Anke van den Berg

List of Publications by Citations

Source: https://exaly.com/author-pdf/6651512/anke-van-den-berg-publications-by-citations.pdf
Version: 2024-04-24

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
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	BIC and miR-155 are highly expressed in Hodgkin, primary mediastinal and diffuse large B cell lymphomas. *Journal of Pathology*, **2005**, 207, 243-9	9.4	584
293	MHC class II transactivator CIITA is a recurrent gene fusion partner in lymphoid cancers. *Nature*, **2011**, 471, 377-81	50.4	467
292	Somatic mutations of the von Hippel-Lindau disease tumour suppressor gene in non-familial clear cell renal carcinoma. *Human Molecular Genetics*, **1994**, 3, 2169-73	5.6	314
291	High expression of the CC chemokine TARC in Reed-Sternberg cells. A possible explanation for the characteristic T-cell infiltratein Hodgkin's lymphoma. *American Journal of Pathology*, **1999**, 154, 1685-91	5.8	313
290	Cigarette smoke-induced emphysema: A role for the B cell?. *American Journal of Respiratory and Critical Care Medicine*, **2006**, 173, 751-8	10.2	242
289	High expression of B-cell receptor inducible gene BIC in all subtypes of Hodgkin lymphoma. *Genes Chromosomes and Cancer*, **2003**, 37, 20-8	5	208
288	Lack of BIC and microRNA miR-155 expression in primary cases of Burkitt lymphoma. *Genes Chromosomes and Cancer*, **2006**, 45, 147-53	5	204
287	MicroRNAs, macrocontrol: regulation of miRNA processing. *Rna*, **2010**, 16, 1087-95	5.8	195
286	Circulating tumor cells in small-cell lung cancer: a predictive and prognostic factor. *Annals of Oncology*, **2012**, 23, 2937-2942	10.3	167
285	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. *BMC Cancer*, **2009**, 9, 163	4.8	164
284	A genome-wide association study of Hodgkin's lymphoma identifies new susceptibility loci at 2p16.1 (REL), 8q24.21 and 10p14 (GATA3). *Nature Genetics*, **2010**, 42, 1126-1130	36.3	158
283	Cytoplasmic p21 expression levels determine cisplatin resistance in human testicular cancer. *Journal of Clinical Investigation*, **2010**, 120, 3594-605	15.9	157
282	Association with HLA class I in Epstein-Barr-virus-positive and with HLA class III in Epstein-Barr-virus-negative Hodgkin's lymphoma. *Lancet, The*, **2005**, 365, 2216-24	40	130
281	Isolation of the human semaphorin III/F gene (SEMA3F) at chromosome 3p21, a region deleted in lung cancer. *Genomics*, **1996**, 32, 39-48	4.3	127
280	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. *Blood*, **2003**, 101, 1149-54	2.2	122
279	Regulation of pri-microRNA BIC transcription and processing in Burkitt lymphoma. *Oncogene*, **2007**, 26, 3769-76	9.2	121
278	Chemokines, cytokines and their receptors in Hodgkin's lymphoma cell lines and tissues. *Annals of Oncology*, **2002**, 13 Suppl 1, 52-6	10.3	118

277	Genome-wide association study of classical Hodgkin lymphoma and Epstein-Barr virus status-defined subgroups. *Journal of the National Cancer Institute*, **2012**, 104, 240-53	9.7	117
276	Hodgkin lymphoma cell lines are characterized by a specific miRNA expression profile. *Neoplasia*, **2009**, 11, 167-76	6.4	111
275	HLA-A*02 is associated with a reduced risk and HLA-A*01 with an increased risk of developing EBV+ Hodgkin lymphoma. *Blood*, **2007**, 110, 3310-5	2.2	111
274	Immuno-miRs: critical regulators of T-cell development, function and ageing. *Immunology*, **2015**, 144, 1-10	7.8	110
273	HLA class II expression by Hodgkin Reed-Sternberg cells is an independent prognostic factor in classical Hodgkin's lymphoma. *Journal of Clinical Oncology*, **2007**, 25, 3101-8	2.2	107
272	Rapid generation of microRNA sponges for microRNA inhibition. *PLoS ONE*, **2012**, 7, e29275	3.7	103
271	Comprehensive analysis of miRNA expression in T-cell subsets of rheumatoid arthritis patients reveals defined signatures of naive and memory Tregs. *Genes and Immunity*, **2014**, 15, 115-25	4.4	100
270	Gene expression profiling of microdissected Hodgkin Reed-Sternberg cells correlates with treatment outcome in classical Hodgkin lymphoma. *Blood*, **2012**, 120, 3530-40	2.2	100
269	miRNA analysis in B-cell chronic lymphocytic leukaemia: proliferation centres characterized by low miR-150 and high BIC/miR-155 expression. *Journal of Pathology*, **2008**, 215, 13-20	9.4	100
268	Single-cell sequencing reveals karyotype heterogeneity in murine and human malignancies. *Genome Biology*, **2016**, 17, 115	18.3	99
267	Proteomics analysis of Hodgkin lymphoma: identification of new players involved in the cross-talk between HRS cells and infiltrating lymphocytes. *Blood*, **2008**, 111, 2339-46	2.2	99
266	A high throughput experimental approach to identify miRNA targets in human cells. *Nucleic Acids Research*, **2009**, 37, e137	20.1	94
265	Serum chemokine levels in Hodgkin lymphoma patients: highly increased levels of CCL17 and CCL22. *British Journal of Haematology*, **2008**, 140, 527-36	4.5	93
264	Plasma vesicle miRNAs for therapy response monitoring in Hodgkin lymphoma patients. *JCI Insight*, **2016**, 1, e89631	9.9	93
263	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. *Journal of Clinical Oncology*, **2011**, 29, 4320-6	2.2	91
262	SETD2: an epigenetic modifier with tumor suppressor functionality. *Oncotarget*, **2016**, 7, 50719-50734	3.3	89
261	miRNA profiling of B-cell subsets: specific miRNA profile for germinal center B cells with variation between centroblasts and centrocytes. *Laboratory Investigation*, **2009**, 89, 708-16	5.9	87
260	The role of microRNAs in normal hematopoiesis and hematopoietic malignancies. *Leukemia*, **2006**, 20, 1931-6	10.7	86

