George F Gao

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

520	66,239 citations	84	253
papers		h-index	g-index
570 ext. papers	84,554 ext. citations	12.9 avg, IF	8.48 L-index

#	Paper	IF	Citations
520	Target-Based Virtual Screening and LC/MS-Guided Isolation Procedure for Identifying Phloroglucinol-Terpenoid Inhibitors of SARS-CoV-2 <i>Journal of Natural Products</i> , 2022 ,	4.9	2
519	SNX27 suppresses SARS-CoV-2 infection by inhibiting viral lysosome/late endosome entry <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022 , 119,	11.5	7
518	Receptor binding and complex structures of human ACE2 to spike RBD from omicron and delta SARS-CoV-2 <i>Cell</i> , 2022 ,	56.2	48
517	A binding-enhanced but enzymatic activity-eliminated human ACE2 efficiently neutralizes SARS-CoV-2 variants <i>Signal Transduction and Targeted Therapy</i> , 2022 , 7, 10	21	0
516	Effects of a Prolonged Booster Interval on Neutralization of Omicron Variant <i>New England Journal of Medicine</i> , 2022 ,	59.2	15
515	A COVID-19 T-Cell Response Detection Method Based on a Newly Identified Human CD8 T Cell Epitope from SARS-CoV-2 - Hubei Province, China, 2021 <i>China CDC Weekly</i> , 2022 , 4, 83-87	4	0
514	SARS-CoV-2 transmissibility compared between variants of concern and vaccination status <i>Briefings in Bioinformatics</i> , 2022 ,	13.4	2
513	Powassan virus: a tick borne flavivirus infecting humans. <i>Biosafety and Health</i> , 2022 , 4, 30-30	4.7	0
512	Etiological and epidemiological features of acute meningitis or encephalitis in China: a nationwide active surveillance study <i>The Lancet Regional Health - Western Pacific</i> , 2022 , 20, 100361	5	3
511	Nasal delivery of thermostable and broadly neutralizing antibodies protects mice against SARS-CoV-2 infection <i>Signal Transduction and Targeted Therapy</i> , 2022 , 7, 55	21	2
510	A tandem-repeat dimeric RBD protein-based COVID-19 vaccine ZF2001 protects mice and nonhuman primates <i>Emerging Microbes and Infections</i> , 2022 , 1-39	18.9	5
509	Long-Lasting Virus-Specific T Cell Response with Divergent Features in Self-Resolved and Chronic Hepatitis C Virus Patients 35 Years Postinfection <i>ImmunoHorizons</i> , 2022 , 6, 191-201	2.7	
508	PD-1 N58-Glycosylation-Dependent Binding of Monoclonal Antibody Cemiplimab for Immune Checkpoint Therapy <i>Frontiers in Immunology</i> , 2022 , 13, 826045	8.4	2
507	Exploration of immunological responses underpinning severe fever with thrombocytopenia syndrome virus infection reveals IL-6 as a therapeutic target in an immunocompromised mouse model. 2022 , 1, pgac024		0
506	Heterologous BBIBP-CorV/ZF2001 vaccination augments neutralization against SARS-CoV-2 variants: A preliminary observation <i>The Lancet Regional Health - Western Pacific</i> , 2022 , 21, 100440	5	O
505	Phosphosite-dependent presentation of dual phosphorylated peptides by MHC class I molecules <i>IScience</i> , 2022 , 25, 104013	6.1	
504	The "Wolf" Is Indeed Coming: Recombinant "Deltacron" SARS-CoV-2 Detected <i>China CDC Weekly</i> , 2022 , 4, 285-287	4	3

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503	Novel Insights Into the Sulfated Glucuronic Acid-Based Anti-SARS-CoV-2 Mechanism of Exopolysaccharides From Halophilic Archaeon <i>Frontiers in Chemistry</i> , 2022 , 10, 871509	5	0
502	Protective prototype-Beta and Delta-Omicron chimeric RBD-dimer vaccines against SARS-CoV-2 <i>Cell</i> , 2022 ,	56.2	8
501	Transcriptome profiling in swine macrophages infected with African swine fever virus at single-cell resolution <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022 , 119, e2201288119	11.5	4
500	Efficacy and Safety of the RBD-Dimer-Based Covid-19 Vaccine ZF2001 in Adults <i>New England Journal of Medicine</i> , 2022 ,	59.2	21
