

# Stuart C Ray

## List of Publications by Year in descending order

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136  
papers

14,429  
citations

30047

54  
h-index

19726

117  
g-index

144  
all docs

144  
docs citations

144  
times ranked

12876  
citing authors

#	ARTICLE	IF	CITATIONS
1	Full-Length Human Immunodeficiency Virus Type 1 Genomes from Subtype C-Infected Seroconverters in India, with Evidence of Intersubtype Recombination. <i>Journal of Virology</i> , 1999, 73, 152-160.	1.5	2,383
2	Determinants of Viral Clearance and Persistence during Acute Hepatitis C Virus Infection. <i>Journal of Experimental Medicine</i> , 2001, 194, 1395-1406.	4.2	1,091
3	Immune evasion by hepatitis C virus NS3/4A protease-mediated cleavage of the Toll-like receptor 3 adaptor protein TRIF. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 2992-2997.	3.3	991
4	Eukaryotic start and stop translation sites. <i>Nucleic Acids Research</i> , 1991, 19, 3185-3192.	6.5	631
5	Protection against persistence of hepatitis C. <i>Lancet, The</i> , 2002, 359, 1478-1483.	6.3	426
6	Residual Human Immunodeficiency Virus Type 1 Viremia in Some Patients on Antiretroviral Therapy Is Dominated by a Small Number of Invariant Clones Rarely Found in Circulating CD4 + T Cells. <i>Journal of Virology</i> , 2006, 80, 6441-6457.	1.5	377
7	Genetic Epidemiology of Hepatitis C Virus throughout Egypt. <i>Journal of Infectious Diseases</i> , 2000, 182, 698-707.	1.9	336
8	Intermittent HIV-1 Viremia (Blips) and Drug Resistance in Patients Receiving HAART. <i>JAMA - Journal of the American Medical Association</i> , 2005, 293, 817.	3.8	323
9	Spontaneous Control of Primary Hepatitis C Virus Infection and Immunity Against Persistent Reinfection. <i>Gastroenterology</i> , 2010, 138, 315-324.	0.6	316
10	Cellular immune selection with hepatitis C virus persistence in humans. <i>Journal of Experimental Medicine</i> , 2005, 201, 1741-1752.	4.2	278
11	Oral antibiotic treatment of right-sided staphylococcal endocarditis in injection drug users: Prospective randomized comparison with parenteral therapy. <i>American Journal of Medicine</i> , 1996, 101, 68-76.	0.6	256
12	Human Immunodeficiency Virus-Related Microbial Translocation and Progression of Hepatitis C. <i>Gastroenterology</i> , 2008, 135, 226-233.	0.6	251
13	Acute hepatitis C. <i>Lancet, The</i> , 2008, 372, 321-332.	6.3	244
14	Clearance of hepatitis C infection is associated with the early appearance of broad neutralizing antibody responses. <i>Hepatology</i> , 2014, 59, 2140-2151.	3.6	230
15	Comprehensive analyses of CD8+ T cell responses during longitudinal study of acute human hepatitis C. <i>Hepatology</i> , 2005, 42, 104-112.	3.6	211
16	Selection Pressure From Neutralizing Antibodies Drives Sequence Evolution During Acute Infection With Hepatitis C Virus. <i>Gastroenterology</i> , 2009, 136, 2377-2386.	0.6	207
17	HIV-1 Drug Resistance Profiles in Children and Adults With Viral Load of <math>\leq 50</math> Copies/mL Receiving Combination Therapy. <i>JAMA - Journal of the American Medical Association</i> , 2001, 286, 196.	3.8	196
18	Prospective Evaluation of Community-Acquired Acute-Phase Hepatitis C Virus Infection. <i>Clinical Infectious Diseases</i> , 2005, 40, 951-958.	2.9	195