259	A gene in the chromosomal region 3p21 with greatly reduced expression in lung cancer is similar to the gene for ubiquitin-activating enzyme. *Proceedings of the National Academy of Sciences of the United States of America*, **1993**, 90, 6071-5	11.5	84
258	Strongly enhanced IL-10 production using stable galectin-1 homodimers. *Molecular Immunology*, **2007**, 44, 506-13	4.3	83
257	Aging disturbs the balance between effector and regulatory CD4+ T cells. *Experimental Gerontology* , **2014**, 60, 190-6	4.5	81
256	The microenvironment in classical Hodgkin lymphoma: an actively shaped and essential tumor component. *Seminars in Cancer Biology*, **2014**, 24, 15-22	12.7	79
255	Interaction between host T cells and Reed-Sternberg cells in Hodgkin lymphomas. *Seminars in Cancer Biology*, **2000**, 10, 345-50	12.7	79
254	MBRS-36. IDENTIFICATION OF TWO PROTEIN-SIGNALING STATES DELINEATING TRANSCRIPTIONALLY HETEROGENEOUS HUMAN MEDULLOBLASTOMA. *Neuro-Oncology*, **2018**, 20, i136-i136	1	78
253	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, 107-16	7	76
252	Dimeric galectin-1 induces IL-10 production in T-lymphocytes: an important tool in the regulation of the immune response. *Journal of Pathology*, **2004**, 204, 511-8	9.4	75
251	Generation of miRNA sponge constructs. *Methods*, **2012**, 58, 113-7	4.6	71
250	Latent Epstein-Barr virus infection of tumor cells in classical Hodgkin's lymphoma predicts adverse outcome in older adult patients. *Journal of Clinical Oncology*, **2009**, 27, 3815-21	2.2	71
249	Specific expression of miR-17-5p and miR-127 in testicular and central nervous system diffuse large B-cell lymphoma. *Modern Pathology*, **2009**, 22, 547-55	9.8	67
248	Involvement of multiple loci on chromosome 3 in renal cell cancer development **1997**, 19, 59-76		67
247	HLA-G protein expression as a potential immune escape mechanism in classical Hodgkin's lymphoma. *Tissue Antigens*, **2008**, 71, 219-26		67
246	Clonal relation in a case of CLL, ALCL, and Hodgkin composite lymphoma. *Blood*, **2002**, 100, 1425-1429	2.2	67
245	An alternative route for multistep tumorigenesis in a novel case of hereditary renal cell cancer and a t(2;3)(q35;q21) chromosome translocation. *American Journal of Human Genetics*, **1998**, 62, 1475-83	11	64
244	MiRNA profiling in B non-Hodgkin lymphoma: a MYC-related miRNA profile characterizes Burkitt lymphoma. *British Journal of Haematology*, **2010**, 149, 896-9	4.5	63
243	HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?. *OncoImmunology*, **2017**, 6, e1295202	7.2	60
242	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?. *Human Pathology*, **2006**, 37, 528-33	3.7	59

241	A homozygous deletion in a small cell lung cancer cell line involving a 3p21 region with a marked instability in yeast artificial chromosomes. *Cancer Research*, **1994**, 54, 4183-7	10.1	58
240	Development of lymphoma in Autoimmune Lymphoproliferative Syndrome (ALPS) and its relationship to Fas gene mutations. *Leukemia and Lymphoma*, **2004**, 45, 423-31	1.9	56
239	Common and differential chemokine expression patterns in rs cells of NLP, EBV positive and negative classical Hodgkin lymphomas. *International Journal of Cancer*, **2002**, 99, 665-72	7.5	56
238	A meta-analysis of Hodgkin lymphoma reveals 19p13.3 TCF3 as a novel susceptibility locus. *Nature Communications*, **2014**, 5, 3856	17.4	55
237	Protease activity of plasma hemopexin. *Kidney International*, **2005**, 68, 603-10	9.9	55
236	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. *PLoS ONE*, **2013**, 8, e76217	3.7	55
235	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. *Leukemia*, **2014**, 28, 2248-51	10.7	54
234	Treatment of the bronchial tree from beginning to end: targeting small airway inflammation in asthma. *Allergy: European Journal of Allergy and Clinical Immunology*, **2013**, 68, 16-26	9.3	53
233	Current smoking-specific gene expression signature in normal bronchial epithelium is enhanced in squamous cell lung cancer. *Journal of Pathology*, **2009**, 218, 182-91	9.4	53
232	The CD4+CD26- T-cell population in classical Hodgkin's lymphoma displays a distinctive regulatory T-cell profile. *Laboratory Investigation*, **2008**, 88, 482-90	5.9	53
231	Mitoxantrone resistance in a small cell lung cancer cell line is associated with ABCA2 upregulation. *British Journal of Cancer*, **2004**, 90, 2411-7	8.7	53
230	Genomic aberrations in squamous cell lung carcinoma related to lymph node or distant metastasis. *Lung Cancer*, **2009**, 66, 372-8	5.9	50
229	Low frequency of FAS mutations in Reed-Sternberg cells of Hodgkin's lymphoma. *American Journal of Pathology*, **2003**, 162, 29-35	5.8	50
228	TP53 gene mutations in Hodgkin lymphoma are infrequent and not associated with absence of Epstein-Barr virus. *International Journal of Cancer*, **2001**, 94, 60-6	7.5	50
227	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. *FEBS Letters*, **1998**, 426, 266-70	3.8	49
226	Dichotomous ALK-IHC Is a Better Predictor for ALK Inhibition Outcome than Traditional ALK-FISH in Advanced Non-Small Cell Lung Cancer. *Clinical Cancer Research*, **2017**, 23, 4251-4258	12.9	48
225	Rheumatoid Arthritis, Immunosenescence and the Hallmarks of Aging. *Current Aging Science*, **2015**, 8, 131-46	2.2	48
224	Expression of the T-cell transcription factors, GATA-3 and T-bet, in the neoplastic cells of Hodgkin lymphomas. *American Journal of Pathology*, **2005**, 166, 127-34	5.8	48