499	Epidemiological and Clinical Characteristics of Respiratory Syncytial Virus Infections in Children Aged <5 Years in China, from 2014\(\begin{align*} \text{2018}. \textit{Vaccines}, \text{2022}, 10, 810 \end{align*}	5.3	О
498	Safety and immunogenicity of heterologous boost immunization with an adenovirus type-5-vectored and protein-subunit-based COVID-19 vaccine (Convidecia/ZF2001): A randomized, observer-blinded, placebo-controlled trial. <i>PLoS Medicine</i> , 2022 , 19, e1003953	11.6	3
497	The emergence, genomic diversity and global spread of SARS-CoV-2. <i>Nature</i> , 2021 ,	50.4	42
496	Molecular basis of pangolin ACE2 engaged by COVID-19 virus. <i>Chinese Science Bulletin</i> , 2021 , 66, 73-84	2.9	3
495	Landscapes and dynamic diversifications of B-cell receptor repertoires in COVID-19 patients. <i>Human Immunology</i> , 2021 , 83, 119-119	2.3	3
494	Establishment of human distal lung organoids for SARS-CoV-2 infection. <i>Cell Discovery</i> , 2021 , 7, 108	22.3	1
493	Machine Learning Approach Effectively Predicts Binding Between SARS-CoV-2 Spike and ACE2 Across Mammalian Species - Worldwide, 2021. <i>China CDC Weekly</i> , 2021 , 3, 967-972	4	0
492	Immune response pattern across the asymptomatic, symptomatic and convalescent periods of COVID-19. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2021 , 1870, 140736	4	2
491	Metagenome-assembled genomes and gene catalog from the chicken gut microbiome aid in deciphering antibiotic resistomes. <i>Communications Biology</i> , 2021 , 4, 1305	6.7	3
490	Changes in notifiable infectious disease incidence in China during the COVID-19 pandemic. <i>Nature Communications</i> , 2021 , 12, 6923	17.4	6
489	Rapid and sustained containment of covid-19 is achievable and worthwhile: implications for pandemic response. <i>BMJ, The</i> , 2021 , 375, e066169	5.9	3
488	Mooring stone-like Arg pulls diverse bulged peptides: first insight into African swine fever virus-derived T cell epitopes presented by swine MHC class I. <i>Journal of Virology</i> , 2021 , JVI0137821	6.6	Ο
487	COVID-19 Expands Its Territories from Humans to Animals. <i>China CDC Weekly</i> , 2021 , 3, 855-858	4	6
486	Eleven COVID-19 Outbreaks with Local Transmissions Caused by the Imported SARS-CoV-2 Delta VOC - China, July-August, 2021. <i>China CDC Weekly</i> , 2021 , 3, 863-868	4	9

485	Molecular insights into receptor binding of recent emerging SARS-CoV-2 variants. <i>Nature Communications</i> , 2021 , 12, 6103	17.4	24
484	Cross-species recognition of SARS-CoV-2 to bat ACE2. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	32
483	Spatiotemporal visualization for the global COVID-19 surveillance by balloon chart. <i>Infectious Diseases of Poverty</i> , 2021 , 10, 21	10.4	2
482	ADAM17 is an essential attachment factor for classical swine fever virus. <i>PLoS Pathogens</i> , 2021 , 17, e10	0 9 .893	3
481	Antibody seroprevalence in the epicenter Wuhan, Hubei, and six selected provinces after containment of the first epidemic wave of COVID-19 in China. <i>The Lancet Regional Health - Western Pacific</i> , 2021 , 8, 100094	5	21
480	Comparative genomic analysis of mobile colistin resistance gene mcr-9 in Salmonella enterica. <i>Journal of Infection</i> , 2021 , 82, e15-e17	18.9	2
479	Comprehensive large-scale nucleic acid-testing strategies support China's sustained containment of COVID-19. <i>Nature Medicine</i> , 2021 , 27, 740-742	50.5	28
478	Retraction notice to "Diurnal rhythm disruptions induced by chronic unpredictable stress relate to depression-like behaviors in rats" [Pharmacol. Biochem. Behav. 194 (2020) 172939]. <i>Pharmacology Biochemistry and Behavior</i> , 2021 , 203, 173130	3.9	
477	Avian influenza A (H7N9) virus: from low pathogenic to highly pathogenic. <i>Frontiers of Medicine</i> , 2021 , 15, 507-527	12	8
