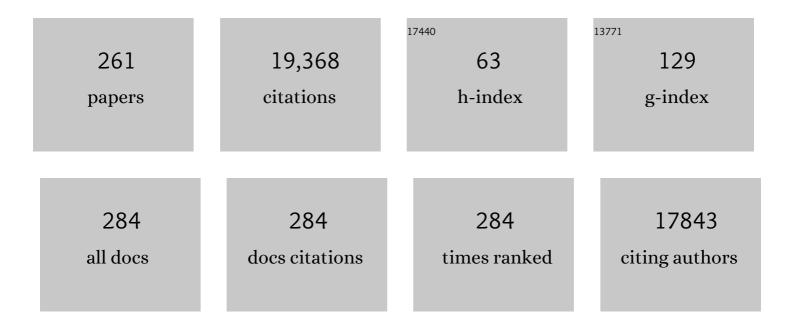
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification and characterization of transmitted and early founder virus envelopes in primary HIV-1 infection. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7552-7557.	7.1	1,708
2	Origin of HIV-1 in the chimpanzee Pan troglodytes troglodytes. Nature, 1999, 397, 436-441.	27.8	1,405
3	Human Immunodeficiency Virus Type 1 env Clones from Acute and Early Subtype B Infections for Standardized Assessments of Vaccine-Elicited Neutralizing Antibodies. Journal of Virology, 2005, 79, 10108-10125.	3.4	1,025
4	Co-evolution of a broadly neutralizing HIV-1 antibody and founder virus. Nature, 2013, 496, 469-476.	27.8	961
5	Diversity Considerations in HIV-1 Vaccine Selection. Science, 2002, 296, 2354-2360.	12.6	731
6	DEG 10, an update of the database of essential genes that includes both protein-coding genes and noncoding genomic elements: Table 1 Nucleic Acids Research, 2014, 42, D574-D580.	14.5	504
7	Human infection by genetically diverse SIVSM-related HIV-2 in West Africa. Nature, 1992, 358, 495-499.	27.8	486
8	Phenotypic properties of transmitted founder HIV-1. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 6626-6633.	7.1	379
9	Genetic and Neutralization Properties of Subtype C Human Immunodeficiency Virus Type 1 Molecular env Clones from Acute and Early Heterosexually Acquired Infections in Southern Africa. Journal of Virology, 2006, 80, 11776-11790.	3.4	334
10	Emergence of SARS-CoV-2 through recombination and strong purifying selection. Science Advances, 2020, 6, .	10.3	307
11	Maturation Pathway from Germline to Broad HIV-1 Neutralizer of a CD4-Mimic Antibody. Cell, 2016, 165, 449-463.	28.9	305
12	Ori-Finder: A web-based system for finding oriC s in unannotated bacterial genomes. BMC Bioinformatics, 2008, 9, 79.	2.6	287
13	A Comprehensive Panel of Near-Full-Length Clones and Reference Sequences for Non-Subtype B Isolates of Human Immunodeficiency Virus Type 1. Journal of Virology, 1998, 72, 5680-5698.	3.4	270
14	Cooperation of B Cell Lineages in Induction of HIV-1-Broadly Neutralizing Antibodies. Cell, 2014, 158, 481-491.	28.9	266
15	Global and regional molecular epidemiology of HIV-1, 1990–2015: a systematic review, global survey, and trend analysis. Lancet Infectious Diseases, The, 2019, 19, 143-155.	9.1	255
16	High-throughput isolation of immunoglobulin genes from single human B cells and expression as monoclonal antibodies. Journal of Virological Methods, 2009, 158, 171-179.	2.1	235
17	The Role of Antibody Polyspecificity and Lipid Reactivity in Binding of Broadly Neutralizing Anti-HIV-1 Envelope Human Monoclonal Antibodies 2F5 and 4E10 to Glycoprotein 41 Membrane Proximal Envelope Epitopes. Journal of Immunology, 2007, 178, 4424-4435.	0.8	230
18	Staged induction of HIV-1 glycan–dependent broadly neutralizing antibodies. Science Translational Medicine, 2017, 9, .	12.4	212

#	Article	IF	CITATIONS
19	Initial antibodies binding to HIV-1 gp41 in acutely infected subjects are polyreactive and highly mutated. Journal of Experimental Medicine, 2011, 208, 2237-2249.	8.5	198
20	Diversion of HIV-1 vaccine–induced immunity by gp41-microbiota cross-reactive antibodies. Science, 2015, 349, aab1253.	12.6	191
21	Antigenicity and Immunogenicity of a Synthetic Human Immunodeficiency Virus Type 1 Group M Consensus Envelope Glycoprotein. Journal of Virology, 2005, 79, 1154-1163.	3.4	189
