

Jonathan K Ball

List of Publications by Year in descending order

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117
papers

5,219
citations

94433

37
h-index

91884

69
g-index

126
all docs

126
docs citations

126
times ranked

5146
citing authors

#	ARTICLE	IF	CITATIONS
1	Broadly neutralizing antibodies protect against hepatitis C virus quasispecies challenge. <i>Nature Medicine</i> , 2008, 14, 25-27.	30.7	556
2	Monoclonal Antibody AP33 Defines a Broadly Neutralizing Epitope on the Hepatitis C Virus E2 Envelope Glycoprotein. <i>Journal of Virology</i> , 2005, 79, 11095-11104.	3.4	262
3	Characterization of host-range and cell entry properties of the major genotypes and subtypes of hepatitis C virus. <i>Hepatology</i> , 2005, 41, 265-274.	7.3	234
4	Identification of Conserved Residues in the E2 Envelope Glycoprotein of the Hepatitis C Virus That Are Critical for CD81 Binding. <i>Journal of Virology</i> , 2006, 80, 8695-8704.	3.4	232
5	Human Adaptation of Ebola Virus during the West African Outbreak. <i>Cell</i> , 2016, 167, 1079-1087.e5.	28.9	180
6	Characterization of the hepatitis C virus E2 epitope defined by the broadly neutralizing monoclonal antibody AP33. <i>Hepatology</i> , 2006, 43, 592-601.	7.3	150
7	Broadly neutralizing human monoclonal antibodies to the hepatitis C virus E2 glycoprotein. <i>Journal of General Virology</i> , 2008, 89, 653-659.	2.9	144
8	Role of Scavenger Receptor Class B Type I in Hepatitis C Virus Entry: Kinetics and Molecular Determinants. <i>Journal of Virology</i> , 2010, 84, 34-43.	3.4	144
9	Genome-to-genome analysis highlights the effect of the human innate and adaptive immune systems on the hepatitis C virus. <i>Nature Genetics</i> , 2017, 49, 666-673.	21.4	129
10	Human combinatorial libraries yield rare antibodies that broadly neutralize hepatitis C virus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 16269-16274.	7.1	127
11	The past, present and future of neutralizing antibodies for hepatitis C virus. <i>Antiviral Research</i> , 2014, 105, 100-111.	4.1	125
12	Neutralizing Monoclonal Antibodies against Hepatitis C Virus E2 Protein Bind Discontinuous Epitopes and Inhibit Infection at a Postattachment Step. <i>Journal of Virology</i> , 2011, 85, 7005-7019.	3.4	120
13	Identification of a Broadly Cross-Reacting and Neutralizing Human Monoclonal Antibody Directed against the Hepatitis C Virus E2 Protein. <i>Journal of Virology</i> , 2008, 82, 1047-1052.	3.4	119
14	Directex vivo comparison of the breadth and specificity of the T cells in the liver and peripheral blood of patients with chronic HCV infection. <i>European Journal of Immunology</i> , 2001, 31, 2388-2394.	2.9	118
15	Definition of a Conserved Immunodominant Domain on Hepatitis C Virus E2 Glycoprotein by Neutralizing Human Monoclonal Antibodies. <i>Journal of Virology</i> , 2008, 82, 6061-6066.	3.4	112
16	Development of a strand-specific RT-PCR based assay to detect the replicative form of hepatitis C virus RNA. <i>Journal of Virological Methods</i> , 2001, 94, 111-120.	2.1	98
17	Structural Flexibility of a Conserved Antigenic Region in Hepatitis C Virus Glycoprotein E2 Recognized by Broadly Neutralizing Antibodies. <i>Journal of Virology</i> , 2015, 89, 2170-2181.	3.4	96
18	Non-Macrophage-Tropic Human Immunodeficiency Virus Type 1 R5 Envelopes Predominate in Blood, Lymph Nodes, and Semen: Implications for Transmission and Pathogenesis. <i>Journal of Virology</i> , 2006, 80, 6324-6332.	3.4	95