476	Etiological, epidemiological, and clinical features of acute diarrhea in China. <i>Nature Communications</i> , 2021 , 12, 2464	17.4	15
475	Serum sample neutralisation of BBIBP-CorV and ZF2001 vaccines to SARS-CoV-2 501Y.V2. <i>Lancet Microbe, The</i> , 2021 , 2, e285	22.2	36
474	Risk Factors for Death Among the First 80 543 COVID-19 Cases in China: Relationships Between Age, Underlying Disease, Case Severity, and Region. <i>Clinical Infectious Diseases</i> , 2021 ,	11.6	2
473	Diurnal rhythm disruptions induced by chronic unpredictable stress relate to depression-like behaviors in rats. <i>Pharmacology Biochemistry and Behavior</i> , 2021 , 204, 173156	3.9	О
472	Identification of antibiotic resistance genes and associated mobile genetic elements in permafrost. <i>Science China Life Sciences</i> , 2021 , 1	8.5	2
471	NF-B-pathway genes expression in chicken erythrocytes infected with avian influenza virus subtype H9N2. <i>British Poultry Science</i> , 2021 , 62, 666-671	1.9	3
470	Emerging H5N8 avian influenza viruses. <i>Science</i> , 2021 , 372, 784-786	33.3	20
469	Mycobacterium tuberculosis protein kinase G acts as an unusual ubiquitinating enzyme to impair host immunity. <i>EMBO Reports</i> , 2021 , 22, e52175	6.5	5
468	Impact of COVID-19 outbreaks and interventions on influenza in China and the United States. Nature Communications, 2021 , 12, 3249	17.4	33

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467	Molecular basis of cross-species ACE2 interactions with SARS-CoV-2-like viruses of pangolin origin. <i>EMBO Journal</i> , 2021 , 40, e107786	13	13
466	Cryo-EM structures of Lassa and Machupo virus polymerases complexed with cognate regulatory Z proteins identify targets for antivirals. <i>Nature Microbiology</i> , 2021 , 6, 921-931	26.6	6
465	Binding and molecular basis of the bat coronavirus RaTG13 virus to ACE2 in humans and other species. <i>Cell</i> , 2021 , 184, 3438-3451.e10	56.2	30
464	Distinct durability of IgM/IgG antibody responses in COVID-19 patients with differing severity. <i>Science China Life Sciences</i> , 2021 , 1	8.5	2
463	One Hundred Days of Coronavirus Disease 2019 Prevention and Control in China. <i>Clinical Infectious Diseases</i> , 2021 , 72, 332-339	11.6	25
462	Safety and immunogenicity of an inactivated SARS-CoV-2 vaccine, BBIBP-CorV: a randomised, double-blind, placebo-controlled, phase 1/2 trial. <i>Lancet Infectious Diseases, The</i> , 2021 , 21, 39-51	25.5	480
461	Crystal structure of the African swine fever virus core shell protein p15. <i>Biosafety and Health</i> , 2021 , 3, 116-123	4.7	1
460	Viral targets for vaccines against COVID-19. <i>Nature Reviews Immunology</i> , 2021 , 21, 73-82	36.5	402
459	A broadly protective antibody that targets the flavivirus NS1 protein. <i>Science</i> , 2021 , 371, 190-194	33.3	25
458	Long Distance Transmission of SARS-CoV-2 from Contaminated Cold Chain Products to Humans - Qingdao City, Shandong Province, China, September 2020. <i>China CDC Weekly</i> , 2021 , 3, 637-644	4	3
457	Critical role of Syk-dependent STAT1 activation in innate antiviral immunity. Cell Reports, 2021, 34, 108	6 2 7.6	13
456			
	One-year sustained cellular and humoral immunities of COVID-19 convalescents. <i>Clinical Infectious Diseases</i> , 2021 ,	11.6	14
455		11.6	8
455 454	Diseases, 2021, Five Independent Cases of Human Infection with Avian Influenza H5N6 - Sichuan Province, China,		
	Diseases, 2021, Five Independent Cases of Human Infection with Avian Influenza H5N6 - Sichuan Province, China, 2021. China CDC Weekly, 2021, 3, 751-756 Spatial Analysis of People Living with HIV/AIDS Transmitted Through Commercial Heterosexual Contact or Non-Marital Non-Commercial Heterosexual Contact - China, 2018. China CDC Weekly,	4	8