22	Relative resistance of HIV-1 founder viruses to control by interferon-alpha. Retrovirology, 2013, 10, 146.	2.0	183
23	Near Full-Length Clones and Reference Sequences for Subtype C Isolates of HIV Type 1 from Three Different Continents. AIDS Research and Human Retroviruses, 2001, 17, 161-168.	1.1	182
24	A group M consensus envelope glycoprotein induces antibodies that neutralize subsets of subtype B and C HIV-1 primary viruses. Virology, 2006, 353, 268-282.	2.4	176
25	Bug mapping and fitness testing of chemically synthesized chromosome X. Science, 2017, 355, .	12.6	173
26	Polyclonal B Cell Responses to Conserved Neutralization Epitopes in a Subset of HIV-1-Infected Individuals. Journal of Virology, 2011, 85, 11502-11519.	3.4	168
27	DatabaseÂResources of the National Genomics Data Center, China National Center for Bioinformation in 2021. Nucleic Acids Research, 2021, 49, D18-D28.	14.5	168
28	Database Resources of the National Genomics Data Center in 2020. Nucleic Acids Research, 2020, 48, D24-D33.	14.5	165
29	Vertical T cell immunodominance and epitope entropy determine HIV-1 escape. Journal of Clinical Investigation, 2013, 123, 380-93.	8.2	165
30	Deep functional analysis of synII, a 770-kilobase synthetic yeast chromosome. Science, 2017, 355, .	12.6	163
31	GC-Profile: a web-based tool for visualizing and analyzing the variation of GC content in genomic sequences. Nucleic Acids Research, 2006, 34, W686-W691.	14.5	162
32	Antibodyâ€virus coâ€evolution in <scp>HIV</scp> infection: paths for <scp>HIV</scp> vaccine development. Immunological Reviews, 2017, 275, 145-160.	6.0	160
33	Early Low-Titer Neutralizing Antibodies Impede HIV-1 Replication and Select for Virus Escape. PLoS Pathogens, 2012, 8, e1002721.	4.7	159
34	Database Resources of the BIG Data Center in 2019. Nucleic Acids Research, 2019, 47, D8-D14.	14.5	157
35	Human Non-neutralizing HIV-1 Envelope Monoclonal Antibodies Limit the Number of Founder Viruses during SHIV Mucosal Infection in Rhesus Macaques. PLoS Pathogens, 2015, 11, e1005042.	4.7	145
36	Transmitted/Founder and Chronic Subtype C HIV-1 Use CD4 and CCR5 Receptors with Equal Efficiency and Are Not Inhibited by Blocking the Integrin α4β7. PLoS Pathogens, 2012, 8, e1002686.	4.7	140

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37	DoriC 5.0: an updated database of oriC regions in both bacterial and archaeal genomes. Nucleic Acids Research, 2012, 41, D90-D93.	14.5	128
38	Immune perturbations in HIV-1–infected individuals who make broadly neutralizing antibodies. Science Immunology, 2016, 1, aag0851.	11.9	120
39	Potent and broad HIV-neutralizing antibodies in memory B cells and plasma. Science Immunology, 2017, 2, .	11.9	119
40	DEG 15, an update of the Database of Essential Genes that includes built-in analysis tools. Nucleic Acids Research, 2021, 49, D677-D686.	14.5	119
41	Recurrent Signature Patterns in HIV-1 B Clade Envelope Glycoproteins Associated with either Early or Chronic Infections. PLoS Pathogens, 2011, 7, e1002209.	4.7	114
42	New Software for the Fast Estimation of Population Recombination Rates (FastEPRR) in the Genomic Era. G3: Genes, Genomes, Genetics, 2016, 6, 1563-1571.	1.8	110
43	Codon Usage Optimization of HIV Type 1 Subtype Cgag,pol,env, andnefGenes:In VitroExpression and Immune Responses in DNA-Vaccinated Mice. AIDS Research and Human Retroviruses, 2003, 19, 817-823.	1.1	100