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19	Genotypic Analysis of HIV-1 Drug Resistance at the Limit of Detection: Virus Production without Evolution in Treated Adults with Undetectable HIV Loads. <i>Journal of Infectious Diseases</i> , 2004, 189, 1452-1465.	1.9	186
20	Divergent and convergent evolution after a common-source outbreak of hepatitis C virus. <i>Journal of Experimental Medicine</i> , 2005, 201, 1753-1759.	4.2	178
21	Cross-Genotype Immunity to Hepatitis C Virus. <i>Journal of Virology</i> , 2004, 78, 1575-1581.	1.5	175
22	Acute Hepatitis C Virus Structural Gene Sequences as Predictors of Persistent Viremia: Hypervariable Region 1 as a Decoy. <i>Journal of Virology</i> , 1999, 73, 2938-2946.	1.5	175
23	Expanded cellular clones carrying replication-competent HIV-1 persist, wax, and wane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E2575-E2584.	3.3	173
24	Humoral Immune Response in Acute Hepatitis C Virus Infection. <i>Clinical Infectious Diseases</i> , 2005, 41, 667-675.	2.9	172
25	Neutralizing Antibodies Do Not Mediate Suppression of Human Immunodeficiency Virus Type 1 in Elite Suppressors or Selection of Plasma Virus Variants in Patients on Highly Active Antiretroviral Therapy. <i>Journal of Virology</i> , 2006, 80, 4758-4770.	1.5	156
26	GATTA Hypermutation in Protease and Reverse Transcriptase Regions of Human Immunodeficiency Virus Type 1 Residing in Resting CD4+ T Cells In Vivo. <i>Journal of Virology</i> , 2005, 79, 1975-1980.	1.5	154
27	A stable latent reservoir for HIV-1 in resting CD4+ T lymphocytes in infected children. <i>Journal of Clinical Investigation</i> , 2000, 105, 995-1003.	3.9	151
28	Broadly neutralizing antibodies with few somatic mutations and hepatitis C virus clearance. <i>JCI Insight</i> , 2017, 2, .	2.3	129
29	Prospective Characterization of Full-Length Hepatitis C Virus NS5A Quasispecies during Induction and Combination Antiviral Therapy. <i>Journal of Virology</i> , 2000, 74, 9028-9038.	1.5	121
30	Persistence of Wild-Type Virus and Lack of Temporal Structure in the Latent Reservoir for Human Immunodeficiency Virus Type 1 in Pediatric Patients with Extensive Antiretroviral Exposure. <i>Journal of Virology</i> , 2002, 76, 9481-9492.	1.5	119
31	Control of HIV-1 in Elite Suppressors despite Ongoing Replication and Evolution in Plasma Virus. <i>Journal of Virology</i> , 2010, 84, 7018-7028.	1.5	116
32	Determinants of the Quantity of Hepatitis C Virus RNA. <i>Journal of Infectious Diseases</i> , 2000, 181, 844-851.	1.9	114
33	High-Programmed Death-1 Levels on Hepatitis C Virus-Specific T Cells during Acute Infection Are Associated with Viral Persistence and Require Preservation of Cognate Antigen during Chronic Infection. <i>Journal of Immunology</i> , 2008, 181, 8215-8225.	0.4	114
34	The Role of Viral Introductions in Sustaining Community-Based HIV Epidemics in Rural Uganda: Evidence from Spatial Clustering, Phylogenetics, and Egocentric Transmission Models. <i>PLoS Medicine</i> , 2014, 11, e1001610.	3.9	114
35	Analysis of Genetic Linkage of HIV From Couples Enrolled in the HIV Prevention Trials Network 052 Trial. <i>Journal of Infectious Diseases</i> , 2011, 204, 1918-1926.	1.9	99
36	Needlestick Transmission of Hepatitis C. <i>JAMA - Journal of the American Medical Association</i> , 2002, 287, 2406.	3.8	98