223	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. *Genes Chromosomes and Cancer*, **1996**, 15, 64-72	5	48
222	Differential expression and distribution of epithelial adhesion molecules in non-small cell lung cancer and normal bronchus. *Journal of Clinical Pathology*, **2007**, 60, 608-14	3.9	47
221	The European Hematology Association Roadmap for European Hematology Research: a consensus document. *Haematologica*, **2016**, 101, 115-208	6.6	46
220	Plasma thymus and activation-regulated chemokine as an early response marker in classical Hodgkin's lymphoma. *Haematologica*, **2012**, 97, 410-5	6.6	45
219	Long noncoding RNAs as a novel component of the Myc transcriptional network. *FASEB Journal*, **2015**, 29, 2338-46	0.9	42
218	HLA associations in classical Hodgkin lymphoma: EBV status matters. *PLoS ONE*, **2012**, 7, e39986	3.7	41
217	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. *Journal of Clinical Endocrinology and Metabolism*, **1996**, 81, 2881-2884	5.6	41
216	Immune reactions in classical Hodgkin's lymphoma. *Seminars in Hematology*, **1999**, 36, 253-9	4	41
215	The role of female sex hormones in the development and severity of allergic and non-allergic asthma. *Clinical and Experimental Allergy*, **2009**, 39, 1477-81	4.1	40
214	A comparison of genomic structures and expression patterns of two closely related flanking genes in a critical lung cancer region at 3p21.3. *European Journal of Human Genetics*, **1999**, 7, 478-86	5.3	40
213	Toll-like receptors in the pathogenesis of human B cell malignancies. *Journal of Hematology and Oncology*, **2014**, 7, 57	22.4	38
212	Splenic marginal zone lymphomas presenting with splenomegaly and typical immunophenotype are characterized by allelic loss in 7q31-32. *Modern Pathology*, **2003**, 16, 1210-7	9.8	38
211	Genetic susceptibility to Hodgkin's lymphoma associated with the human leukocyte antigen region. *European Journal of Haematology*, **2005**, 75, 34-41	3.8	38
210	Partial 3q duplication syndrome and assignment of D3S5 to 3q25-3q28. *Human Genetics*, **1991**, 87, 151-4	6.3	37
209	Combining genomewide association study and lung eQTL analysis provides evidence for novel genes associated with asthma. *Allergy: European Journal of Allergy and Clinical Immunology*, **2016**, 71, 1712-1720	9.3	36
208	Building Bridges for Innovation in Ageing: Synergies between Action Groups of the EIP on AHA. *Journal of Nutrition, Health and Aging*, **2017**, 21, 92-104	5.2	36
207	Induction of glomerular alkaline phosphatase after challenge with lipopolysaccharide. *International Journal of Experimental Pathology*, **2003**, 84, 135-44	2.8	36
206	Cytokine gene expression profile distinguishes CD4+/CD57+ T cells of the nodular lymphocyte predominance type of Hodgkin's lymphoma from their tonsillar counterparts. *Journal of Pathology*, **2006**, 208, 423-30	9.4	35

205	Normal FHIT transcripts in renal cell cancer- and lung cancer-derived cell lines, including a cell line with a homozygous deletion in the FRA3B region. *Genes Chromosomes and Cancer*, **1997**, 19, 220-7	5	34
204	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. *Oncotarget*, **2016**, 7, 2391-400	3.3	34
203	A gene from human chromosome region 3p21 with reduced expression in small cell lung cancer. *Cancer Research*, **1992**, 52, 1536-41	10.1	33
202	miR-24-3p Is Overexpressed in Hodgkin Lymphoma and Protects Hodgkin and Reed-Sternberg Cells from Apoptosis. *American Journal of Pathology*, **2017**, 187, 1343-1355	5.8	32
201	Analysis of multiple renal cell adenomas and carcinomas suggests allelic loss at 3p21 to be a prerequisite for malignant development. *Genes Chromosomes and Cancer*, **1997**, 19, 228-32	5	32
200	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. *Cancer Epidemiology Biomarkers and Prevention*, **2006**, 15, 2280-4	4	32
199	High expression of calcium-binding proteins, S100A10, S100A11 and CALM2 in anaplastic large cell lymphoma. *British Journal of Haematology*, **2005**, 131, 596-608	4.5	32
198	Analysis of chromosomal copy number changes and oncoprotein expression in primary central nervous system lymphomas: frequent loss of chromosome arm 6q. *Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin*, **2003**, 443, 164-9	5.1	31
197	P53 mutation analysis of colorectal liver metastases: relation to actual survival, angiogenic status, and p53 overexpression. *Clinical Cancer Research*, **2005**, 11, 4067-73	12.9	31
196	Intricate crosstalk between MYC and non-coding RNAs regulates hallmarks of cancer. *Molecular Oncology*, **2019**, 13, 26-45	7.9	31
195	An 80 Kb P1 clone from chromosome 3p21.3 suppresses tumor growth in vivo. *Oncogene*, **1996**, 13, 2387-96	9.2	31
194	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	3.3	30
193	Genetic associations in classical hodgkin lymphoma: a systematic review and insights into susceptibility mechanisms. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 2737-47	4	30
192	Analysis of serum immune markers in seropositive and seronegative rheumatoid arthritis and in high-risk seropositive arthralgia patients. *Scientific Reports*, **2016**, 6, 26021	4.9	30
191	Prognostic Model to Predict Post-Autologous Stem-Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. *Journal of Clinical Oncology*, **2017**, 35, 3722-3733	2.2	29
190	The Microenvironment in Epstein-Barr Virus-Associated Malignancies. *Pathogens*, **2018**, 7,	4.5	29
189	MiR-17/106b seed family regulates p21 in Hodgkin's lymphoma. *Journal of Pathology*, **2011**, 225, 609-17	9.4	29
188	Germline FAS gene mutation in a case of ALPS and NLP Hodgkin lymphoma. *Blood*, **2002**, 99, 1492-4	2.2	29