44	Depolarized GABAergic Signaling in Subicular Microcircuits Mediates Generalized Seizure in Temporal Lobe Epilepsy. Neuron, 2017, 95, 92-105.e5.	8.1	97
45	In Vivo gp41 Antibodies Targeting the 2F5 Monoclonal Antibody Epitope Mediate Human Immunodeficiency Virus Type 1 Neutralization Breadth. Journal of Virology, 2009, 83, 3617-3625.	3.4	94
46	An autoreactive antibody from an SLE/HIV-1 individual broadly neutralizes HIV-1. Journal of Clinical Investigation, 2014, 124, 1835-1843.	8.2	93
47	Ancestral and consensus envelope immunogens for HIV-1 subtype C. Virology, 2006, 352, 438-449.	2.4	92
48	DoriC: a database of oriC regions in bacterial genomes. Bioinformatics, 2007, 23, 1866-1867.	4.1	92
49	Completeness of HIV-1 Envelope Glycan Shield at Transmission Determines Neutralization Breadth. Cell Reports, 2018, 25, 893-908.e7.	6.4	91
50	DoriC 10.0: an updated database of replication origins in prokaryotic genomes including chromosomes and plasmids. Nucleic Acids Research, 2019, 47, D74-D77.	14.5	91
51	Therapeutic potential of an anti-high mobility group box-1 monoclonal antibody in epilepsy. Brain, Behavior, and Immunity, 2017, 64, 308-319.	4.1	90
52	Initiation of immune tolerance–controlled HIV gp41 neutralizing B cell lineages. Science Translational Medicine, 2016, 8, 336ra62.	12.4	86
53	Antigenicity and Immunogenicity of Transmitted/Founder, Consensus, and Chronic Envelope Glycoproteins of Human Immunodeficiency Virus Type 1. Journal of Virology, 2013, 87, 4185-4201.	3.4	83
54	Tracking HIV-1 recombination to resolve its contribution to HIV-1 evolution in natural infection. Nature Communications, 2018, 9, 1928.	12.8	83

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55	Evidence of Two Distinct Subsubtypes within the HIV-1 Subtype A Radiation. AIDS Research and Human Retroviruses, 2001, 17, 675-688.	1.1	82
56	Complete genome sequence of Acinetobacter baumannii MDR-TJ and insights into its mechanism of antibiotic resistance. Journal of Antimicrobial Chemotherapy, 2012, 67, 2825-2832.	3.0	82
57	Ori-Finder 2, an integrated tool to predict replication origins in the archaeal genomes. Frontiers in Microbiology, 2014, 5, 482.	3.5	81
58	Features of Recently Transmitted HIV-1 Clade C Viruses that Impact Antibody Recognition: Implications for Active and Passive Immunization. PLoS Pathogens, 2016, 12, e1005742.	4.7	81
59	Genetic Signatures in the Envelope Glycoproteins of HIV-1 that Associate with Broadly Neutralizing Antibodies. PLoS Computational Biology, 2010, 6, e1000955.	3.2	78
60	Initiation of HIV neutralizing B cell lineages with sequential envelope immunizations. Nature Communications, 2017, 8, 1732.	12.8	76
61	A centralized gene-based HIV-1 vaccine elicits broad cross-clade cellular immune responses in rhesus monkeys. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 10489-10494.	7.1	75
62	Impact of Clade, Geography, and Age of the Epidemic on HIV-1 Neutralization by Antibodies. Journal of Virology, 2014, 88, 12623-12643.	3.4	75
63	An Alternative and Effective HIV Vaccination Approach Based on Inhibition of Antigen Presentation Attenuators in Dendritic Cells. PLoS Medicine, 2006, 3, e11.	8.4	74
64	Affinity maturation in an HIV broadly neutralizing B-cell lineage through reorientation of variable domains. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 10275-10280.	7.1	73
