

# Lydia Visser

## List of Publications by Year in descending order

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83  
papers

2,362  
citations

201674

27  
h-index

223800

46  
g-index

85  
all docs

85  
docs citations

85  
times ranked

3476  
citing authors

#	ARTICLE	IF	CITATIONS
1	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, 240-253.	6.3	141
2	HLA-A*02 is associated with a reduced risk and HLA-A*01 with an increased risk of developing EBV+ Hodgkin lymphoma. <i>Blood</i> , 2007, 110, 3310-3315.	1.4	131
3	Plasma vesicle miRNAs for therapy response monitoring in Hodgkin lymphoma patients. <i>JCI Insight</i> , 2016, 1, e89631.	5.0	121
4	Proteomics analysis of Hodgkin lymphoma: identification of new players involved in the cross-talk between HRS cells and infiltrating lymphocytes. <i>Blood</i> , 2008, 111, 2339-2346.	1.4	114
5	Serum chemokine levels in Hodgkin lymphoma patients: highly increased levels of CCL17 and CCL22. <i>British Journal of Haematology</i> , 2008, 140, 527-536.	2.5	110
6	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
7	Morphologic, immunologic, enzyme histochemical and chromosomal analysis of a cell line derived from Hodgkin's disease. Evidence for a B-cell origin of sternberg-reed cells. <i>Cancer</i> , 1985, 55, 683-690.	4.1	95
8	Strongly enhanced IL-10 production using stable galectin-1 homodimers. <i>Molecular Immunology</i> , 2007, 44, 506-513.	2.2	93
9	Dimeric galectin-1 induces IL-10 production in T-lymphocytes: an important tool in the regulation of the immune response. <i>Journal of Pathology</i> , 2004, 204, 511-518.	4.5	87
10	HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?. <i>Oncimmunology</i> , 2017, 6, e1295202.	4.6	84
11	Long noncoding RNAs as a novel component of the Myc transcriptional network. <i>FASEB Journal</i> , 2015, 29, 2338-2346.	0.5	67
12	Common and differential chemokine expression patterns in rs cells of NLP, EBV positive and negative classical hodgkin lymphomas. <i>International Journal of Cancer</i> , 2002, 99, 665-672.	5.1	66
13	The CD4+CD26 <sup>hi</sup> T-cell population in classical Hodgkin's lymphoma displays a distinctive regulatory T-cell profile. <i>Laboratory Investigation</i> , 2008, 88, 482-490.	3.7	62
14	Plasma thymus and activation-regulated chemokine as an early response marker in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012, 97, 410-415.	3.5	56
15	HLA Associations in Classical Hodgkin Lymphoma: EBV Status Matters. <i>PLoS ONE</i> , 2012, 7, e39986.	2.5	52
16	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
17	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
18	Biomarkers for evaluation of treatment response in classical Hodgkin lymphoma: comparison of $\alpha$ -galactin-1, $\alpha$ -CD163 and $\alpha$ -CD30 with TARC. <i>British Journal of Haematology</i> , 2016, 175, 868-875.	2.5	44