454	Piseases, 2021, Five Independent Cases of Human Infection with Avian Influenza H5N6 - Sichuan Province, China, 2021. China CDC Weekly, 2021, 3, 751-756 Spatial Analysis of People Living with HIV/AIDS Transmitted Through Commercial Heterosexual Contact or Non-Marital Non-Commercial Heterosexual Contact - China, 2018. China CDC Weekly, 2021, 3, 316-319 Interpretation of the Protocol for Prevention and Control of COVID-19 in China (Edition 8). China	4	8 1 10
454 453	Piseases, 2021, Five Independent Cases of Human Infection with Avian Influenza H5N6 - Sichuan Province, China, 2021. China CDC Weekly, 2021, 3, 751-756 Spatial Analysis of People Living with HIV/AIDS Transmitted Through Commercial Heterosexual Contact or Non-Marital Non-Commercial Heterosexual Contact - China, 2018. China CDC Weekly, 2021, 3, 316-319 Interpretation of the Protocol for Prevention and Control of COVID-19 in China (Edition 8). China CDC Weekly, 2021, 3, 527-530 Re-emergence of H5N8 highly pathogenic avian influenza virus in wild birds, China. Emerging	4 4	8 1 10

449	Downregulated miR-451a as a feature of the plasma cfRNA landscape reveals regulatory networks of IL-6/IL-6R-associated cytokine storms in COVID-19 patients. <i>Cellular and Molecular Immunology</i> , 2021 , 18, 1064-1066	15.4	8
448	Structural basis for the inhibition of the SARS-CoV-2 main protease by the anti-HCV drug narlaprevir. <i>Signal Transduction and Targeted Therapy</i> , 2021 , 6, 51	21	7
447	Myocarditis and heart function impairment occur in neonatal mice following in utero exposure to the Zika virus. <i>Journal of Cellular and Molecular Medicine</i> , 2021 , 25, 2730-2733	5.6	1
446	Susceptibility and Attenuated Transmissibility of SARS-CoV-2 in Domestic Cats. <i>Journal of Infectious Diseases</i> , 2021 , 223, 1313-1321	7	28
445	mRNA vaccines: A matter of delivery. <i>EClinicalMedicine</i> , 2021 , 32, 100746	11.3	13
444	Recombinant SARS-CoV-2 RBD with a built in T helper epitope induces strong neutralization antibody response. <i>Vaccine</i> , 2021 , 39, 1241-1247	4.1	8
443	Two immunogenic recombinant protein vaccine candidates showed disparate protective efficacy against Zika virus infection in rhesus macaques. <i>Vaccine</i> , 2021 , 39, 915-925	4.1	0
442	Asymptomatic SARS-CoV-2 Infections Among Persons Entering China From April 16 to October 12, 2020. <i>JAMA - Journal of the American Medical Association</i> , 2021 , 325, 489-492	27.4	20
441	The molecular basis for SARS-CoV-2 binding to dog ACE2. <i>Nature Communications</i> , 2021 , 12, 4195	17.4	17
440	Protective Zika vaccines engineered to eliminate enhancement of dengue infection via immunodominance switch. <i>Nature Immunology</i> , 2021 , 22, 958-968	19.1	4
439	Mobile Colistin Resistance Enzyme MCR-3 Facilitates Bacterial Evasion of Host Phagocytosis. <i>Advanced Science</i> , 2021 , 8, e2101336	13.6	2
438	Overview of SARS-CoV-2 genome-encoded proteins. <i>Science China Life Sciences</i> , 2021 , 1	8.5	11
437	Lessons Learnt From the COVID-19 Pandemic. Frontiers in Public Health, 2021, 9, 694705	6	4
436	O-glycosylation pattern of the SARS-CoV-2 spike protein reveals an "O-Follow-N" rule. <i>Cell Research</i> , 2021 , 31, 1123-1125	24.7	19
435	Safety and immunogenicity of a recombinant tandem-repeat dimeric RBD-based protein subunit vaccine (ZF2001) against COVID-19 in adults: two randomised, double-blind, placebo-controlled, phase 1 and 2 trials. <i>Lancet Infectious Diseases, The</i> , 2021 , 21, 1107-1119	25.5	145
434	Assessing the extent of community spread caused by mink-derived SARS-CoV-2 variants. <i>Innovation(China)</i> , 2021 , 2, 100128	17.8	5
433	More diversified antibiotic resistance genes in chickens and workers of the live poultry markets. <i>Environment International</i> , 2021 , 153, 106534	12.9	11