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37	Continued Production of Drug-Sensitive Human Immunodeficiency Virus Type 1 in Children on Combination Antiretroviral Therapy Who Have Undetectable Viral Loads. <i>Journal of Virology</i> , 2004, 78, 968-979.	1.5	98
38	Transmission of Human Immunodeficiency Virus Type 1 from a Patient Who Developed AIDS to an Elite Suppressor. <i>Journal of Virology</i> , 2008, 82, 7395-7410.	1.5	90
39	Broadly Neutralizing Antibody Mediated Clearance of Human Hepatitis C Virus Infection. <i>Cell Host and Microbe</i> , 2018, 24, 717-730.e5.	5.1	78
40	Naturally selected hepatitis C virus polymorphisms confer broad neutralizing antibody resistance. <i>Journal of Clinical Investigation</i> , 2015, 125, 437-447.	3.9	78
41	Durable SARS-CoV-2 B cell immunity after mild or severe disease. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	76
42	Healthy donor T cell responses to common cold coronaviruses and SARS-CoV-2. <i>Journal of Clinical Investigation</i> , 2020, 130, 6631-6638.	3.9	75
43	Use of Laser Capture Microdissection to Map Hepatitis C Virus-Positive Hepatocytes in Human Liver. <i>Gastroenterology</i> , 2013, 145, 1404-1413.e10.	0.6	74
44	Acceleration of Hepatitis C Virus Envelope Evolution in Humans Is Consistent with Progressive Humoral Immune Selection during the Transition from Acute to Chronic Infection. <i>Journal of Virology</i> , 2010, 84, 5067-5077.	1.5	70
45	Comprehensive Genetic and Epigenetic Analysis of Occult Hepatitis B from Liver Tissue Samples. <i>Clinical Infectious Diseases</i> , 2008, 46, 1227-1236.	2.9	69
46	Genome Sequencing and Analysis of Geographically Diverse Clinical Isolates of Herpes Simplex Virus 2. <i>Journal of Virology</i> , 2015, 89, 8219-8232.	1.5	68
47	Convergent evolution within the V3 loop domain of human immunodeficiency virus type 1 in association with disease progression. <i>Journal of Virology</i> , 1995, 69, 7548-7558.	1.5	68
48	Hypervariable Region 1 Sequence Stability during Hepatitis C Virus Replication in Chimpanzees. <i>Journal of Virology</i> , 2000, 74, 3058-3066.	1.5	65
49	CMPK2 and BCL-G are associated with type 1 interferon-induced HIV restriction in humans. <i>Science Advances</i> , 2018, 4, eaat0843.	4.7	64
50	Spontaneous clearance of primary acute hepatitis C virus infection correlated with high initial viral RNA level and rapid HVR1 evolution. <i>Hepatology</i> , 2012, 55, 1684-1691.	3.6	63
51	Hepatitis C Virus Immune Escape via Exploitation of a Hole in the T Cell Repertoire. <i>Journal of Immunology</i> , 2008, 181, 6435-6446.	0.4	61
52	Slow Human Immunodeficiency Virus Type 1 Evolution in Viral Reservoirs in Infants Treated with Effective Antiretroviral Therapy. <i>AIDS Research and Human Retroviruses</i> , 2007, 23, 381-390.	0.5	59
53	Hepatitis C Virus Infection of Neuroepithelioma Cell Lines. <i>Gastroenterology</i> , 2010, 139, 1365-1374.e2.	0.6	59
54	Early Archiving and Predominance of Nonnucleoside Reverse Transcriptase Inhibitor-Resistant HIV-1 among Recently Infected Infants Born in the United States. <i>Journal of Infectious Diseases</i> , 2007, 195, 1402-1410.	1.9	58