187	Direct molecular analysis of a deletion of 3p in tumors from patients with sporadic renal cell carcinoma. *Cancer Genetics and Cytogenetics*, **1988**, 32, 281-5		29
186	Identification of transforming growth factor-beta-regulated microRNAs and the microRNA-targetomes in primary lung fibroblasts. *PLoS ONE*, **2017**, 12, e0183815	3.7	28
185	MicroRNA profiling of human primary macrophages exposed to dengue virus identifies miRNA-3614-5p as antiviral and regulator of ADAR1 expression. *PLoS Neglected Tropical Diseases*, **2017**, 11, e0005981	4.8	27
184	Prolonged protection of the new inhaled corticosteroid fluticasone furoate against AMP hyperresponsiveness in patients with asthma. *Allergy: European Journal of Allergy and Clinical Immunology*, **2010**, 65, 1531-5	9.3	27
183	Brain death induces inflammation in the donor intestine. *Transplantation*, **2008**, 86, 148-54	1.8	27
182	Biomarkers for evaluation of treatment response in classical Hodgkin lymphoma: comparison of sGalectin-1, sCD163 and sCD30 with TARC. *British Journal of Haematology*, **2016**, 175, 868-875	4.5	27
181	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cell-Specific Long Noncoding RNAs. *American Journal of Pathology*, **2016**, 186, 2462-72	5.8	27
180	Budesonide and fluticasone propionate differentially affect the airway epithelial barrier. *Respiratory Research*, **2016**, 17, 2	7.3	26
179	Expression of the c-Met oncogene by tumor cells predicts a favorable outcome in classical Hodgkin's lymphoma. *Haematologica*, **2012**, 97, 572-8	6.6	26
178	Production of hemopexin by TNF-alpha stimulated human mesangial cells. *Kidney International*, **2003**, 63, 1681-6	9.9	26
177	T-cell Activation Induces Dynamic Changes in miRNA Expression Patterns in CD4 and CD8 T-cell Subsets. *MicroRNA (Shariqah, United Arab Emirates)*, **2015**, 4, 117-22	2.9	26
176	Emerging roles for long noncoding RNAs in B-cell development and malignancy. *Critical Reviews in Oncology/Hematology*, **2017**, 120, 77-85	7	25
175	Mutations in EMT-Related Genes in ALK Positive Crizotinib Resistant Non-Small Cell Lung Cancers. *Cancers*, **2018**, 10,	6.6	25
174	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. *Genes Chromosomes and Cancer*, **2013**, 52, 165-73	5	25
173	Maternal smoking during pregnancy decreases Wnt signalling in neonatal mice. *Thorax*, **2010**, 65, 553-4	7.3	25
172	BCL6 alternative translocation breakpoint cluster region associated with follicular lymphoma grade 3B. *Genes Chromosomes and Cancer*, **2005**, 44, 301-4	5	25
171	ZDHHC11 and ZDHHC11B are critical novel components of the oncogenic MYC-miR-150-MYB network in Burkitt lymphoma. *Leukemia*, **2017**, 31, 1470-1473	10.7	24
170	Shifted T-cell polarisation after agricultural dust exposure in mice and men. *Thorax*, **2014**, 69, 630-7	7.3	24

169	SF Treg cells transcribing high levels of Bcl-2 and microRNA-21 demonstrate limited apoptosis in RA. *Rheumatology*, **2015**, 54, 950-8	3.9	24
168	BCL6 alternative breakpoint region break and homozygous deletion of 17q24 in the nodular lymphocyte predominance type of Hodgkin's lymphoma-derived cell line DEV. *Human Pathology*, **2006**, 37, 675-83	3.7	24
167	Identification of chromosomal copy number changes associated with transformation of follicular lymphoma to diffuse large B-cell lymphoma. *Human Pathology*, **2003**, 34, 915-23	3.7	24
166	A chronic obstructive pulmonary disease related signature in squamous cell lung cancer. *Lung Cancer*, **2011**, 72, 177-83	5.9	23
165	HLA-A*02:07 is a protective allele for EBV negative and a susceptibility allele for EBV positive classical Hodgkin lymphoma in China. *PLoS ONE*, **2012**, 7, e31865	3.7	22
164	Gene expression analysis of dendritic/Langerhans cells and Langerhans cell histiocytosis. *Journal of Pathology*, **2006**, 209, 474-83	9.4	22
163	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, 363-6	4.4	22
162	Involvement of multiple loci on chromosome 3 in renal cell cancer development. *Genes Chromosomes and Cancer*, **1997**, 19, 59-76	5	22
161	Non-Coding RNAs in Cancer Radiosensitivity: MicroRNAs and lncRNAs as Regulators of Radiation-Induced Signaling Pathways. *Cancers*, **2020**, 12,	6.6	21
160	Global correlation of genome and transcriptome changes in classical Hodgkin lymphoma. *Hematological Oncology*, **2007**, 25, 21-9	1.3	21
159	Epidemiology of classical Hodgkin lymphoma and its association with Epstein Barr virus in Northern China. *PLoS ONE*, **2011**, 6, e21152	3.7	21
158	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. *PLoS ONE*, **2015**, 10, e0137556	3.7	21
157	Mutation patterns in small cell and non-small cell lung cancer patients suggest a different level of heterogeneity between primary and metastatic tumors. *Carcinogenesis*, **2017**, 38, 144-151	4.6	21
156	Expression of CD1d and presence of invariant NKT cells in classical Hodgkin lymphoma. *American Journal of Hematology*, **2010**, 85, 539-41	7.1	20
155	Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma. *Cancer Treatment Reviews*, **2020**, 82, 101931	14.4	20
154	Brain death causes structural and inflammatory changes in donor intestine. *Transplantation Proceedings*, **2005**, 37, 448-9	1.1	19
153	Cellular localization and processing of primary transcripts of exonic microRNAs. *PLoS ONE*, **2013**, 8, e76647	3.7	19
152	Insulin-like growth factor 1 receptor is a prognostic factor in classical Hodgkin lymphoma. *PLoS ONE* , **2014**, 9, e87474	3.7	19