432	Structural basis of malarial parasite RIFIN-mediated immune escape against LAIR1. <i>Cell Reports</i> , 2021 , 36, 109600	10.6	1

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431	Etiological and epidemiological features of acute respiratory infections in China. <i>Nature Communications</i> , 2021 , 12, 5026	17.4	12
430	A Novel Potentially Recombinant Rodent Coronavirus with a Polybasic Cleavage Site in the Spike Protein. <i>Journal of Virology</i> , 2021 , 95, e0117321	6.6	6
429	Three Novel Avastroviruses Identified in Dead Wild Crows. Virologica Sinica, 2021, 1	6.4	
428	ERASE: a novel nucleic-acid based antiviral mechanism. <i>Cell Research</i> , 2021 , 31, 1142-1143	24.7	
427	Mechanism of Microbial Metabolite Leupeptin in the Treatment of COVID-19 by Traditional Chinese Medicine Herbs. <i>MBio</i> , 2021 , 12, e0222021	7.8	5
426	Tracing the origins of SARS-CoV-2: lessons learned from the past. <i>Cell Research</i> , 2021 , 31, 1139-1141	24.7	7
425	Safety and immunogenicity of an inactivated COVID-19 vaccine, BBIBP-CorV, in people younger than 18 years: a randomised, double-blind, controlled, phase 1/2 trial. <i>Lancet Infectious Diseases, The</i> , 2021 ,	25.5	45
424	Atypical TNF-TNFR superfamily binding interface in the GITR-GITRL complex for Thell activation. <i>Cell Reports</i> , 2021 , 36, 109734	10.6	1
423	Virome of bat-infesting arthropods: highly divergent viruses in different vectors. <i>Journal of Virology</i> , 2021 , JVI0146421	6.6	6
422	Humoral and cellular immunity and the safety of COVID-19 vaccines: a summary of data published by 21 May 2021. <i>International Immunology</i> , 2021 , 33, 529-540	4.9	5
421	The origins of viruses: discovery takes time, international resources, and cooperation. <i>Lancet, The</i> , 2021 , 398, 1401-1402	40	2
420	Peptide Presentations of Marsupial MHC Class I Visualize Immune Features of Lower Mammals Paralleled with Bats. <i>Journal of Immunology</i> , 2021 , 207, 2167-2178	5-3	0
419	Cell entry by SARS-CoV-2. <i>Trends in Biochemical Sciences</i> , 2021 , 46, 848-860	10.3	34
418	Neutralisation of ZF2001-elicited antisera to SARS-CoV-2 variants. <i>Lancet Microbe, The</i> , 2021 , 2, e494	22.2	16
417	Potent inhibition of Severe Acute Respiratory Syndrome Coronavirus 2 by photosensitizers compounds. <i>Dyes and Pigments</i> , 2021 , 194, 109570	4.6	2
416	The changing pattern of enteric pathogen infections in China during the COVID-19 pandemic: a nation-wide observational study. <i>The Lancet Regional Health - Western Pacific</i> , 2021 , 16, 100268	5	7
4 ¹ 5	Recombinant chimpanzee adenovirus AdC7 expressing dimeric tandem-repeat spike protein RBD protects mice against COVID-19. <i>Emerging Microbes and Infections</i> , 2021 , 10, 1574-1588	18.9	3
414	Heterologous prime-boost immunizations with chimpanzee adenoviral vectors elicit potent and protective immunity against SARS-CoV-2 infection <i>Cell Discovery</i> , 2021 , 7, 123	22.3	2

413	N-glycosylation of PD-1 promotes binding of camrelizumab. <i>EMBO Reports</i> , 2020 , 21, e51444	6.5	14
412	Immune restoration in HIV-1-infected patients after 12 years of antiretroviral therapy: a real-world observational study. <i>Emerging Microbes and Infections</i> , 2020 , 9, 2550-2561	18.9	9
411	Plasma IP-10 and MCP-3 levels are highly associated with disease severity and predict the progression of COVID-19. <i>Journal of Allergy and Clinical Immunology</i> , 2020 , 146, 119-127.e4	11.5	358
410	Diversity and abundance of resistome in rhizosphere soil. <i>Science China Life Sciences</i> , 2020 , 63, 1946-194	19 .5	1
409	A noncompeting pair of human neutralizing antibodies block COVID-19 virus binding to its receptor ACE2. <i>Science</i> , 2020 , 368, 1274-1278	33.3	682