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55	PCR Detection of Adenovirus in a Bone Marrow Transplant Recipient: Hemorrhagic Cystitis as a Presenting Manifestation of Disseminated Disease. <i>Journal of Clinical Microbiology</i> , 1999, 37, 686-689.	1.8	57
56	Kupffer cells are depleted with HIV immunodeficiency and partially recovered with antiretroviral immune reconstitution. <i>Aids</i> , 2009, 23, 2397-2404.	1.0	55
57	Synergistic anti-HCV broadly neutralizing human monoclonal antibodies with independent mechanisms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E82-E91.	3.3	52
58	Prevalence of Congenital Homocystinuria in Denmark. <i>New England Journal of Medicine</i> , 1999, 340, 1513-1513.	13.9	50
59	Extra-epitopic hepatitis C virus polymorphisms confer resistance to broadly neutralizing antibodies by modulating binding to scavenger receptor B1. <i>PLoS Pathogens</i> , 2017, 13, e1006235.	2.1	47
60	Presence in India of HIV Type 1 Similar to North American Strains. <i>AIDS Research and Human Retroviruses</i> , 1994, 10, 1039-1041.	0.5	45
61	Inferring Viral Dynamics in Chronically HCV Infected Patients from the Spatial Distribution of Infected Hepatocytes. <i>PLoS Computational Biology</i> , 2014, 10, e1003934.	1.5	45
62	Human Immunodeficiency Virus Seroconversion and Evolution of the Hepatitis C Virus Quasispecies. <i>Journal of Virology</i> , 2001, 75, 3259-3267.	1.5	42
63	Identification of Ongoing Human Immunodeficiency Virus Type 1 (HIV-1) Replication in Residual Viremia during Recombinant HIV-1 Poxvirus Immunizations in Patients with Clinically Undetectable Viral Loads on Durable Suppressive Highly Active Antiretroviral Therapy. <i>Journal of Virology</i> , 2009, 83, 9731-9742.	1.5	41
64	Evolution of the HIV-1 nefgene in HLA-B*57 Positive Elite Suppressors. <i>Retrovirology</i> , 2010, 7, 94.	0.9	39
65	A sensitive genotyping assay for detection of drug resistance mutations in reverse transcriptase of HIV-1 subtypes B and C in samples stored as dried blood spots or frozen RNA extracts. <i>Journal of Virological Methods</i> , 2006, 136, 238-247.	1.0	37
66	Global Diversity within and between Human Herpesvirus 1 and 2 Glycoproteins. <i>Journal of Virology</i> , 2015, 89, 8206-8218.	1.5	37
67	Assessment of Hepatitis C Virus Sequence Complexity by Electrophoretic Mobilities of Both Single-and Double-Stranded DNAs. <i>Journal of Clinical Microbiology</i> , 1998, 36, 2982-2989.	1.8	36
68	Dynamics of SEN Virus Infection among Injection Drug Users. <i>Journal of Infectious Diseases</i> , 2001, 184, 1315-1319.	1.9	35
69	Identification of Nevirapine-Resistant HIV-1 in the Latent Reservoir after Single-Dose Nevirapine to Prevent Mother-to-Child Transmission of HIV-1. <i>Journal of Infectious Diseases</i> , 2009, 199, 1301-1309.	1.9	35
70	High diversity of hepatitis C viral quasispecies is associated with early virological response in patients undergoing antiviral therapy. <i>Hepatology</i> , 2009, 50, 1765-1772.	3.6	35
71	Plasma deconvolution identifies broadly neutralizing antibodies associated with hepatitis C virus clearance. <i>Journal of Clinical Investigation</i> , 2019, 129, 4786-4796.	3.9	33
72	A Novel Assay Allows Genotyping of the Latent Reservoir for Human Immunodeficiency Virus Type 1 in the Resting CD4+ T Cells of Viremic Patients. <i>Journal of Virology</i> , 2005, 79, 5185-5202.	1.5	32