151	The microenvironment of classical Hodgkin lymphoma: heterogeneity by Epstein-Barr virus presence and location within the tumor. *Blood Cancer Journal*, **2016**, 6, e417	7	19
150	Combined osimertinib, dabrafenib and trametinib treatment for advanced non-small-cell lung cancer patients with an osimertinib-induced BRAF V600E mutation. *Lung Cancer*, **2020**, 146, 358-361	5.9	18
149	Relevance and Effectiveness of Molecular Tumor Board Recommendations for Patients With Non-Small-Cell Lung Cancer With Rare or Complex Mutational Profiles.. *JCO Precision Oncology*, **2020**, 4, 393-410	3.6	18
148	Age-related gene and miRNA expression changes in airways of healthy individuals. *Scientific Reports* , **2019**, 9, 3765	4.9	17
147	Microarray amplification bias: loss of 30% differentially expressed genes due to long probe - poly(A)-tail distances. *BMC Genomics*, **2007**, 8, 277	4.5	17
146	High expression of Mcl-1 in ALK positive and negative anaplastic large cell lymphoma. *Journal of Clinical Pathology*, **2005**, 58, 520-4	3.9	17
145	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	3.9	17
144	Functional Studies on Primary Tubular Epithelial Cells Indicate a Tumor Suppressor Role of SETD2 in Clear Cell Renal Cell Carcinoma. *Neoplasia*, **2016**, 18, 339-46	6.4	16
143	Impact of acute exposure to cigarette smoke on airway gene expression. *Physiological Genomics*, **2018**, 50, 705-713	3.6	16
142	A Novel Risk Locus at 6p21.3 for Epstein-Barr Virus-Positive Hodgkin Lymphoma. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 1838-43	4	16
141	High expression of TIAF-1 in chronic kidney and liver allograft rejection and in activated T-helper cells. *Transplantation*, **2003**, 75, 2076-82	1.8	16
140	An evolutionary rearrangement of the Xp11.3-11.23 region in 3p21.3, a region frequently deleted in a variety of cancers. *Genomics*, **1999**, 60, 238-40	4.3	16
139	Chromosomal findings and p53-mutation analysis in chromophilic renal-cell carcinomas. *International Journal of Cancer*, **1996**, 68, 47-50	7.5	16
138	Meta-analysis of genome-wide association studies reveals genetic overlap between Hodgkin lymphoma and multiple sclerosis. *International Journal of Epidemiology*, **2016**, 45, 728-40	7.8	16
137	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	3.5	15
136	Genetics as a diagnostic tool in sarcomatoid renal-cell cancer. *International Journal of Cancer*, **1997**, 72, 265-9	7.5	15
135	Upregulation of ADAM19 in chronic allograft nephropathy. *American Journal of Transplantation*, **2006**, 6, 1673-81	8.7	15
134	Report: workshop on mediastinal grey zone lymphoma. *European Journal of Haematology*, **2005**, 75, 45-53	3.8	15

133	Expression of HLA class I and HLA class II by tumor cells in Chinese classical Hodgkin lymphoma patients. *PLoS ONE*, **2010**, 5, e10865	3.7	15
132	Overall survival in EGFR mutated non-small-cell lung cancer patients treated with afatinib after EGFR TKI and resistant mechanisms upon disease progression. *PLoS ONE*, **2017**, 12, e0182885	3.7	15
131	The genomic structure of the human UBE1L gene. *Gene Expression*, **1995**, 4, 163-75	3.4	15
130	Clonal relation in a case of CLL, ALCL, and Hodgkin composite lymphoma. *Blood*, **2002**, 100, 1425-9	2.2	15
129	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. *Annals of Oncology*, **2017**, 28, 1668-1670	10.3	14
128	Genomic aberrations guiding treatment of non-small cell lung cancer patients. *Cancer Treatment Communications*, **2015**, 4, 23-33		14
127	Array comparative genomic hybridization reveals a very high frequency of deletions of the long arm of chromosome 6 in testicular lymphoma. *Genes Chromosomes and Cancer*, **2006**, 45, 976-81	5	14
126	Identification of relevant drugable targets in diffuse large B-cell lymphoma using a genome-wide unbiased CD20 guilt-by association approach. *PLoS ONE*, **2018**, 13, e0193098	3.7	14
125	Recommendations for the clinical interpretation and reporting of copy number gains using gene panel NGS analysis in routine diagnostics. *Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin*, **2019**, 474, 673-680	5.1	13
124	CX Chemokine Receptor 7 Contributes to Survival of -Mutant Non-Small Cell Lung Cancer upon Loss of Epidermal Growth Factor Receptor. *Cancers*, **2019**, 11,	6.6	13
123	Similar efficacy of ciclesonide versus prednisolone to treat asthma worsening after steroid tapering. *Respiratory Medicine*, **2009**, 103, 1216-23	4.6	13
122	Composite lymphoma: EBV-positive classic Hodgkin lymphoma and peripheral T-cell lymphoma: a case report. *Applied Immunohistochemistry and Molecular Morphology*, **2009**, 17, 72-6	1.9	13
121	Hodgkin's lymphoma associated T-cells exhibit a transcription factor profile consistent with distinct lymphoid compartments. *Journal of Clinical Pathology*, **2007**, 60, 1092-7	3.9	13
120	Differential miRNA Expression Profiles in Cumulus and Mural Granulosa Cells from Human Pre-ovulatory Follicles. *MicroRNA (Shariqah, United Arab Emirates)*, **2019**, 8, 61-67	2.9	13
119	PTTG1IP and MAML3, novel genomewide association study genes for severity of hyperresponsiveness in adult asthma. *Allergy: European Journal of Allergy and Clinical Immunology*, **2017**, 72, 792-801	9.3	12
118	Studying microRNAs in lymphoma. *Methods in Molecular Biology*, **2013**, 971, 265-76	1.4	12
117	Analysis of Released Circulating Tumor Cells During Surgery for Non-Small Cell Lung Cancer. *Clinical Cancer Research*, **2020**, 26, 1656-1666	12.9	12
116	Embryological signaling pathways in Barrett's metaplasia development and malignant transformation; mechanisms and therapeutic opportunities. *Critical Reviews in Oncology/Hematology*, **2014**, 92, 25-37	7	11