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19	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. <i>Oncotarget</i> , 2016, 7, 2391-2400.	1.8	43
20	CD45 (leucocyte common antigen) expression in T and B lymphocyte subsets. <i>Leukemia and Lymphoma</i> , 1996, 20, 217-222.	1.3	42
21	Cytokine gene expression profile distinguishes CD4+/CD57+ T cells of the nodular lymphocyte predominance type of Hodgkin's lymphoma from their tonsillar counterparts. <i>Journal of Pathology</i> , 2006, 208, 423-430.	4.5	41
22	The Microenvironment in Epstein-Barr Virus-Associated Malignancies. <i>Pathogens</i> , 2018, 7, 40.	2.8	40
23	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cell-Specific Long Noncoding RNAs. <i>American Journal of Pathology</i> , 2016, 186, 2462-2472.	3.8	36
24	Enrichment of the tumour immune microenvironment in patients with desmoplastic colorectal liver metastasis. <i>British Journal of Cancer</i> , 2020, 123, 196-206.	6.4	35
25	Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma. <i>Cancer Treatment Reviews</i> , 2020, 82, 101931.	7.7	33
26	ADAM17 up-regulation in renal transplant dysfunction and non-transplant-related renal fibrosis. <i>Nephrology Dialysis Transplantation</i> , 2012, 27, 2114-2122.	0.7	31
27	Expression of the c-Met oncogene by tumor cells predicts a favorable outcome in classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012, 97, 572-578.	3.5	29
28	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
29	HLA-A*02:07 Is a Protective Allele for EBV Negative and a Susceptibility Allele for EBV Positive Classical Hodgkin Lymphoma in China. <i>PLoS ONE</i> , 2012, 7, e31865.	2.5	25
30	Expression of CD1d and presence of invariant NKT cells in classical Hodgkin lymphoma. <i>American Journal of Hematology</i> , 2010, 85, 539-541.	4.1	24
31	Epstein-barr virus positivity in hodgkin's disease does not correlate with an hla a2-negative phenotype. <i>Cancer</i> , 1994, 73, 3059-3063.	4.1	23
32	Paediatric nodal marginal zone B-cell lymphadenopathy of the neck: a <i>Haemophilus influenzae</i> -driven immune disorder?. <i>Journal of Pathology</i> , 2015, 236, 302-314.	4.5	23
33	Characterization of the Microenvironment of Nodular Lymphocyte Predominant Hodgkin Lymphoma. <i>International Journal of Molecular Sciences</i> , 2016, 17, 2127.	4.1	23
34	Insulin-Like Growth Factor 1 Receptor Is a Prognostic Factor in Classical Hodgkin Lymphoma. <i>PLoS ONE</i> , 2014, 9, e87474.	2.5	22
35	Neoplastic Changes Involving Follicles: Morphological, Immunophenotypic and Genetic Diversity of Lymphoproliferations Derived from Germinal Center and Mantle Zone. <i>Immunological Reviews</i> , 1992, 126, 163-178.	6.0	21
36	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.	1.6	20

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37	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.	2.5	20
38	Heparin binding epidermal growth factor in renal ischaemia/reperfusion injury. Journal of Pathology, 2010, 221, 183-192.	4.5	19
39	Induction of B-cell chronic lymphocytic leukaemia and hairy cell leukaemia like phenotypes by phorbol ester treatment of normal peripheral blood B-cells. British Journal of Haematology, 1990, 75, 359-365.	2.5	18
40	Mechanisms of induction of renal allograft tolerance in CD45RB-treated mice. Kidney International, 1999, 55, 1303-1310.	5.2	17
41	Expression of HLA Class I and HLA Class II by Tumor Cells in Chinese Classical Hodgkin Lymphoma Patients. PLoS ONE, 2010, 5, e10865.	2.5	16
42	Interim thymus and activation regulated chemokine versus interim 18 F-fluorodeoxyglucose positron emission tomography in classical Hodgkin lymphoma response evaluation. British Journal of Haematology, 2020, 190, 40-44.	2.5	15
43	Postnatal changes of CD45 expression in peripheral blood T and B cells. British Journal of Haematology, 1994, 87, 251-257.	2.5	13
44	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
45	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
46	Prolonged survival of rat islet xenografts in mice after CD45RB monotherapy. Transplantation, 2004, 77, 386-391.	1.0	12
47	WEE1 Inhibition Enhances Anti-Apoptotic Dependency as a Result of Premature Mitotic Entry and DNA Damage. Cancers, 2019, 11, 1743.	3.7	12
48	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, 204062071989837.	2.5	12
49	Comparison of the ZAP70+ and ZAP70 <sup>~</sup> B-CLL Kinome: Higher Kinase Activity in ZAP70+ B-CLL Cells.. Blood, 2006, 108, 2804-2804.	1.4	11
50	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.	2.5	9
51	Gene expression-based model predicts outcome in children with intermediate-risk classical Hodgkin lymphoma. Blood, 2021, , .	1.4	9
52	CD4+ T cells in classical Hodgkin lymphoma express exhaustion associated transcription factors TOX and TOX2. Oncolmmunology, 2022, 11, 2033433.	4.6	9
53	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.	6.2	8
54	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