65	Evolutionary conservation analysis between the essential and nonessential genes in bacterial genomes. Scientific Reports, 2015, 5, 13210.	3.3	72
66	Maternal HIV-1 envelope–specific antibody responses and reduced risk of perinatal transmission. Journal of Clinical Investigation, 2015, 125, 2702-2706.	8.2	68
67	Antigenicity and immunogenicity of HIV-1 consensus subtype B envelope glycoproteins. Virology, 2007, 360, 218-234.	2.4	67
68	Unselected Mutations in the Human Immunodeficiency Virus Type 1 Genome Are Mostly Nonsynonymous and Often Deleterious. Journal of Virology, 2004, 78, 2426-2433.	3.4	66
69	Strain-Specific V3 and CD4 Binding Site Autologous HIV-1 Neutralizing Antibodies Select Neutralization-Resistant Viruses. Cell Host and Microbe, 2015, 18, 354-362.	11.0	66
70	Rare HIV-1 transmitted/founder lineages identified by deep viral sequencing contribute to rapid shifts in dominant quasispecies during acute and early infection. PLoS Pathogens, 2017, 13, e1006510.	4.7	63
71	Prediction of proteinase cleavage sites in polyproteins of coronaviruses and its applications in analyzing SARS-CoV genomes. FEBS Letters, 2003, 553, 451-456.	2.8	62
72	Cross-Subtype T-Cell Immune Responses Induced by a Human Immunodeficiency Virus Type 1 Group M Consensus Env Immunogen. Journal of Virology, 2006, 80, 6745-6756.	3.4	62

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73	Progress in HIV-1 vaccine development. Journal of Allergy and Clinical Immunology, 2014, 134, 3-10.	2.9	62
74	Comparison of various algorithms for recognizing short coding sequences of human genes. Bioinformatics, 2004, 20, 673-681.	4.1	61
75	Detection of minor drug-resistant populations by parallel allele-specific sequencing. Nature Methods, 2007, 4, 123-125.	19.0	56
76	Dynamic Antibody Specificities and Virion Concentrations in Circulating Immune Complexes in Acute to Chronic HIV-1 Infection. Journal of Virology, 2011, 85, 11196-11207.	3.4	56
77	Significant improvement of oxidase activity through the genetic incorporation of a redox-active unnatural amino acid. Chemical Science, 2015, 6, 3881-3885.	7.4	55
78	Centralized immunogens as a vaccine strategy to overcome HIV-1 diversity. Expert Review of Vaccines, 2004, 3, S161-S168.	4.4	54
79	Recent development of Ori-Finder system and DoriC database for microbial replication origins. Briefings in Bioinformatics, 2019, 20, 1114-1124.	6.5	54
80	Detection of Diverse Variants of Human Immunodeficiency Virus–1 Groups M, N, and O and Simian Immunodeficiency Viruses from Chimpanzees by Using GenericpolandenvPrimer Pairs. Journal of Infectious Diseases, 2000, 181, 1791-1795.	4.0	51
81	Primary Infection by a Human Immunodeficiency Virus with Atypical Coreceptor Tropism. Journal of Virology, 2011, 85, 10669-10681.	3.4	51
82	Impact of immune escape mutations on HIV-1 fitness in the context of the cognate transmitted/founder genome. Retrovirology, 2012, 9, 89.	2.0	50
83	Presence of Diverse Human Immunodeficiency Virus Type 1 Viral Variants in Cameroon. AIDS Research and Human Retroviruses, 2000, 16, 1319-1324.	1.1	49
84	Analysis of Low-Frequency Mutations Associated with Drug Resistance to Raltegravir before Antiretroviral Treatment. Antimicrobial Agents and Chemotherapy, 2011, 55, 1114-1119.	3.2	49