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19	Variation in HIV-1 R5 macrophage-tropism correlates with sensitivity to reagents that block envelope: CD4 interactions but not with sensitivity to other entry inhibitors. <i>Retrovirology</i> , 2008, 5, 5.	2.0	71
20	An alpaca nanobody inhibits hepatitis C virus entry and cell-to-cell transmission. <i>Hepatology</i> , 2013, 58, 932-939.	7.3	69
21	Hepatitis C Virus (HCV) Infection May Elicit Neutralizing Antibodies Targeting Epitopes Conserved in All Viral Genotypes. <i>PLoS ONE</i> , 2009, 4, e8254.	2.5	64
22	Identification of New Functional Regions in Hepatitis C Virus Envelope Glycoprotein E2. <i>Journal of Virology</i> , 2011, 85, 1777-1792.	3.4	64
23	A Diverse Panel of Hepatitis C Virus Glycoproteins for Use in Vaccine Research Reveals Extremes of Monoclonal Antibody Neutralization Resistance. <i>Journal of Virology</i> , 2016, 90, 3288-3301.	3.4	62
24	Determination of the human antibody response to the epitope defined by the hepatitis C virus-neutralizing monoclonal antibody AP33. <i>Journal of General Virology</i> , 2007, 88, 2991-3001.	2.9	61
25	The role of neutralizing antibodies in hepatitis C virus infection. <i>Journal of General Virology</i> , 2012, 93, 1-19.	2.9	58
26	Naturally Occurring Antibodies That Recognize Linear Epitopes in the Amino Terminus of the Hepatitis C Virus E2 Protein Confer Noninterfering, Additive Neutralization. <i>Journal of Virology</i> , 2012, 86, 2739-2749.	3.4	54
27	Hepatitis C Virus Vaccine: Challenges and Prospects. <i>Vaccines</i> , 2020, 8, 90.	4.4	53
28	Studentsâ€™ Views towards Sars-Cov-2 Mass Asymptomatic Testing, Social Distancing and Self-Isolation in a University Setting during the COVID-19 Pandemic: A Qualitative Study. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 4182.	2.6	53
29	Specific interaction of hepatitis C virus glycoproteins with mannan binding lectin inhibits virus entry. <i>Protein and Cell</i> , 2010, 1, 664-674.	11.0	52
30	Hepatitis C Patient-Derived Glycoproteins Exhibit Marked Differences in Susceptibility to Serum Neutralizing Antibodies: Genetic Subtype Defines Antigenic but Not Neutralization Serotype. <i>Journal of Virology</i> , 2011, 85, 4246-4257.	3.4	51
31	Antigenicity and Immunogenicity of Differentially Glycosylated Hepatitis C Virus E2 Envelope Proteins Expressed in Mammalian and Insect Cells. <i>Journal of Virology</i> , 2019, 93, .	3.4	51
32	Intercompartmental Recombination of HIV-1 Contributes to <i>env</i> Intrahost Diversity and Modulates Viral Tropism and Sensitivity to Entry Inhibitors. <i>Journal of Virology</i> , 2011, 85, 6024-6037.	3.4	50
33	Evolutionary trends of the first hypervariable region of the hepatitis C virus E2 protein in individuals with differing liver disease severity. <i>Journal of General Virology</i> , 2002, 83, 11-23.	2.9	49
34	A Conserved Determinant in the V1 Loop of HIV-1 Modulates the V3 Loop To Prime Low CD4 Use and Macrophage Infection. <i>Journal of Virology</i> , 2011, 85, 2397-2405.	3.4	49
35	All Surfaces Are Not Equal in Contact Transmission of SARS-CoV-2. <i>Matter</i> , 2020, 3, 1433-1441.	10.0	49
36	Discovery of Novel Alphacoronaviruses in European Rodents and Shrews. <i>Viruses</i> , 2016, 8, 84.	3.3	45