115	CD57+ T-cells are a subpopulation of T-follicular helper cells in nodular lymphocyte predominant Hodgkin lymphoma. *Experimental Hematology and Oncology*, **2015**, 4, 27	7.8	11
114	RT-PCR and immunohistochemical evaluation of sentinel lymph nodes after in vivo mapping with Patent Blue V in colon cancer patients. *Scandinavian Journal of Gastroenterology*, **2006**, 41, 1073-8	2.4	11
113	Defining the position of the breakpoint of the constitutional t(3;6) occurring in a family with renal cell carcinoma. *Genes Chromosomes and Cancer*, **1995**, 12, 224-8	5	11
112	Genomic Aberrations in Crizotinib Resistant Lung Adenocarcinoma Samples Identified by Transcriptome Sequencing. *PLoS ONE*, **2016**, 11, e0153065	3.7	11
111	Novel genes and insights in complete asthma remission: A genome-wide association study on clinical and complete asthma remission. *Clinical and Experimental Allergy*, **2018**, 48, 1286-1296	4.1	11
110	Targeted sequencing of circulating cell-free DNA in stage II-III resectable oesophageal squamous cell carcinoma patients. *BMC Cancer*, **2019**, 19, 818	4.8	10
109	Increased miR-142-3p Expression Might Explain Reduced Regulatory T Cell Function in Granulomatosis With Polyangiitis. *Frontiers in Immunology*, **2019**, 10, 2170	8.4	10
108	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. *Cancers*, **2020**, 12,	6.6	10
107	Treatment of patients with MYC rearrangement positive large B-cell lymphoma with R-CHOP plus lenalidomide: results of a multicenter HOVON phase II trial. *Haematologica*, **2020**, 105, 2805-2812	6.6	10
106	Identification of Two Protein-Signaling States Delineating Transcriptionally Heterogeneous Human Medulloblastoma. *Cell Reports*, **2018**, 22, 3206-3216	10.6	10
105	FAS gene mutation in a case of autoimmune lymphoproliferative syndrome type IA with accumulation of gammadelta+ T cells. *American Journal of Surgical Pathology*, **2003**, 27, 546-53	6.7	10
104	Enhanced ecto-apyrase activity of stimulated endothelial or mesangial cells is downregulated by glucocorticoids in vitro. *European Journal of Pharmacology*, **2004**, 501, 191-8	5.3	10
103	Prolonged survival of rat islet xenografts in mice after CD45RB monotherapy. *Transplantation*, **2004**, 77, 386-91	1.8	10
102	Rosetting T cells in Hodgkin lymphoma are activated by immunological synapse components HLA class II and CD58. *Blood*, **2020**, 136, 2437-2441	2.2	10
101	KRAS Mutation as a Resistance Mechanism to BRAF/MEK Inhibition in NSCLC. *Journal of Thoracic Oncology*, **2018**, 13, e249-e251	8.9	10
100	Interim thymus and activation regulated chemokine versus interim F-fluorodeoxyglucose positron-emission tomography in classical Hodgkin lymphoma response evaluation. *British Journal of Haematology*, **2020**, 190, 40-44	4.5	9
99	GATA6 expression in Barrett's oesophagus and oesophageal adenocarcinoma. *Digestive and Liver Disease*, **2015**, 47, 73-80	3.3	9
98	Characterization of the Microenvironment of Nodular Lymphocyte Predominant Hodgkin Lymphoma. *International Journal of Molecular Sciences*, **2016**, 17,	6.3	9

97	Marked TGF-β regulated miRNA expression changes in both COPD and control lung fibroblasts. *Scientific Reports*, **2019**, 9, 18214	4.9	9
96	Mutational Evolution in Relapsed Diffuse Large B-Cell Lymphoma. *Cancers*, **2018**, 10,	6.6	9
95	Latrophilin receptors: novel bronchodilator targets in asthma. *Thorax*, **2017**, 72, 74-82	7.3	8
94	Involvement of MicroRNAs in the Aging-Related Decline of CD28 Expression by Human T Cells. *Frontiers in Immunology*, **2018**, 9, 1400	8.4	8
93	Current Smoking is Associated with Decreased Expression of miR-335-5p in Parenchymal Lung Fibroblasts. *International Journal of Molecular Sciences*, **2019**, 20,	6.3	8
92	Analysis of in vivo endothelial cell activation applying RT-PCR following endothelial cell isolation by laser dissection microscopy. *Annals of the New York Academy of Sciences*, **2002**, 973, 586-9	6.5	8
91	Correlation of microRNA-16, microRNA-21 and microRNA-101 expression with cyclooxygenase-2 expression and angiogenic factors in cirrhotic and noncirrhotic human hepatocellular carcinoma. *PLoS ONE*, **2014**, 9, e95826	3.7	8
90	Chronic Obstructive Pulmonary Disease Is Not Associated with KRAS Mutations in Non-Small Cell Lung Cancer. *PLoS ONE*, **2016**, 11, e0152317	3.7	8
89	Small RNA sequencing reveals a comprehensive miRNA signature of BRCA1-associated high-grade serous ovarian cancer. *Journal of Clinical Pathology*, **2016**,	3.9	8
88	Circulating miRNAs in patients with Barrett's esophagus, high-grade dysplasia and esophageal adenocarcinoma. *Journal of Gastrointestinal Oncology*, **2018**, 9, 1150-1156	2.8	8
87	Pim1 kinase activity preserves airway epithelial integrity upon house dust mite exposure. *American Journal of Physiology - Lung Cellular and Molecular Physiology*, **2015**, 309, L1344-53	5.8	7
86	Frequent lack of translation of antigen presentation-associated molecules MHC class I, CD1a and Beta(2)-microglobulin in Reed-Sternberg cells. *International Journal of Cancer*, **2000**, 86, 548-52	7.5	7
85	Ordering of polymorphic markers in the chromosome region 3p21. *Cytogenetic and Genome Research*, **1996**, 72, 225-8	1.9	7
84	Hyaline vascular Castleman's disease: a case report and brief review of the literature. *Netherlands Journal of Medicine*, **2002**, 60, 444-7	0.5	7
83	Argonaute 2 immunoprecipitation revealed large tumor suppressor kinase 1 as a novel proapoptotic target of miR-21 in T cells. *FEBS Journal*, **2017**, 284, 555-567	5.7	6
82	WEE1 Inhibition Enhances Anti-Apoptotic Dependency as a Result of Premature Mitotic Entry and DNA Damage. *Cancers*, **2019**, 11,	6.6	6
81	Autoimmune lymphoproliferative syndrome in a patient with a new minimal deletion in the death domain of the FAS gene. *Human Pathology*, **2008**, 39, 137-41	3.7	6
80	TIMP-1 expression in anaplastic large cell lymphoma is usually restricted to macrophages and only seldom observed in tumour cells. *Journal of Pathology*, **2005**, 206, 445-50	9.4	6