85	Selection and environmental adaptation along a path to speciation in the Tibetan frog <i>Nanorana parkeri</i> . Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E5056-E5065.	7.1	49
86	Toward a high-quality pan-genome landscape of <i>Bacillus subtilis</i> by removal of confounding strains. Briefings in Bioinformatics, 2021, 22, 1951-1971.	6.5	46
87	High throughput functional analysis of HIV-1 env genes without cloning. Journal of Virological Methods, 2007, 143, 104-111.	2.1	45
88	Low-frequency stimulation in anterior nucleus of thalamus alleviates kainate-induced chronic epilepsy and modulates the hippocampal EEG rhythm. Experimental Neurology, 2016, 276, 22-30.	4.1	44
89	A Comprehensive Overview of Online Resources to Identify and Predict Bacterial Essential Genes. Frontiers in Microbiology, 2017, 8, 2331.	3.5	44
90	Zisland Explorer: detect genomic islands by combining homogeneity and heterogeneity properties. Briefings in Bioinformatics, 2017, 18, bbw019.	6.5	43

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91	Transient Receptor Potential Vanilloid 1 Activation by Dietary Capsaicin Promotes Urinary Sodium Excretion by Inhibiting Epithelial Sodium Channel α Subunit–Mediated Sodium Reabsorption. Hypertension, 2014, 64, 397-404.	2.7	42
92	Development of a contemporary globally diverse HIV viral panel by the EQAPOL program. Journal of Immunological Methods, 2014, 409, 117-130.	1.4	42
93	Tissue plasminogen activator (tPA) signal sequence enhances immunogenicity of MVA-based vaccine against tuberculosis. Immunology Letters, 2017, 190, 51-57.	2.5	41
94	Heat shock protein 90 protects rat mesenchymal stem cells against hypoxia and serum deprivation-induced apoptosis via the PI3K/Akt and ERK1/2 pathways. Journal of Zhejiang University: Science B, 2010, 11, 608-617.	2.8	40
95	Functionality of essential genes drives gene strand-bias in bacterial genomes. Biochemical and Biophysical Research Communications, 2010, 396, 472-476.	2.1	40
96	Gene Essentiality Analysis Based on DEG 10, an Updated Database of Essential Genes. Methods in Molecular Biology, 2015, 1279, 219-233.	0.9	40
97	First demonstration of the FLASH effect with ultrahigh dose rate high-energy X-rays. Radiotherapy and Oncology, 2022, 166, 44-50.	0.6	40
98	Postnatally-transmitted HIV-1 Envelope variants have similar neutralization-sensitivity and function to that of nontransmitted breast milk variants. Retrovirology, 2013, 10, 3.	2.0	39
99	An overview of potential inhibitors targeting non-structural proteins 3 (PLpro and Mac1) and 5 (3CLpro/Mpro) of SARS-CoV-2. Computational and Structural Biotechnology Journal, 2021, 19, 4868-4883.	4.1	39
100	HIV-1 did not contribute to the 2019-nCoV genome. Emerging Microbes and Infections, 2020, 9, 378-381.	6.5	38
101	Genome Sequence of Acinetobacter baumannii MDR-TJ. Journal of Bacteriology, 2011, 193, 2365-2366.	2.2	37
102	Protection Principle for a DC Distribution System with a Resistive Superconductive Fault Current Limiter. Energies, 2015, 8, 4839-4852.	3.1	34
103	Amino Acid Changes in the HIV-1 gp41 Membrane Proximal Region Control Virus Neutralization Sensitivity. EBioMedicine, 2016, 12, 196-207.	6.1	34
104	Segmentation algorithm for DNA sequences. Physical Review E, 2005, 72, 041917.	2.1	33
105	Anterior thalamic nucleus stimulation modulates regional cerebral metabolism: An FDG-MicroPET study in rats. Neurobiology of Disease, 2009, 34, 477-483.	4.4	33