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37	The Role of Humoral Innate Immunity in Hepatitis C Virus Infection. <i>Viruses</i> , 2012, 4, 1-27.	3.3	43
38	Immunogenicity of a new gorilla adenovirus vaccine candidate for COVID-19. <i>Molecular Therapy</i> , 2021, 29, 2412-2423.	8.2	41
39	Two doses of the SARS-CoV-2 BNT162b2 vaccine enhance antibody responses to variants in individuals with prior SARS-CoV-2 infection. <i>Science Translational Medicine</i> , 2021, 13, eabj0847.	12.4	40
40	Severe fibrosis in hepatitis C virus-infected patients is associated with increased activity of the mannan-binding lectin (MBL)/MBL-associated serine protease 1 (MASP-1) complex. <i>Clinical and Experimental Immunology</i> , 2006, 147, 061127015327009-???	2.6	38
41	B-cell receptors expressed by lymphomas of hepatitis C virus (HCV)â€“infected patients rarely react with the viral proteins. <i>Blood</i> , 2014, 123, 1512-1515.	1.4	37
42	Evolutionary dynamics of hepatitis C virus envelope genes during chronic infection. <i>Journal of General Virology</i> , 2005, 86, 1931-1942.	2.9	36
43	An ancestral host defence peptide within human Î²-defensin 3 recapitulates the antibacterial and antiviral activity of the full-length molecule. <i>Scientific Reports</i> , 2016, 5, 18450.	3.3	35
44	Shared Common Ancestry of Rodent Alphacoronaviruses Sampled Globally. <i>Viruses</i> , 2019, 11, 125.	3.3	35
45	Novel functional hepatitis C virus glycoprotein isolates identified using an optimized viral pseudotype entry assay. <i>Journal of General Virology</i> , 2016, 97, 2265-2279.	2.9	33
46	Hepatitis C Virus Envelope Glycoprotein Fitness Defines Virus Population Composition following Transmission to a New Host. <i>Journal of Virology</i> , 2012, 86, 11956-11966.	3.4	31
47	Dendritic cells cultured from mononuclear cells and CD34 cells in myeloma do not harbour human herpesvirus 8. <i>British Journal of Haematology</i> , 1998, 100, 793-796.	2.5	28
48	Analysis of the binding of hepatitis C virus genotype 1a and 1b E2 glycoproteins to peripheral blood mononuclear cell subsets. <i>Journal of General Virology</i> , 2005, 86, 2507-2512.	2.9	28
49	Recombinant Human L-Ficolin Directly Neutralizes Hepatitis C Virus Entry. <i>Journal of Innate Immunity</i> , 2014, 6, 676-684.	3.8	28
50	HIV-1 in semen: Determination of proviral and viral titres compared to blood, and quantification of semen leukocyte populations. , 1999, 59, 356-363.		27
51	Cross-genotype AR3-specific neutralizing antibodies confer long-term protection in injecting drug users after HCV clearance. <i>Journal of Hepatology</i> , 2019, 71, 14-24.	3.7	27
52	Cross-genotype characterization of genetic diversity and molecular adaptation in hepatitis C virus envelope glycoprotein genes. <i>Journal of General Virology</i> , 2007, 88, 458-469.	2.9	25
53	Cloning, Expression, and Functional Analysis of Patient-Derived Hepatitis C Virus Glycoproteins. <i>Methods in Molecular Biology</i> , 2007, 379, 177-197.	0.9	25
54	The Impact of Real-Time Whole-Genome Sequencing in Controlling Healthcare-Associated SARS-CoV-2 Outbreaks. <i>Journal of Infectious Diseases</i> , 2022, 225, 10-18.	4.0	25