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73	Genomic diversity of SARS-CoV-2 during early introduction into the Baltimore–Washington metropolitan area. <i>JCI Insight</i> , 2021, 6, .	2.3	31
74	Progression of Fibrosis during Chronic Hepatitis C Is Associated with Rapid Virus Evolution. <i>Journal of Virology</i> , 2007, 81, 6513-6522.	1.5	30
75	Sex and the genetic diversity of HIV-1. <i>Nature Medicine</i> , 2000, 6, 23-25.	15.2	29
76	CD4+T Cell Depletion in an Untreated HIV Type 1–Infected Human Leukocyte Antigen–B*5801–Positive Patient with an Undetectable Viral Load. <i>Clinical Infectious Diseases</i> , 2008, 46, e78-e82.	2.9	29
77	Hepatitis C virus epitope exposure and neutralization by antibodies is affected by time and temperature. <i>Virology</i> , 2012, 422, 174-184.	1.1	29
78	Immunogenicity and Cross-Reactivity of a Representative Ancestral Sequence in Hepatitis C Virus Infection. <i>Journal of Immunology</i> , 2012, 188, 5177-5188.	0.4	28
79	No evidence of SARS-CoV-2 reverse transcription and integration as the origin of chimeric transcripts in patient tissues. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	28
80	Hepatitis C virus resistance to broadly neutralizing antibodies measured using replication-competent virus and pseudoparticles. <i>Journal of General Virology</i> , 2016, 97, 2883-2893.	1.3	27
81	Molecular epidemiology of HIV-1 subtypes in southern China. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2005, 38, 356-62.	0.9	26
82	Accurate Representation of the Hepatitis C Virus Quasispecies in 5.2-Kilobase Amplicons. <i>Journal of Clinical Microbiology</i> , 2004, 42, 4223-4229.	1.8	25
83	Maternal Neutralizing Antibody and Transmission of Hepatitis C Virus to Infants. <i>Journal of Infectious Diseases</i> , 2008, 198, 1651-1655.	1.9	25
84	Analysis of HIV Diversity Using a High-Resolution Melting Assay. <i>AIDS Research and Human Retroviruses</i> , 2010, 26, 913-918.	0.5	24
85	A Hepatitis C Virus Envelope Polymorphism Confers Resistance to Neutralization by Polyclonal Sera and Broadly Neutralizing Monoclonal Antibodies. <i>Journal of Virology</i> , 2016, 90, 3773-3782.	1.5	24
86	An Update on Severe Acute Respiratory Syndrome Coronavirus 2 Diversity in the US National Capital Region: Evolution of Novel and Variants of Concern. <i>Clinical Infectious Diseases</i> , 2022, 74, 1419-1428.	2.9	24
87	Constraints on Viral Evolution during Chronic Hepatitis C Virus Infection Arising from a Common-Source Exposure. <i>Journal of Virology</i> , 2012, 86, 12582-12590.	1.5	23
88	IgM anti-ACE2 autoantibodies in severe COVID-19 activate complement and perturb vascular endothelial function. <i>JCI Insight</i> , 2022, 7, .	2.3	23
89	Analyses of HIV-1 Drug-Resistance Profiles Among Infected Adolescents Experiencing Delayed Antiretroviral Treatment Switch After Initial Nonsuppressive Highly Active Antiretroviral Therapy. <i>AIDS Patient Care and STDs</i> , 2008, 22, 545-552.	1.1	22
90	Computational Reconstruction of Bole1a, a Representative Synthetic Hepatitis C Virus Subtype 1a Genome. <i>Journal of Virology</i> , 2012, 86, 5915-5921.	1.5	21