106	DeOri: a database of eukaryotic DNA replication origins. Bioinformatics, 2012, 28, 1551-1552.	4.1	32
107	Salicin inhibits AGE-induced degradation of type II collagen and aggrecan in human SW1353 chondrocytes: therapeutic potential in osteoarthritis. Artificial Cells, Nanomedicine and Biotechnology, 2019, 47, 1043-1049.	2.8	30
108	Infant transmitted/founder HIV-1 viruses from peripartum transmission are neutralization resistant to paired maternal plasma. PLoS Pathogens, 2018, 14, e1006944.	4.7	29

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109	Cross-reactive monoclonal antibodies to multiple HIV-1 subtype and SIVcpz envelope glycoproteins. Virology, 2009, 394, 91-98.	2.4	28
110	Complete Sequence of pABTJ2, A Plasmid from Acinetobacter baumannii MDR-TJ, Carrying Many Phage-like Elements. Genomics, Proteomics and Bioinformatics, 2014, 12, 172-177.	6.9	28
111	Centralized HIV-1 Envelope Immunogens and Neutralizing Antibodies. Current HIV Research, 2007, 5, 572-577.	0.5	27
112	Recombination-mediated escape from primary CD8+ T cells in acute HIV-1 infection. Retrovirology, 2014, 11, 69.	2.0	27
113	Protein Localization Analysis of Essential Genes in Prokaryotes. Scientific Reports, 2014, 4, 6001.	3.3	27
114	Quantitative analysis of correlation between AT and GC biases among bacterial genomes. PLoS ONE, 2017, 12, e0171408.	2.5	27
115	Coronavirus phylogeny based on a geometric approach. Molecular Phylogenetics and Evolution, 2005, 36, 224-232.	2.7	26
116	Evolution of Drug-Resistant Viral Populations during Interruption of Antiretroviral Therapy. Journal of Virology, 2011, 85, 6403-6415.	3.4	26
117	Longitudinal Antigenic Sequences and Sites from Intra-Host Evolution (LASSIE) Identifies Immune-Selected HIV Variants. Viruses, 2015, 7, 5443-5475.	3.3	26
118	Comparison of the binding characteristics of SARS-CoV and SARS-CoV-2 RBDs to ACE2 at different temperatures by MD simulations. Briefings in Bioinformatics, 2021, 22, 1122-1136.	6.5	26
119	Identification of Horizontally-transferred Genomic Islands and Genome Segmentation Points by Using the GC Profile Method. Current Genomics, 2014, 15, 113-121.	1.6	26
120	Antiviral Effects of ABMA against Herpes Simplex Virus Type 2 In Vitro and In Vivo. Viruses, 2018, 10, 119.	3.3	25
121	Maternal Broadly Neutralizing Antibodies Can Select for Neutralization-Resistant, Infant-Transmitted/Founder HIV Variants. MBio, 2020, 11, .	4.1	25
122	Lupus gut microbiota transplants cause autoimmunity and inflammation. Clinical Immunology, 2021, 233, 108892.	3.2	25
123	Direct Prediction of Bioaccumulation of Organic Contaminants in Plant Roots from Soils with Machine Learning Models Based on Molecular Structures. Environmental Science & Technology, 2021, 55, 16358-16368.	10.0	25
124	Simultaneous Detection of Major Drug Resistance Mutations in the Protease and Reverse Transcriptase Genes for HIV-1 Subtype C by Use of a Multiplex Allele-Specific Assay. Journal of Clinical Microbiology, 2013, 51, 3666-3674.	3.9	24
125	Detection of Donor's HIV Strain in HIV-Positive Kidney-Transplant Recipient. New England Journal of Medicine, 2020, 382, 195-197.	27.0	24
126	Phosphodiesterase 5 inhibitor, zaprinast, selectively increases cerebral blood flow in the ischemic penumbra in the rat brain. Neurological Research, 2005, 27, 638-643.	1.3	23