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55	Concordance between semen-derived HIV-1 proviral DNA and viral RNA hypervariable region 3 (V3) envelope sequences in cases where semen populations are distinct from those present in blood. <i>Journal of Medical Virology</i> , 2002, 67, 9-19.	5.0	24
56	Variation in the biological properties of HIV-1 R5 envelopes: implications of envelope structure, transmission and pathogenesis. <i>Future Virology</i> , 2010, 5, 435-451.	1.8	24
57	Long-lasting viability of HIV after patient's death. <i>Lancet, The</i> , 1991, 338, 63.	13.7	22
58	Targeting a host-cell entry factor barricades antiviral-resistant HCV variants from on-therapy breakthrough in human-liver mice. <i>Cut</i> , 2016, 65, 2029-2034.	12.1	21
59	Interferon-Induced Transmembrane Proteins Mediate Viral Evasion in Acute and Chronic Hepatitis C Virus Infection. <i>Hepatology</i> , 2019, 70, 1506-1520.	7.3	21
60	Adjuvant formulated virus-like particles expressing native-like forms of the Lassa virus envelope surface glycoprotein are immunogenic and induce antibodies with broadly neutralizing activity. <i>Npj Vaccines</i> , 2020, 5, 71.	6.0	21
61	Semen characteristics in HIV-1 positive men and the effect of semen washing.. <i>Sexually Transmitted Infections</i> , 1997, 73, 303-305.	1.9	20
62	An Antigenically Diverse, Representative Panel of Envelope Glycoproteins for Hepatitis C Virus Vaccine Development. <i>Gastroenterology</i> , 2022, 162, 562-574.	1.3	20
63	HIV coreceptor and chemokine ligand gene expression in the male urethra and female cervix. <i>Aids</i> , 2005, 19, 1257-1265.	2.2	19
64	A novel neutralizing human monoclonal antibody broadly abrogates hepatitis C virus infection in vitro and in vivo. <i>Antiviral Research</i> , 2017, 148, 53-64.	4.1	18
65	A next generation vaccine against human rabies based on a single dose of a chimpanzee adenovirus vector serotype C. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008459.	3.0	18
66	Dramatic Potentiation of the Antiviral Activity of HIV Antibodies by Cholesterol Conjugation. <i>Journal of Biological Chemistry</i> , 2014, 289, 35015-35028.	3.4	17
67	Structure-Based Design of Hepatitis C Virus E2 Glycoprotein Improves Serum Binding and Cross-Neutralization. <i>Journal of Virology</i> , 2020, 94, .	3.4	17
68	Standardized Method for the Study of Antibody Neutralization of HCV Pseudoparticles (HCVpp). <i>Methods in Molecular Biology</i> , 2019, 1911, 441-450.	0.9	17
69	Novel human anti-claudin 1 mAbs inhibit hepatitis C virus infection and may synergize with anti-SRB1 mAb. <i>Journal of General Virology</i> , 2016, 97, 82-94.	2.9	16
70	Detection of HIV-1 by digoxigenin-labelled PCR and microtitre plate solution hybridisation assay and prevention of PCR carry-over by uracil-N-glycosylase. <i>Journal of Virological Methods</i> , 1993, 44, 67-76.	2.1	14
71	Incorporation of single-stranded DNA binding protein early in polymerase chain reaction product sequencing reactions prevents enzyme pausing. <i>Analytical Biochemistry</i> , 1992, 207, 349-351.	2.4	13
72	Characterisation of a series of human immunodeficiency virus isolates derived sequentially from a single patient. <i>Journal of Medical Virology</i> , 1991, 34, 104-113.	5.0	12

