-like growth factor 1 receptor is a prognostic factor in classical Hodgkin lymphoma. <i>PLoS ONE</i> , 2014 , 9, e87474	3.7	19
166	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	3.7	8
165	Meta-Analysis of Hodgkin Lymphoma and Asthma Genome-Wide Association Scans reveals common variants in GATA3. <i>Blood</i> , 2014 , 124, 135-135	2.2	
164	GATA6 expression in Barrett's metaplasia development and malignant progression.. <i>Journal of Clinical Oncology</i> , 2014 , 32, 26-26	2.2	
163	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, 165-73	5	25
162	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	9.3	53
161	Studying microRNAs in lymphoma. <i>Methods in Molecular Biology</i> , 2013 , 971, 265-76	1.4	12
160	The Role of MicroRNAs in Hodgkin's Lymphoma 2013 , 435-447		
159	PML nuclear bodies and SATB1 are associated with HLA class I expression in EBV+ Hodgkin lymphoma. <i>PLoS ONE</i> , 2013 , 8, e72930	3.7	5
158	Long Non-Coding RNAs Are Commonly Deregulated In Hodgkin Lymphoma. <i>Blood</i> , 2013 , 122, 628-628	2.2	2
157	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	3.7	55
156	Cellular localization and processing of primary transcripts of exonic microRNAs. <i>PLoS ONE</i> , 2013 , 8, e76647	3.7	19
155	Long Non-Coding RNAs As Components Of The MYC Network In B Cell Lymphoma. <i>Blood</i> , 2013 , 122, 1260-1260	2.2	
154	A Meta-Analysis Of Hodgkin Lymphoma Reveals 19p13.3 (TCF3) As a Novel Susceptibility Loc. <i>Blood</i> , 2013 , 122, 626-626	2.2	
153	Circulating tumor cells in small-cell lung cancer: a predictive and prognostic factor. <i>Annals of Oncology</i> , 2012 , 23, 2937-2942	10.3	167
152	Generation of miRNA sponge constructs. <i>Methods</i> , 2012 , 58, 113-7	4.6	71

151	Genome-wide association study of classical Hodgkin lymphoma and Epstein-Barr virus status-defined subgroups. <i>Journal of the National Cancer Institute</i> , 2012 , 104, 240-53	9.7	117
150	Rapid generation of microRNA sponges for microRNA inhibition. <i>PLoS ONE</i> , 2012 , 7, e29275	3.7	103
149	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	3.7	22
148	HLA associations in classical Hodgkin lymphoma: EBV status matters. <i>PLoS ONE</i> , 2012 , 7, e39986	3.7	41
147	Gene expression profiling of microdissected Hodgkin Reed-Sternberg cells correlates with treatment outcome in classical Hodgkin lymphoma. <i>Blood</i> , 2012 , 120, 3530-40	2.2	100
146	Plasma thymus and activation-regulated chemokine as an early response marker in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012 , 97, 410-5	6.6	45
145	Expression of the c-Met oncogene by tumor cells predicts a favorable outcome in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012 , 97, 572-8	6.6	26
144	Expression, Prognostic Value and Function of IGF1R in Hodgkin Lymphoma.. <i>Blood</i> , 2012 , 120, 2640-2640.	2.2	1
143	A chronic obstructive pulmonary disease related signature in squamous cell lung cancer. <i>Lung Cancer</i> , 2011 , 72, 177-83	5.9	23
142	Multiple HLA class I and II associations in classical Hodgkin lymphoma and EBV status defined subgroups. <i>Blood</i> , 2011 , 118, 5211-7	2.2	1
141	MHC class II transactivator CIITA is a recurrent gene fusion partner in lymphoid cancers. <i>Nature</i> , 2011 , 471, 377-81	50.4	467
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