

Ruth F Jarrett

List of Publications by Year in descending order

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100
papers

7,758
citations

101384

36
h-index

62479

80
g-index

103
all docs

103
docs citations

103
times ranked

10775
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluating the Effects of SARS-CoV-2 Spike Mutation D614G on Transmissibility and Pathogenicity. <i>Cell</i> , 2021, 184, 64-75.e11.	13.5	843
2	HTLV-III expression and production involve complex regulation at the levels of splicing and translation of viral RNA. <i>Cell</i> , 1986, 46, 807-817.	13.5	832
3	Human Tissues Contain CD141hi Cross-Presenting Dendritic Cells with Functional Homology to Mouse CD103+ Nonlymphoid Dendritic Cells. <i>Immunity</i> , 2012, 37, 60-73.	6.6	643
4	Circulating SARS-CoV-2 spike N439K variants maintain fitness while evading antibody-mediated immunity. <i>Cell</i> , 2021, 184, 1171-1187.e20.	13.5	541
5	Epstein-Barr virus-associated Hodgkin's disease: Epidemiologic characteristics in international data. , 1997, 70, 375-382.		424
6	Mutations in the I κ B α gene in Hodgkin's disease suggest a tumour suppressor role for I κ B α . <i>Oncogene</i> , 1999, 18, 3063-3070.	2.6	330
7	Long-term inhibition of human T-lymphotropic virus type III/lymphadenopathy-associated virus (human) Tj ETQq1 1 0.784314 rgBT /Over 2',3'-dideoxynucleosides in vitro.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1987, 84, 2033-2037.	3.3	203
8	Impact of tumor Epstein-Barr virus status on presenting features and outcome in age-defined subgroups of patients with classic Hodgkin lymphoma: a population-based study. <i>Blood</i> , 2005, 106, 2444-2451.	0.6	184
9	A genome-wide association study of Hodgkin's lymphoma identifies new susceptibility loci at 2p16.1 (REL), 8q24.21 and 10p14 (GATA3). <i>Nature Genetics</i> , 2010, 42, 1126-1130.	9.4	177
10	Detection of Epstein-Barr virus (EBV) genomes in the serum of patients with EBV-associated Hodgkin's disease. , 1999, 84, 442-448.		146
11	Detection of human herpesvirus-6 DNA in peripheral blood and saliva. <i>Journal of Medical Virology</i> , 1990, 32, 73-76.	2.5	141
12	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.	3.0	141
13	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.	0.6	131
14	Prevalence of Antibody to Human Herpesvirus 7 by Age. <i>Journal of Infectious Diseases</i> , 1993, 168, 251-252.	1.9	128
15	A prenylated dsRNA sensor protects against severe COVID-19. <i>Science</i> , 2021, 374, eabj3624.	6.0	124
16	Risk factors for Hodgkin's disease by Epstein-Barr virus (EBV) status: prior infection by EBV and other agents. <i>British Journal of Cancer</i> , 2000, 82, 1117-1121.	2.9	116
17	1 The epidemiology of Hodgkin's disease. <i>Best Practice and Research: Clinical Haematology</i> , 1996, 9, 401-416.	1.1	111
18	Serum chemokine levels in Hodgkin lymphoma patients: highly increased levels of CCL17 and CCL22. <i>British Journal of Haematology</i> , 2008, 140, 527-536.	1.2	110