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127	Insights into mutualism mechanism and versatile metabolism of Ketogulonicigenium vulgare Hbe602 based on comparative genomics and metabolomics studies. Scientific Reports, 2016, 6, 23068.	3.3	23
128	Comparison of immunogenicity, efficacy and transcriptome changes of inactivated rabies virus vaccine with different adjuvants. Vaccine, 2018, 36, 5020-5029.	3.8	23
129	Extensive Recombination Due to Heteroduplexes Generates Large Amounts of Artificial Gene Fragments during PCR. PLoS ONE, 2014, 9, e106658.	2.5	23
130	Distinct mechanisms of long-term virologic control in two HIV-infected individuals after treatment interruption of anti-retroviral therapy. Nature Medicine, 2021, 27, 1893-1898.	30.7	23
131	Differential response in levels of high-density lipoprotein cholesterol to one-year metformin treatment in prediabetic patients by race/ethnicity. Cardiovascular Diabetology, 2015, 14, 79.	6.8	22
132	Pan-genomic analysis provides novel insights into the association of <i>E.coli</i> with human host and its minimal genome. Bioinformatics, 2019, 35, 1987-1991.	4.1	22
133	Bacteria may have multiple replication origins. Frontiers in Microbiology, 2015, 6, 324.	3.5	21
134	Antiviral effects of Retro-2 cycl and Retro-2.1 against Enterovirus 71 inÂvitro and inÂvivo. Antiviral Research, 2017, 144, 311-321.	4.1	21
135	Identification of HIV-1 genitourinary tract compartmentalization by analyzing the env gene sequences in urine. Aids, 2015, 29, 1651-1657.	2.2	20
136	Genome Sequence of Bacillus endophyticus and Analysis of Its Companion Mechanism in the Ketogulonigenium vulgare-Bacillus Strain Consortium. PLoS ONE, 2015, 10, e0135104.	2.5	20
137	Molecular cloning and recombinant expression of a gene encoding a fungal immunomodulatory protein from Ganoderma lucidum in Pichia pastoris. World Journal of Microbiology and Biotechnology, 2009, 25, 383-390.	3.6	19
138	Recent advances in the genome-wide study of DNA replication origins in yeast. Frontiers in Microbiology, 2015, 6, 117.	3.5	19
139	Recent Advances in the Identification of Replication Origins Based on the Z-curve Method. Current Genomics, 2014, 15, 104-112.	1.6	19
140	CdTe QDs@ZIF-8 composite-based recyclable ratiometric fluorescent sensor for rapid and sensitive detection of chlortetracycline. Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy, 2022, 270, 120785.	3.9	19
141	Identification of amino acid substitutions associated with neutralization phenotype in the human immunodeficiency virus type-1 subtype C gp120. Virology, 2011, 409, 163-174.	2.4	18
142	Transmission of Multiple HIV-1 Subtype C Transmitted/founder Viruses into the Same Recipients Was not Determined by Modest Phenotypic Differences. Scientific Reports, 2016, 6, 38130.	3.3	18
143	HIV-1 Consensus Envelope-Induced Broadly Binding Antibodies. AIDS Research and Human Retroviruses, 2017, 33, 859-868.	1.1	18
144	Exosome-Mediated Delivery of Inducible miR-423-5p Enhances Resistance of MRC-5 Cells to Rabies Virus Infection. International Journal of Molecular Sciences, 2019, 20, 1537.	4.1	18

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145	Exosomes Released from Rabies Virus-Infected Cells May be Involved in the Infection Process. Virologica Sinica, 2019, 34, 59-65.	3.0	18