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73	Development of a high-throughput pyrosequencing assay for monitoring temporal evolution and resistance associated variant emergence in the Hepatitis C virus protease coding-region. <i>Antiviral Research</i> , 2014, 110, 52-59.	4.1	12
74	A modified alkaline phosphatase enzyme amplification system and its application in an HIV antigen ELISA. <i>Journal of Virological Methods</i> , 1992, 37, 149-153.	2.1	11
75	GBV-C/HCV coinfection in HIV-1-positive men: Frequent detection of viral RNA in blood plasma but absence from seminal fluid plasma. , 1998, 56, 321-326.		11
76	A polymerase chain reaction method for the amplification of full-length envelope genes of HIV-1 from DNA samples containing single molecules of HIV-1 provirus. <i>Journal of Virological Methods</i> , 2000, 88, 73-80.	2.1	11
77	Flexible and rapid construction of viral chimeras applied to hepatitis C virus. <i>Journal of General Virology</i> , 2016, 97, 2187-2193.	2.9	11
78	Retrospective screening of routine respiratory samples revealed undetected community transmission and missed intervention opportunities for SARS-CoV-2 in the United Kingdom. <i>Journal of General Virology</i> , 2021, 102, .	2.9	10
79	Elevated serum activity of MBL and ficolin-2 as biomarkers for progression to hepatocellular carcinoma in chronic HCV infection. <i>Virology</i> , 2019, 530, 99-106.	2.4	9
80	Discovery and Prevalence of Divergent RNA Viruses in European Field Voles and Rabbits. <i>Viruses</i> , 2020, 12, 47.	3.3	9
81	Production of Single-Stranded DNA Using a Uracil-N-glycosylase-Mediated Asymmetric Polymerase Chain Reaction Method. <i>Analytical Biochemistry</i> , 1997, 253, 264-267.	2.4	8
82	Poor reduction of HIV-1 RNA titres in nucleoside reverse transcriptase inhibitor experienced patients treated with indinavir combination therapy. <i>Sexually Transmitted Infections</i> , 1999, 75, 337-339.	1.9	8
83	Non-ionic detergents facilitate non-specific binding of M13 bacteriophage to polystyrene surfaces. <i>Journal of Virological Methods</i> , 2015, 221, 1-8.	2.1	8
84	Broad neutralization of hepatitis C virus-resistant variants by Civacir hepatitis C immunoglobulin. <i>Hepatology</i> , 2016, 64, 1495-1506.	7.3	8
85	Development and characterization of a human monoclonal antibody targeting the N-terminal region of hepatitis C virus envelope glycoprotein E1. <i>Virology</i> , 2018, 514, 30-41.	2.4	8
86	Immunization with a synthetic consensus hepatitis C virus E2 glycoprotein ectodomain elicits virus-neutralizing antibodies. <i>Antiviral Research</i> , 2018, 160, 25-37.	4.1	8
87	Real-World Outcomes of Direct-Acting Antiviral Treatment and Retreatment in United Kingdom-Based Patients Infected With Hepatitis C Virus Genotypes/Subtypes Endemic in Africa. <i>Journal of Infectious Diseases</i> , 2022, 226, 995-1004.	4.0	8
88	The use of uracil-N-glycosylase in the preparation of PCR products of direct sequencing. <i>Nucleic Acids Research</i> , 1992, 20, 3255-3255.	14.5	7
89	A bivalent HCV peptide vaccine elicits pan-genotypic neutralizing antibodies in mice. <i>Vaccine</i> , 2020, 38, 6864-6867.	3.8	7
90	Trichodysplasia Spinulosa Polyomavirus in Respiratory Tract of Immunocompromised Child. <i>Emerging Infectious Diseases</i> , 2018, 24, 1744-1746.	4.3	6