79	HLA expression and HLA type associations in relation to EBV status in Hispanic Hodgkin lymphoma patients. *PLoS ONE*, **2017**, 12, e0174457	3.7	6
78	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. *European Journal of Radiology*, **2021**, 138, 109646	4.7	6
77	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,	6.3	6
76	Tuberous sclerosis complex is required for tumor maintenance in MYC-driven Burkitt's lymphoma. *EMBO Journal*, **2018**, 37,	13	6
75	Serum periostin does not reflect type 2-driven inflammation in COPD. *Respiratory Research*, **2018**, 19, 112	7.3	6
74	Molecular imaging in lymphoma beyond F-FDG-PET: understanding the biology and its implications for diagnostics and therapy. *Lancet Haematology,the*, **2020**, 7, e479-e489	14.6	5
73	WEE1 inhibition synergizes with CHOP chemotherapy and radiation therapy through induction of premature mitotic entry and DNA damage in diffuse large B-cell lymphoma. *Therapeutic Advances in Hematology*, **2020**, 11, 2040620719898373	5.7	5
72	PML nuclear bodies and SATB1 are associated with HLA class I expression in EBV+ Hodgkin lymphoma. *PLoS ONE*, **2013**, 8, e72930	3.7	5
71	Pathogenesis and medical therapy of primary sclerosing cholangitis. Any news?. *European Journal of Gastroenterology and Hepatology*, **1999**, 11, 121-4	2.2	5
70	A PCR-aided transcript titration assay revealing very low expression of a gene at band 3p21 in 33 cells lines derived from all types of lung cancer. *European Journal of Human Genetics*, **1993**, 1, 156-63	5.3	5
69	Proteomics Based Identification of Proteins with Deregulated Expression in B Cell Lymphomas. *PLoS ONE*, **2016**, 11, e0146624	3.7	5
68	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	5.8	4
67	Seriai analysis of gene expression: rapid RT-PCR analysis of unknown SAGE tags. *Nucleic Acids Research*, **1999**, 27, i-iii	20.1	4
66	Tumour necrosis as assessed with F-FDG PET is a potential prognostic marker in diffuse large B cell lymphoma independent of MYC rearrangements. *European Radiology*, **2019**, 29, 6018-6028	8	3
65	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. *PLoS ONE*, **2019**, 14, e0223260	3.7	3
64	Genetic analysis of 2 cases of clear cell renal cancer in 2 sisters. *International Journal of Cancer*, **1998**, 77, 494-7	7.5	3
63	Cell-of-origin classification using the Hans and Lymph2Cx algorithms in primary cutaneous large B-cell lymphomas.. *Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin*, **2022**, 480, 667	5.1	3
62	Mir-155 Enhances B-Cell Lymphoma Growth By Targeting TBRG1. *Blood*, **2015**, 126, 4820-4820	2.2	3

61	A High Throughput Experimental Approach to Identify miRNA Target Genes in Hodgkin Lymphoma.. *Blood*, **2008**, 112, 1461-1461	2.2	3
60	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	7.3	3
59	Frequent mutated B2M, EZH2, IRF8, and TNFRSF14 in primary bone diffuse large B-cell lymphoma reflect a GCB phenotype. *Blood Advances*, **2021**, 5, 3760-3775	7.8	3
58	Combined PD-1 and JAK1/2 inhibition in refractory primary mediastinal B-cell lymphoma. *Annals of Hematology*, **2018**, 97, 905-907	3	2
57	Lymphadenopathy driven by TCR-V8V1 T-cell expansion in FAS-related autoimmune lymphoproliferative syndrome. *Blood Advances*, **2017**, 1, 1101-1106	7.8	2
56	131 Are chromosome 3 deletions a consistent abnormality in mesotheliomas?. *Cancer Genetics and Cytogenetics*, **1989**, 38, 203		2
55	Long Non-Coding RNAs Are Commonly Deregulated In Hodgkin Lymphoma. *Blood*, **2013**, 122, 628-628	2.2	2
54	Immune Dysfunction in Classical Hodgkin Lymphoma **2004**, 315-334		2
53	An All-In-One Transcriptome-Based Assay to Identify Therapy-Guiding Genomic Aberrations in Nonsmall Cell Lung Cancer Patients. *Cancers*, **2020**, 12,	6.6	2
52	Interaction between ERAP Alleles and HLA Class I Types Support a Role of Antigen Presentation in Hodgkin Lymphoma Development. *Cancers*, **2021**, 13,	6.6	2
51	Clinical Value of EGFR Copy Number Gain Determined by Amplicon-Based Targeted Next Generation Sequencing in Patients with EGFR-Mutated NSCLC. *Targeted Oncology*, **2021**, 16, 215-226	5	2
50	Actionability of on-target ALK Resistance Mutations in Patients With Non-Small Cell Lung Cancer: Local Experience and Review of the Literature. *Clinical Lung Cancer*, **2021**,	4.9	2
49	Identification of the estrogen receptor beta as a possible new tamoxifen-sensitive target in diffuse large B-cell lymphoma.. *Blood Cancer Journal*, **2022**, 12, 36	7	2
48	NGS-Based High-Throughput Screen to Identify MicroRNAs Regulating Growth of B-Cell Lymphoma. *Methods in Molecular Biology*, **2019**, 1956, 269-282	1.4	1
47	Multiple HLA class I and II associations in classical Hodgkin lymphoma and EBV status defined subgroups. *Blood*, **2011**, 118, 5211-7	2.2	1
46	5'-UTR FAS mutation pattern and frequency in nodal diffuse large B-cell lymphoma similar to normal germinal center B cells. *Leukemia and Lymphoma*, **2007**, 48, 851-2	1.9	1
45	A new RFLP identified at the D3S48 locus. *Nucleic Acids Research*, **1991**, 19, 4797	20.1	1
44	Understanding the role of long-acting muscarinic antagonists in asthma treatment.. *Annals of Allergy, Asthma and Immunology*, **2022**,	3.2	1