146	Remote Diffusion-Weighted Imaging Lesions in Intracerebral Hemorrhage: Characteristics, Mechanisms, Outcomes, and Therapeutic Implications. Frontiers in Neurology, 2017, 8, 678.	2.4	17
147	Variable epilepsy phenotypes associated with heterozygous mutation in the SCN9A gene: report of two cases. Neurological Sciences, 2018, 39, 1113-1115.	1.9	17
148	Development of broad neutralization activity in simian/human immunodeficiency virus-infected rhesus macaques after long-term infection. Aids, 2018, 32, 555-563.	2.2	17
149	Origins of replication in <i>Cyanothece</i> 51142. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, E125; author reply E126-7.	7.1	16
150	Enzymes Are Enriched in Bacterial Essential Genes. PLoS ONE, 2011, 6, e21683.	2.5	16
151	Novel intranasal pertussis vaccine based on bacterium-like particles as a mucosal adjuvant. Immunology Letters, 2018, 198, 26-32.	2.5	16
152	Antiviral Effect of Retro-2.1 against Herpes Simplex Virus Type 2 In Vitro. Journal of Microbiology and Biotechnology, 2018, 28, 849-859.	2.1	16
153	Predicting crop root concentration factors of organic contaminants with machine learning models. Journal of Hazardous Materials, 2022, 424, 127437.	12.4	16
154	Identification of the Replication Origins from Cyanothece ATCC 51142 and Their Interactions with the DnaA Protein: From In Silico to In Vitro Studies. Frontiers in Microbiology, 2015, 6, 1370.	3.5	15
155	Fast Dissemination of New HIV-1 CRF02/A1 Recombinants in Pakistan. PLoS ONE, 2016, 11, e0167839.	2.5	15
156	Comparative genomics analysis of the companion mechanisms of Bacillus thuringiensis Bc601 and Bacillus endophyticus Hbe603 in bacterial consortium. Scientific Reports, 2016, 6, 28794.	3.3	15
157	Complete genome sequencing and antibiotics biosynthesis pathways analysis of Streptomyces lydicus 103. Scientific Reports, 2017, 7, 44786.	3.3	15
158	Determination of the cleavage site of enterovirus 71 VPO and the effect of this cleavage on viral infectivity and assembly. Microbial Pathogenesis, 2019, 134, 103568.	2.9	15
159	Thrombin disrupts vascular endothelialâ€cadherin and leads to hydrocephalus via proteaseâ€activated receptorsâ€1 pathway. CNS Neuroscience and Therapeutics, 2019, 25, 1142-1150.	3.9	15
160	Molecular Characterization of a Highly Divergent HIV Type 1 Isolate Obtained Early in the AIDS Epidemic from the Democratic Republic of Congo. AIDS Research and Human Retroviruses, 2001, 17, 1217-1222.	1.1	14
161	Simultaneous Detection of Major Drug Resistance Mutations of HIV-1 Subtype B Viruses from Dried Blood Spot Specimens by Multiplex Allele-Specific Assay. Journal of Clinical Microbiology, 2016, 54, 220-222.	3.9	14
162	An HIV-1 vaccine based on bacterium-like particles elicits Env-specific mucosal immune responses. Immunology Letters, 2020, 222, 29-39.	2.5	14

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163	Increased predominance of HIV-1 CRF01_AE and its recombinants in the Philippines. Journal of General Virology, 2019, 100, 511-522.	2.9	14
164	A reduction in both visceral and subcutaneous fats contributes to increased adiponectin by lifestyle intervention in the Diabetes Prevention Program. Acta Diabetologica, 2015, 52, 625-628.	2.5	13
165	Bioinformatics analysis of molecular mechanisms involved in intervertebral disc degeneration induced by TNF-α and IL-1β. Molecular Medicine Reports, 2016, 13, 2925-2931.	2.4	13
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