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19	“Cytomegalovirus disease” in renal allograft recipients: Is human herpesvirus 7 a co-factor for disease progression?. , 1996, 48, 295-301.		105
20	HLA-A alleles and infectious mononucleosis suggest a critical role for cytotoxic T-cell response in EBV-related Hodgkin lymphoma. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 6400-6405.	3.3	102
21	HHV-8 and multiple myeloma in the UK. Lancet, The, 1997, 350, 1144-1145.	6.3	95
22	HLA class I polymorphisms are associated with development of infectious mononucleosis upon primary EBV infection. Journal of Clinical Investigation, 2007, 117, 3042-3048.	3.9	92
23	Viruses and lymphoma/leukaemia. Journal of Pathology, 2006, 208, 176-186.	2.1	89
24	Viral involvement in Hodgkin's disease: detection of clonal type A Epstein-Barr virus genomes in tumour samples. British Journal of Cancer, 1991, 64, 227-232.	2.9	85
25	Association Between Simian Virus 40 DNA and Lymphoma in the United Kingdom. Journal of the National Cancer Institute, 2003, 95, 1001-1003.	3.0	85
26	Mutations of <i>NFKBIA</i> , encoding I κ B α , are a recurrent finding in classical Hodgkin lymphoma but are not a unifying feature of non-EBV-associated cases. International Journal of Cancer, 2009, 125, 1334-1342.	2.3	85
27	The seroepidemiology of human herpesvirus-6 (HHV-6) from a case-control study of leukaemia and lymphoma. International Journal of Cancer, 1990, 45, 829-833.	2.3	84
28	The molecular pathogenesis of Hodgkin lymphoma. Histopathology, 2011, 58, 15-25.	1.6	74
29	Sensitive ELISA for the gp120 and gp160 Surface Glycoproteins of HIV-1. AIDS Research and Human Retroviruses, 1988, 4, 369-379.	0.5	69
30	Primary Epstein-Barr virus infection with and without infectious mononucleosis. PLoS ONE, 2019, 14, e0226436.	1.1	67
31	Nucleotide sequence of transforming human c-srcDNA clones with homology to platelet-derived growth factor. Nucleic Acids Research, 1985, 13, 5007-5018.	6.5	63
32	Detection of SARS-CoV-2 in respiratory samples from cats in the UK associated with human-to-cat transmission. Veterinary Record, 2021, 188, e247.	0.2	63
33	Variation at 3p24.1 and 6q23.3 influences the risk of Hodgkin's lymphoma. Nature Communications, 2013, 4, 2549.	5.8	62
34	Risk Factors for Hodgkin's Lymphoma by EBV Status and Significance of Detection of EBV Genomes in Serum of Patients with EBV-Associated Hodgkin's Lymphoma. Leukemia and Lymphoma, 2003, 44, S27-S32.	0.6	55
35	An epidemiologic study of index and family infectious mononucleosis and adult Hodgkin's disease (HD): Evidence for a specific association with EBV+ve HD in young adults. International Journal of Cancer, 2003, 107, 298-302.	2.3	51
36	Validation of the safety of MDCK cells as a substrate for the production of a cell-derived influenza vaccine. Biologicals, 2010, 38, 544-551.	0.5	51

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37	Classical Hodgkin lymphoma is associated with frequent gains of 17q. <i>Genes Chromosomes and Cancer</i> , 2003, 38, 126-136.	1.5	49
38	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. <i>Nature Communications</i> , 2017, 8, 1892.	5.8	40
39	Hodgkin lymphoma and Epstein-Barr virus (EBV): No evidence to support hit-and-run mechanism in cases classified as non-EBV-associated. <i>International Journal of Cancer</i> , 2003, 104, 624-630.	2.3	38
40	The Human Leukocyte Antigen Class I Region Is Associated with EBV-Positive Hodgkin's Lymphoma: HLA-A and HLA Complex Group 9 Are Putative Candidate Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 2280-2284.	1.1	36
41	Human leukocyte antigens and genetic susceptibility to lymphoma. <i>Tissue Antigens</i> , 2015, 86, 98-113.	1.0	36
42	Inherited Chromosomally Integrated Human Herpesvirus 6 Genomes Are Ancient, Intact, and Potentially Able To Reactivate from Telomeres. <i>Journal of Virology</i> , 2017, 91, .	1.5	36
43	Viruses and Hodgkin disease: No evidence of novel herpesviruses in non-EBV-associated lesions. <i>International Journal of Cancer</i> , 2002, 101, 259-264.	2.3	35
44	Effect of IL-6 promoter polymorphism on incidence and outcome in Hodgkin's lymphoma. <i>British Journal of Haematology</i> , 2005, 128, 493-495.	1.2	35
45	Lack of involvement of known oncogenic DNA viruses in Epstein-Barr virus-negative Hodgkin's disease. <i>British Journal of Cancer</i> , 1998, 77, 1045-1047.	2.9	33
46	The expression of the EBV latent membrane protein (LMP1) is independent of CD23 and bcl2 in Reed-Sternberg cells in Hodgkin's disease. <i>Histopathology</i> , 1992, 21, 72-73.	1.6	31
47	JC and BK virus sequences are not detectable in leukaemic samples from children with common acute lymphoblastic leukaemia. <i>British Journal of Cancer</i> , 1999, 81, 898-899.	2.9	30
48	Modeling HLA associations with EBV-positive and -negative Hodgkin lymphoma suggests distinct mechanisms in disease pathogenesis. <i>International Journal of Cancer</i> , 2015, 137, 1066-1075.	2.3	30
49	Optimisation and validation of a PCR for antigen receptor rearrangement (PARR) assay to detect clonality in canine lymphoid malignancies. <i>Veterinary Immunology and Immunopathology</i> , 2016, 182, 115-124.	0.5	30
50	POSTTRANSPLANT LYMPHOPROLIFERATIVE DISORDER ASSOCIATED WITH PRIMATE GAMMA-HERPESVIRUS IN CYNOMOLGUS MONKEYS USED IN PIG-TO-PRIMATE RENAL XENOTRANSPLANTATION AND PRIMATE RENAL ALLOTRANSPLANTATION. <i>Transplantation</i> , 2002, 73, 44-52.	0.5	28
51	Measles virus and classical Hodgkin lymphoma: No evidence for a direct association. <i>International Journal of Cancer</i> , 2007, 121, 442-447.	2.3	25
52	Expression and function of T cell homing molecules in Hodgkin's lymphoma. <i>Cancer Immunology, Immunotherapy</i> , 2009, 58, 85-94.	2.0	22
53	Germ-Line Transmitted, Chromosomally Integrated HHV-6 and Classical Hodgkin Lymphoma. <i>PLoS ONE</i> , 2014, 9, e112642.	1.1	22
54	Phenotype and frequency of Epstein-Barr virus-infected cells in pretreatment blood samples from patients with Hodgkin lymphoma. <i>British Journal of Haematology</i> , 2005, 129, 511-519.	1.2	21