43	Killer Cell Immunoglobulin-Like Receptor Haplotype B Modulates Susceptibility to EBV-Associated Classic Hodgkin Lymphoma.. *Frontiers in Immunology*, **2022**, 13, 829943	8.4	1
42	High Expression of Micro-RNA BIC / miR155 in All Subtypes of Hodgkin Lymphoma.. *Blood*, **2004**, 104, 430-430	2.2	1
41	Polymorphisms and Lack of or Aberrant Expression of HLA Class I and II May Influence Antigen Presentation in Classical Hodgkin Lymphoma.. *Blood*, **2005**, 106, 20-20	2.2	1
40	Mid-Treatment Plasma Levels of Thymus Activated and Regulated Chemokine (TARC) Predict Treatment Outcome In Classical Hodgkin Lymphoma Patients. *Blood*, **2010**, 116, 748-748	2.2	1
39	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	2.2	1
38	Application of RIP-Chip for the Identification of miRNA Targets. *Neuromethods*, **2011**, 159-169	0.4	1
37	Microenvironment, Cross-Talk, and Immune Escape Mechanisms **2011**, 49-61		1
36	MiR-378a-3p Is Critical for Burkitt Lymphoma Cell Growth. *Cancers*, **2020**, 12,	6.6	1
35	Soluble PD-L1 is a promising disease biomarker but does not reflect tissue expression in classic Hodgkin lymphoma. *British Journal of Haematology*, **2021**, 193, 506-514	4.5	1
34	The relation between age and airway epithelial barrier function.. *Respiratory Research*, **2022**, 23, 43	7.3	1
33	Frequent 4EBP1 Amplification Induces Synthetic Dependence on FGFR Signaling in Cancer. *Cancers*, **2022**, 14, 2397	6.6	1
32	Management of Asthma with ICS and LABAs: Different treatment strategies. *Clinical Medicine Therapeutics*, **2009**, 1, CMT.S2283		0
31	CD4+ T cells in classical Hodgkin lymphoma express exhaustion associated transcription factors TOX and TOX2: Characterizing CD4+ T cells in Hodgkin lymphoma.. *OncoImmunology*, **2022**, 11, 2033433	7.2	0
30	The Microbiome in Bronchial Biopsies from Smokers and Ex-Smokers with Stable COPD - A Metatranscriptomic Approach.. *COPD: Journal of Chronic Obstructive Pulmonary Disease*, **2022**, 1-7	2	0
29	Targeting the Microenvironment in Hodgkin Lymphoma: Opportunities and Challenges. *Molecular Pathology Library*, **2018**, 59-90		
28	Long Non-coding RNAs in the Development and Maintenance of Lymphoid Malignancies **2019**, 127-149		
27	Microenvironment, Crosstalk, and Immune Escape Mechanisms. *Hematologic Malignancies*, **2015**, 65-78	0	
26	The Role of MicroRNAs in Hodgkin's Lymphoma **2013**, 435-447		

25	Deletion of C-MYC in the spectrum of transformed follicular lymphoma grade 3B: reply. *Human Pathology*, **2007**, 38, 192-193	3.7
24	Low Mutational Burden of Extra Nodal Marginal Zone Lymphoma of Mucosa-Associated Lymphoid Tissue in Patients with Primary Sjogren's Syndrome. *Blood*, **2021**, 138, 1331-1331	2.2
23	Responsivity and Reproducibility of Sputum Inflammatory Biomarkers During COPD Exacerbation and Stable Phases - A Pilot Study. *International Journal of COPD*, **2021**, 16, 3055-3064	3
22	Serial Analysis of Gene Expression Revealed Consistent Downregulation of More Than 100 Genes in Hodgkin Lymphoma.. *Blood*, **2004**, 104, 4288-4288	2.2
21	Haplotype-Based Sequencing To Delineate the Associated HLA Class I Region for EBV Positive Hodgkin Lymphoma.. *Blood*, **2005**, 106, 971-971	2.2
20	Lack of BIC and microRNA miR-155 Expression in Primary Cases of Burkitt Lymphoma.. *Blood*, **2005**, 106, 1922-1922	2.2
19	Regulation of pri-miRNA BIC Transcription and Processing in Burkitt Lymphoma.. *Blood*, **2006**, 108, 2380-2380	2.2
18	Discovery of Biomarkers for Hodgkin Lymphoma Using Proteomics Technology.. *Blood*, **2006**, 108, 2261-2261	2.2
17	miRNA Expression Profile of B-SLL Consistent with Normal Memory B Cells:BIC/miR155 Specific Location in Proliferation Center.. *Blood*, **2007**, 110, 2081-2081	2.2
16	Microenvironment, Cross-Talk, and Immune Escape Mechanisms. *Hematologic Malignancies*, **2020**, 69-86	0
15	Meta-Analysis of Hodgkin Lymphoma and Asthma Genome-Wide Association Scans reveals common variants in GATA3. *Blood*, **2014**, 124, 135-135	2.2
14	Mutations in CD58 and Other Immune System Related Genes in Hodgkin Lymphoma. *Blood*, **2015**, 126, 1439-1439	2.2
13	Impact of Radiotherapy and Rituximab on Outcome of Stage I(E) Diffuse Large B-Cell Lymphoma. *Blood*, **2015**, 126, 1503-1503	2.2
12	Expression of CD1c, CD1d and Presence of Invariant NKT Cells in Hodgkin Lymphoma.. *Blood*, **2009**, 114, 3659-3659	2.2
11	The HGF/c-Met Signaling Pathway in Hodgkin Lymphoma.. *Blood*, **2009**, 114, 1551-1551	2.2
10	Serum TARC Levels Correspond with Tumor Load and Response to Chemotherapy in Classical Hodgkin Lymphoma Patients.. *Blood*, **2009**, 114, 3663-3663	2.2
9	Expression of the c-Met Oncogene Correlates with Favorable Progression Free Survival In Classical Hodgkin Lymphoma. *Blood*, **2010**, 116, 3880-3880	2.2
8	HLA Class I and EBV Positive Classical Hodgkin Lymphoma In the Chinese Population. *Blood*, **2010**, 116, 2688-2688	2.2

7	Protective and Predisposing HLA Alleles In Dutch Classical Hodgkin Lymphoma Patients. *Blood*, **2010**, 116, 749-749	2.2
6	CSF1R Expression of Hodgkin Reed Sternberg Cells Is Associated with the Number of Macrophages in the Tumor Microenvironment and Is Correlated with Treatment Outcome. *Blood*, **2011**, 118, 427-427	2.2
5	CCR4 Expression in Hodgkin Lymphoma. *Blood*, **2011**, 118, 2626-2626	2.2
4	Expression, Prognostic Value and Function of IGF1R in Hodgkin Lymphoma.. *Blood*, **2012**, 120, 2640-2640	2.2
3	Long Non-Coding RNAs As Components Of The MYC Network In B Cell Lymphoma. *Blood*, **2013**, 122, 1260-1260	2.2
2	A Meta-Analysis Of Hodgkin Lymphoma Reveals 19p13.3 (TCF3) As a Novel Susceptibility Loc. *Blood* , **2013**, 122, 626-626	2.2
1	GATA6 expression in Barrett's metaplasia development and malignant progression.. *Journal of Clinical Oncology*, **2014**, 32, 26-26	2.2