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91	Human Immunodeficiency Virus Type 1 env and p17gag Sequence Variation in Polymerase Chain Reaction-Positive, Seronegative Injection Drug Users. <i>Journal of Infectious Diseases</i> , 1995, 171, 797-804.	1.9	19
92	Acute Hepatitis C Virus Infection Induces Consistent Changes in Circulating MicroRNAs That Are Associated with Nonlytic Hepatocyte Release. <i>Journal of Virology</i> , 2015, 89, 9454-9464.	1.5	19
93	Rapid and sensitive detection of viral nucleic acids using silicon microchips. <i>Analyst, The</i> , 2018, 143, 2596-2603.	1.7	19
94	Predominance of defective proviral sequences in an HIV + long-term non-progressor. <i>Immunology Letters</i> , 1996, 51, 3-6.	1.1	18
95	Autologous Strain-Specific Cytolytic T Lymphocyte Responses Directed against Human Immunodeficiency Virus Type 1 Env. <i>AIDS Research and Human Retroviruses</i> , 1998, 14, 3-13.	0.5	18
96	CD4 <sup>+</sup> T-Cell-Dependent Reduction in Hepatitis C Virus-Specific Neutralizing Antibody Responses After Coinfection With Human Immunodeficiency Virus. <i>Journal of Infectious Diseases</i> , 2015, 212, 914-923.	1.9	18
97	Interferon-free treatment of chronic hepatitis C with faldaprevir, deleobuvir and ribavirin: <scp>SOUND</scp>â€ƒ3, a Phase 2b study. <i>Liver International</i> , 2015, 35, 417-421.	1.9	18
98	Accelerating Drug Development Through Collaboration: The Hepatitis C Drug Development Advisory Group. <i>Clinical Pharmacology and Therapeutics</i> , 2014, 96, 162-165.	2.3	17
99	Epidemiology of hepatitis C virus infection & liver disease among injection drug users (IDUs) in Chennai, India. <i>Indian Journal of Medical Research</i> , 2010, 132, 706-14.	0.4	17
100	The transhepatic endotoxin gradient is present despite liver cirrhosis and is attenuated after transjugular portosystemic shunt (TIPS).. <i>BMC Gastroenterology</i> , 2011, 11, 107.	0.8	15
101	Laser captured hepatocytes show association of butyrylcholinesterase gene loss and fibrosis progression in hepatitis C-infected drug users. <i>Hepatology</i> , 2012, 56, 544-554.	3.6	15
102	Molecular Epidemiology of HIV Type 1 in Singapore and Identification of Novel CRF01_AE/B Recombinant Forms. <i>AIDS Research and Human Retroviruses</i> , 2011, 27, 1135-1137.	0.5	14
103	Antiretroviral therapy, interferon sensitivity, and virologic setpoint in human immunodeficiency virus/hepatitis C virus coinfecting patients. <i>Hepatology</i> , 2014, 60, 477-486.	3.6	14
104	Characterization of novel recombinant HIV-1 genomes using the branching index. <i>Virology</i> , 2003, 316, 116-125.	1.1	13
105	Genetic Divergence of Hepatitis C Virus: The Role of HIV-Related Immunosuppression. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2008, 49, 136-141.	0.9	12
106	Complex patterns of Hepatitis-C virus longitudinal clustering in a high-risk population. <i>Infection, Genetics and Evolution</i> , 2018, 58, 77-82.	1.0	12
107	Monomer sequence determination of carbohydrates using fast-atom bombardment mass spectrometry of periodate-oxidized acetate ester derivatives. <i>Carbohydrate Research</i> , 1990, 197, 1-14.	1.1	11
108	Correlates of hepatitis C viral clustering among people who inject drugs in Baltimore. <i>Infection, Genetics and Evolution</i> , 2020, 77, 104078.	1.0	11