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55	HHV-8-unrelated primary effusion-like lymphoma associated with clonal loss of inherited chromosomally-integrated human herpesvirus-6A from the telomere of chromosome 19q. <i>Scientific Reports</i> , 2016, 6, 22730.	1.6	21
56	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.	1.1	20
57	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.	0.9	20
58	Configuration and expression of the T cell receptor beta chain gene in human T-lymphotropic virus I-infected cells.. <i>Journal of Experimental Medicine</i> , 1986, 163, 383-399.	4.2	19
59	Germline mutations and polymorphisms in the NFKBIA gene in Hodgkin lymphoma. <i>International Journal of Cancer</i> , 2005, 116, 646-651.	2.3	19
60	Viruses and Hodgkin lymphoma: No evidence of polyomavirus genomes in tumor biopsies. <i>Leukemia and Lymphoma</i> , 2006, 47, 1315-1321.	0.6	18
61	Analysis of Epstein-Barr virus (EBV) nuclear antigen 1 subtypes in EBV-associated lymphomas from Brazil and the United Kingdom. <i>Journal of General Virology</i> , 1999, 80, 2741-2745.	1.3	18
62	Analysis of T-cell receptor and immunoglobulin gene rearrangements in the diagnosis of Hodgkin's and non-Hodgkin's lymphoma. <i>Journal of Pathology</i> , 1990, 161, 245-254.	2.1	17
63	Infection of human T lymphotropic virus-I-specific immune T cell clones by human T lymphotropic virus-I.. <i>Journal of Clinical Investigation</i> , 1986, 78, 1302-1310.	3.9	16
64	Uterine perforation by a Copper 7 intrauterine contraceptive device with subsequent penetration of the appendix.. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1983, 90, 774-776.	1.1	15
65	Evaluation of the antibody response to the EBV proteome in EBV-associated classical Hodgkin lymphoma. <i>International Journal of Cancer</i> , 2020, 147, 608-618.	2.3	15
66	B-lymphotropic viruses in a novel tropical splenic lymphoma. <i>British Journal of Haematology</i> , 2001, 112, 161-166.	1.2	14
67	SLC6A4 expression and anti-proliferative responses to serotonin transporter ligands clomipramine and fluoxetine in primary B-cell malignancies. <i>Leukemia Research</i> , 2010, 34, 1103-1106.	0.4	14
68	HVMNE, a novel lymphocryptovirus related to Epstein-Barr virus, induces lymphoma in New Zealand White rabbits. <i>Blood</i> , 2001, 98, 2193-2199.	0.6	13
69	Mannose-Binding Lectin Genotypes and Susceptibility to Epstein-Barr Virus Infection in Infancy. <i>Vaccine Journal</i> , 2010, 17, 1484-1487.	3.2	13
70	A susceptibility locus for classical Hodgkin lymphoma at 8q24 near <i>MYC</i> predicts patient outcome in two independent cohorts. <i>British Journal of Haematology</i> , 2018, 180, 286-290.	1.2	13
71	Determination of HLA-A*02 antigen status in Hodgkin's disease and analysis of an HLA-A*02-restricted epitope of the Epstein-Barr virus LMP-2 protein. , 1997, 72, 614-618.		11
72	Genetically Raised Circulating Bilirubin Levels and Risk of Ten Cancers: A Mendelian Randomization Study. <i>Cells</i> , 2021, 10, 394.	1.8	11