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91	OUP accepted manuscript. Journal of Infectious Diseases, 2022, , .	4.0	6
92	Association of antibodies to hepatitis C virus glycoproteins 1 and 2 (anti-E1E2) with HCV disease. Journal of Viral Hepatitis, 2008, 15, 339-345.	2.0	5
93	A case of hepatitis C virus transmission acquired through sharing a haemodialysis machine. CKJ: Clinical Kidney Journal, 2011, 4, 32-35.	2.9	5
94	HIV-1 co-receptor expression and epithelial immune cells of the cervix in asymptomatic women attending a genitourinary medicine clinic. HIV Medicine, 2013, 14, 108-114.	2.2	5
95	Cholesterol conjugation potentiates the antiviral activity of an HIV immunoadhesin. Journal of Peptide Science, 2015, 21, 743-749.	1.4	5
96	Polymer microarrays rapidly identify competitive adsorbents of virus-like particles. Biointerphases, 2020, 15, 061005.	1.6	5
97	Challenges on the development of a pseudotyping assay for Zika glycoproteins. Journal of Medical Microbiology, 2021, 70, .	1.8	5
98	Retrieval of the Complete Coding Sequence of the UK-Endemic Tatenale Orthohantavirus Reveals Extensive Strain Variation and Supports Its Classification as a Novel Species. Viruses, 2020, 12, 454.	3.3	4
99	Expression of human ficolin-2 in hepatocytes confers resistance to infection by diverse hepatotropic viruses. Journal of Medical Microbiology, 2019, 68, 642-648.	1.8	4
100	SARS-CoV-2 proteins (version 2020.2) in the IUPHAR/BPS Guide to Pharmacology Database. IUPHAR/BPS Guide To Pharmacology CITE, 2020, 2020, .	0.2	4
101	Discovery of Novel Coronaviruses in Rodents. Methods in Molecular Biology, 2020, 2203, 33-40.	0.9	4
102	Hepatitis C virus quasispecies and pseudotype analysis from acute infection to chronicity in HIV-1 co-infected individuals. Virology, 2016, 492, 213-224.	2.4	3
103	Identification of Infectious Agents in High-Throughput Sequencing Data Sets Is Easily Achievable Using Free, Cloud-Based Bioinformatics Platforms. Journal of Clinical Microbiology, 2019, 57, .	3.9	3
104	Role of HVR1 sequence similarity in the cross-genotypic neutralization of HCV. Virology Journal, 2020, 17, 140.	3.4	3
105	Cloning and Analysis of Authentic Patient-Derived HCV E1/E2 Glycoproteins. Methods in Molecular Biology, 2019, 1911, 275-294.	0.9	3
106	Simultaneous determination of HCV genotype and NS5B resistance associated substitutions using dried serum spots from São Paulo state, Brazil. Access Microbiology, 2022, 4, .	0.5	3
107	Human parainfluenza 2 & 4: Clinical and genetic epidemiology in the UK, 2013–2017, reveals distinct disease features and co-circulating genomic subtypes. Influenza and Other Respiratory Viruses, 2022, 16, 1122-1132.	3.4	3
108	Analysis of Serine Codon Conservation Reveals Diverse Phenotypic Constraints on Hepatitis C Virus Glycoprotein Evolution. Journal of Virology, 2014, 88, 667-678.	3.4	2

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109	Use of short tandem repeat fingerprinting to validate sample origins in hepatitis C virus molecular epidemiology studies. Journal of General Virology, 2014, 95, 66-70.	2.9	2
110	Enterovirus D68 epidemic, UK, 2018, was caused by subclades B3 and D1, predominantly in children and adults, respectively, with both subclades exhibiting extensive genetic diversity. Microbial Genomics, 2022, 8, .	2.0	2
111	Tagged polymerase chain reaction subtractive hybridization for the enrichment of phage display random peptide libraries. Analytical Biochemistry, 2005, 339, 61-68.	2.4	1
112	InFusion Cloning for the Generation of Biologically Relevant HCV Chimeric Molecular Clones. Methods in Molecular Biology, 2019, 1911, 93-104.	0.9	1
113	Coronavirus (CoV) proteins (version 2020.4) in the IUPHAR/BPS Guide to Pharmacology Database. IUPHAR/BPS Guide To Pharmacology CITE, 2020, 2020, .	0.2	1
114	The HCV Envelope Glycoprotein Down-Modulates NF- κ B Signalling and Associates With Stimulation of the Host Endoplasmic Reticulum Stress Pathway. Frontiers in Immunology, 2022, 13, 831695.	4.8	1
115	Coronavirus (CoV) proteins in GtoPdb v.2021.3. IUPHAR/BPS Guide To Pharmacology CITE, 2021, 2021, .	0.2	0
116	Discovery of novel highly divergent RNA viruses in European rodents and rabbits. Access Microbiology, 2019, 1, .	0.5	0
117	Coronavirus (CoV) proteins in GtoPdb v.2022.2. IUPHAR/BPS Guide To Pharmacology CITE, 2022, 2022, .	0.2	0