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55	B Cells as Prognostic Biomarker After Surgery for Colorectal Liver Metastases. <i>Frontiers in Oncology</i> , 2020, 10, 249.	2.8	7
56	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
57	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
58	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
59	Patterns of Leucocyte Common Antigen Expression in Peripheral Blood T Cell Populations. <i>Cellular Immunology</i> , 1993, 151, 218-224.	3.0	5
60	PML Nuclear Bodies and SATB1 Are Associated with HLA Class I Expression in EBV+ Hodgkin Lymphoma. <i>PLoS ONE</i> , 2013, 8, e72930.	2.5	5
61	Combined loss of <scp>HLA</scp> I and <scp>HLA II</scp> expression is more common in the nonâ€<scp>GCB</scp> type of diffuse large B cell lymphoma. <i>Histopathology</i> , 2018, 72, 886-888.	2.9	4
62	Mid-Treatment Plasma Levels of Thymus Activated and Regulated Chemokine (TARC) Predict Treatment Outcome In Classical Hodgkin Lymphoma Patients. <i>Blood</i> , 2010, 116, 748-748.	1.4	4
63	Validation of Novel Molecular Imaging Targets Identified by Functional Genomic mRNA Profiling to Detect Dysplasia in Barrettâ€™s Esophagus. <i>Cancers</i> , 2022, 14, 2462.	3.7	4
64	Combined PD-1 and JAK1/2 inhibition in refractory primary mediastinal B-cell lymphoma. <i>Annals of Hematology</i> , 2018, 97, 905-907.	1.8	3
65	The Nature of the Lymphocytes in Hodgkinâ€™s Disease. , 1995, , 161-171.		3
66	Evidence for a B-cell origin of the proliferating cells. <i>Cancer Treatment and Research</i> , 1989, 41, 5-27.	0.5	3
67	Computational study, synthesis and evaluation of active peptides derived from Parasporin-2 and spike protein from Alphacoronavirus against colon cancer cells. <i>Bioscience Reports</i> , 2021, 41, .	2.4	3
68	Microenvironment, Cross-Talk, and Immune Escape Mechanisms. , 2011, , 49-61.		1
69	The HGF/c-Met Signaling Pathway in Hodgkin Lymphoma.. <i>Blood</i> , 2009, 114, 1551-1551.	1.4	1
70	Protective and Predisposing HLA Alleles In Dutch Classical Hodgkin Lymphoma Patients. <i>Blood</i> , 2010, 116, 749-749.	1.4	1
71	Microenvironment, Cross-Talk, and Immune Escape Mechanisms. <i>Hematologic Malignancies</i> , 2020, , 69-86.	0.2	1
72	Genetic Modification Approaches for Parasporins Bacillus thuringiensis Proteins with Anticancer Activity. <i>Molecules</i> , 2021, 26, 7476.	3.8	1

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73	Anti-CD37 radioimmunotherapy with <sup>177</sup> Lu-NNV003 synergizes with the PARP inhibitor olaparib in treatment of non-Hodgkin's lymphoma in vitro. PLoS ONE, 2022, 17, e0267543.	2.5	1
74	Microenvironment, Crosstalk, and Immune Escape Mechanisms. Hematologic Malignancies, 2015, , 65-78.	0.2	0
75	Targeting the Microenvironment in Hodgkin Lymphoma: Opportunities and Challenges. Molecular Pathology Library, 2018, , 59-90.	0.1	0
76	Plasma cells in classical Hodgkin lymphoma: a new player in the microenvironment?. British Journal of Haematology, 2019, 184, 119-120.	2.5	0
77	miRNA Expression Profile of B-SLL Consistent with Normal Memory B Cells:BIC/miR-155 Specific Location in Proliferation Center.. Blood, 2007, 110, 2081-2081.	1.4	0
78	Expression of CD1c, CD1d and Presence of Invariant NKT Cells in Hodgkin Lymphoma.. Blood, 2009, 114, 3659-3659.	1.4	0
79	Expression of the c-Met Oncogene Correlates with Favorable Progression Free Survival In Classical Hodgkin Lymphoma. Blood, 2010, 116, 3880-3880.	1.4	0
80	HLA Class I and EBV Positive Classical Hodgkin Lymphoma In the Chinese Population. Blood, 2010, 116, 2688-2688.	1.4	0
81	CCR4 Expression in Hodgkin Lymphoma. Blood, 2011, 118, 2626-2626.	1.4	0
82	EBV and HLA Associations In Classical Hodgkin Lymphoma Patients From Brazil. Blood, 2011, 118, 4858-4858.	1.4	0
83	Strong HLA Class I Expression Is Positively Correlated with the Number of PML Nuclear Bodies and Negatively with the Percentage of SATB1 Positive HRS Cells in EBV+ Classical Hodgkin Lymphoma (cHL). Blood, 2012, 120, 3633-3633.	1.4	0