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73	Epstein-Barr virus immune response in high-risk nasopharyngeal carcinoma families in Greenland. <i>Journal of Medical Virology</i> , 2007, 79, 1877-1881.	2.5	9
74	Letters to the editor. <i>Journal of Pathology</i> , 1991, 164, 345-346.	2.1	7
75	Viral involvement in Hodgkin's disease. <i>International Journal of Cell Cloning</i> , 1992, 10, 315-322.	1.6	7
76	Gammaherpesviruses and canine lymphoma: no evidence for direct involvement in commonly occurring lymphomas. <i>Journal of General Virology</i> , 2015, 96, 1863-1872.	1.3	7
77	The prevalence and characterisation of TRAF3 and POT1 mutations in canine B-cell lymphoma. <i>Veterinary Journal</i> , 2020, 266, 105575.	0.6	7
78	Hodgkin's disease. <i>Best Practice and Research: Clinical Haematology</i> , 1992, 5, 57-79.	1.1	6
79	The Retrovirus XMRV Is not Directly Involved in the Pathogenesis of Common Types of Lymphoid Malignancy. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2232-2236.	1.1	6
80	Molecular Methods of Virus Detection in Lymphoma. <i>Methods in Molecular Biology</i> , 2013, 971, 277-293.	0.4	6
81	Development of an Electrochemical CCL17/TARC Biosensor toward Rapid Triage and Monitoring of Classic Hodgkin Lymphoma. <i>ACS Sensors</i> , 2021, 6, 3262-3272.	4.0	5
82	Infectious mononucleosis, immune genotypes, and non-Hodgkin lymphoma (NHL): an InterLymph Consortium study. <i>Cancer Causes and Control</i> , 2020, 31, 451-462.	0.8	4
83	Epidemiology of Hodgkin Lymphoma. <i>Hematologic Malignancies</i> , 2020, , 3-23.	0.2	4
84	Demonstration of Epstein-Barr viral DNA in formalin-fixed, paraffin-embedded samples of Hodgkin's disease. <i>Journal of Pathology</i> , 1991, 163, 149-151.	2.1	3
85	Chapter 2 The epidemiology of human herpesvirus-6. <i>Perspectives in Medical Virology</i> , 1992, 4, 9-23.	0.1	3
86	Send cat and dog samples to test for SARS-CoV-2. <i>Veterinary Record</i> , 2020, 186, 571-571.	0.2	3
87	Detection of Epstein-Barr virus (EBV) genomes in the serum of patients with EBV-associated Hodgkin's disease. , 1999, 84, 442.		3
88	The Role of Viruses in the Genesis of Hodgkin Lymphoma. , 2011, , 21-32.		3
89	Evidence of an Increased Frequency of HLA-DPB1*0301 in Hodgkin's Disease Supports an Infectious Aetiology. , 1995, , 15-25.		2
90	Chapter 17 Detection of HHV-6 using polymerase chain reaction amplification. <i>Perspectives in Medical Virology</i> , 1992, 4, 227-234.	0.1	1

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91	Human herpesvirus-8. Perspectives in Medical Virology, 2001, 5, 253-290.	0.1	1
92	The Role of Viruses in the Genesis of Hodgkin Lymphoma. Hematologic Malignancies, 2020, , 25-45.	0.2	1
93	HLA Expression in Relation to HLA Type in Classic Hodgkin Lymphoma Patients. Cancers, 2021, 13, 5833.	1.7	1
94	Pathogenesis of retroviral infections. Journal of Pathology, 1987, 153, 199-200.	2.1	0
95	Haplotype-Based Sequencing To Delineate the Associated HLA Class I Region for EBV Positive Hodgkin Lymphoma.. Blood, 2005, 106, 971-971.	0.6	0
96	TARC and MDC Are the Only Chemokines with Highly Increased Levels in Serum of Hodgkin Lymphoma Patients.. Blood, 2006, 108, 2268-2268.	0.6	0
97	Risk of EBV-Positive Hodgkin Lymphoma Varies Over 30-Fold by HLA Class I Genotype and History of Infectious Mononucleosis.. Blood, 2009, 114, 269-269.	0.6	0
98	A SCID Mouse Model of Hodgkinâ€™s Disease? Transplantation of Hodgkinâ€™s and Non-Hodgkinâ€™s Lymphomas Into Severe Combined Immunodeficient Mice. , 1995, , 187-195.		0
99	The Role of Viruses in the Genesis of Hodgkin Lymphoma. Hematologic Malignancies, 2015, , 27-43.	0.2	0
100	Identifying Epstein-Barr virus peptide sequences associated with differential IgG antibody response. International Journal of Infectious Diseases, 2021, 114, 65-71.	1.5	0