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109	Recrudescence of Treated Neurosyphilis in a Patient With Human Immunodeficiency Virus. <i>Mayo Clinic Proceedings</i> , 1999, 74, 53-56.	1.4	10
110	HIV Evolution Following Transmission to an HLA-B*5801-Positive Patient. <i>Journal of Infectious Diseases</i> , 2009, 200, 1820-1824.	1.9	9
111	Intracellular HIV-1 RNA and CD4+ T-cell activation in patients starting antiretrovirals. <i>Aids</i> , 2017, 31, 1405-1414.	1.0	9
112	Factors associated with phylogenetic clustering of hepatitis C among people who inject drugs in Baltimore. <i>BMC Infectious Diseases</i> , 2020, 20, 815.	1.3	9
113	Plasma virome and the risk of blood-borne infection in persons with substance use disorder. <i>Nature Communications</i> , 2021, 12, 6909.	5.8	8
114	Ecchymoses and eschars at sites of injection. <i>Lancet</i> , The, 1997, 349, 1364.	6.3	7
115	Generative optical modeling of whole blood for detecting platelets in lens-free images. <i>Biomedical Optics Express</i> , 2020, 11, 1808.	1.5	7
116	Drivers and barriers to workplace-based HIV self-testing among high-risk men in Uganda: a qualitative study. <i>BMC Public Health</i> , 2021, 21, 1002.	1.2	6
117	Spatiotemporal Phylodynamics of Hepatitis C Among People Who Inject Drugs in India. <i>Hepatology</i> , 2021, 74, 1782-1794.	3.6	6
118	B cell overexpression of FCRL5 and PD-1 is associated with low antibody titers in HCV infection. <i>PLoS Pathogens</i> , 2022, 18, e1010179.	2.1	6
119	Repeated exposure to heterologous hepatitis C viruses associates with enhanced neutralizing antibody breadth and potency. <i>Journal of Clinical Investigation</i> , 2022, 132, .	3.9	5
120	Interleukin-18 and tumor necrosis factor- $\alpha$ are elevated in solid organ transplant recipients with possible cytomegalovirus end-organ disease. <i>Transplant Infectious Disease</i> , 2021, 23, e13682.	0.7	4
121	Severe pruritus after completing pegylated interferon for hepatitis C. <i>Aids Reader</i> , 2008, 18, 562-5.	0.3	3
122	Inconsistent temporal patterns of genetic variation of HCV among high-risk subjects may impact inference of transmission networks. <i>Infection, Genetics and Evolution</i> , 2019, 71, 1-6.	1.0	2
123	Building Leadership Capacity for Mission Execution in a Large Academic Department of Medicine. <i>American Journal of Medicine</i> , 2019, 132, 535-543.	0.6	2
124	Decreased Activated CD4 <sup>+</sup> T Cell Repertoire Diversity After Antiretroviral Therapy in HIV-1/HCV Coinfection Correlates with CD4 <sup>+</sup> T Cell Recovery. <i>Viral Immunology</i> , 2021, 34, 622-631.	0.6	2
125	A 61-Year-Old Female with a Prior History of Tuberculosis Presenting with Hemoptysis. <i>Clinical Infectious Diseases</i> , 2011, 52, 910-910.	2.9	1
126	Molecular epidemiology of GB type C virus among individuals exposed to hepatitis C virus in Cameroon. <i>Mental Illness</i> , 2013, 4, 1.	0.8	1



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127	Impact of Point-of-Care Diagnostics on Clinical Decision-making in Low- and Middle-Income Countries. <i>Journal of applied laboratory medicine</i> , The, 2018, 3, 456-459.	0.6	1
128	An Optical Model of Whole Blood for Detecting Platelets in Lens-Free Images. <i>Lecture Notes in Computer Science</i> , 2019, , 140-150.	1.0	1
129	Severe acute respiratory coronavirus virus 2 (SARS-CoV-2) exposure investigations using genomic sequencing among healthcare workers and patients in a large academic center. <i>Infection Control and Hospital Epidemiology</i> , 2022, , 1-4.	1.0	1
130	Volatilization of mercury from natural water by a broad-spectrum Hg-resistant <i>Bacillus pasteurii</i> strain DR2. <i>The Environmentalist</i> , 1996, 16, 179-185.	0.7	0
131	Characterization of novel recombinant HIV-1 genomes using the branching index. <i>Virology</i> , 2003, 316, 116-116.	1.1	0
132	Genetic Linkage of Hepatitis B Virus in Peripheral Blood Leukocytes Provides Evidence for Contamination. <i>Journal of Virology</i> , 2010, 84, 2184-2186.	1.5	0
133	It's 10 pm ; Do You Know Where Your Data Are?. <i>Circulation Research</i> , 2017, 120, 1551-1554.	2.0	0
134	Joint Holographic Detection and Reconstruction. <i>Lecture Notes in Computer Science</i> , 2019, , 664-672.	1.0	0
135	The Brief Case: The Fly Who Cried Wohlf. <i>Journal of Clinical Microbiology</i> , 2022, 60, .	1.8	0
136	Closing the Brief Case: The Fly Who Cried Wohlf. <i>Journal of Clinical Microbiology</i> , 2022, 60, .	1.8	0