408	Combining metagenomics and metatranscriptomics to study human, animal and environmental resistomes. <i>Medicine in Microecology</i> , 2020 , 3, 100014	4.3	4
407	Structural basis of HCoV-19 fusion core and an effective inhibition peptide against virus entry. <i>Emerging Microbes and Infections</i> , 2020 , 9, 1238-1241	18.9	12
406	Structural and Biochemical Characterization of the nsp12-nsp7-nsp8 Core Polymerase Complex from SARS-CoV-2. <i>Cell Reports</i> , 2020 , 31, 107774	10.6	130
405	Development of an Inactivated Vaccine Candidate, BBIBP-CorV, with Potent Protection against SARS-CoV-2. <i>Cell</i> , 2020 , 182, 713-721.e9	56.2	381
404	Transfer cells mediate nitrate uptake to control root nodule symbiosis. <i>Nature Plants</i> , 2020 , 6, 800-808	11.5	14
403	Active case finding with case management: the key to tackling the COVID-19 pandemic. <i>Lancet, The</i> , 2020 , 396, 63-70	40	155
402	Structural insight into arenavirus replication machinery. <i>Nature</i> , 2020 , 579, 615-619	50.4	23
401	Tumor cell-intrinsic PD-1 receptor is a tumor suppressor and mediates resistance to PD-1 blockade therapy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 664	0-665) ⁵¹
400	Integrated metagenomic and metatranscriptomic profiling reveals differentially expressed resistomes in human, chicken, and pig gut microbiomes. <i>Environment International</i> , 2020 , 138, 105649	12.9	21
399	Metagenomic analysis reveals the microbiome and resistome in migratory birds. <i>Microbiome</i> , 2020 , 8, 26	16.6	49
398	A Universal Design of Betacoronavirus Vaccines against COVID-19, MERS, and SARS. <i>Cell</i> , 2020 , 182, 722	2-₹6.3.€	.121 27
397	Recombinant SARS-CoV-2 spike S1-Fc fusion protein induced high levels of neutralizing responses in nonhuman primates. <i>Vaccine</i> , 2020 , 38, 5653-5658	4.1	34
396	Prevalent Eurasian avian-like H1N1 swine influenza virus with 2009 pandemic viral genes facilitating human infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 17204-17210	11.5	98

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395	A distinct name is needed for the new coronavirus. Lancet, The, 2020, 395, 949	40	216
394	A novel coronavirus outbreak of global health concern. <i>Lancet, The</i> , 2020 , 395, 470-473	40	3695
393	A Novel Coronavirus from Patients with Pneumonia in China, 2019. <i>New England Journal of Medicine</i> , 2020 , 382, 727-733	59.2	14511
392	Comparative genomic analysis reveals an 'open' pan-genome of African swine fever virus. Transboundary and Emerging Diseases, 2020 , 67, 1553-1562	4.2	11
391	Genomic characterisation and epidemiology of 2019 novel coronavirus: implications for virus origins and receptor binding. <i>Lancet, The</i> , 2020 , 395, 565-574	40	6394
390	Early Transmission Dynamics in Wuhan, China, of Novel Coronavirus-Infected Pneumonia. <i>New England Journal of Medicine</i> , 2020 , 382, 1199-1207	59.2	8694
389	Structural Analysis of Rabies Virus Glycoprotein Reveals pH-Dependent Conformational Changes and Interactions with a Neutralizing Antibody. <i>Cell Host and Microbe</i> , 2020 , 27, 441-453.e7	23.4	18
388	Adaption and parallel evolution of human-isolated H5 avian influenza viruses. <i>Journal of Infection</i> , 2020 , 80, 630-638	18.9	4
387	The structural basis of African swine fever virus pA104R binding to DNA and its inhibition by stilbene derivatives. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 11000-11009	11.5	13
386	Early Detection of Severe Acute Respiratory Syndrome Coronavirus 2 Antibodies as a Serologic Marker of Infection in Patients With Coronavirus Disease 2019. <i>Clinical Infectious Diseases</i> , 2020 , 71, 200	56-207	2 79
385	CRISPR-Cas13 Inhibitors Block RNA Editing in Bacteria and Mammalian Cells. <i>Molecular Cell</i> , 2020 , 78, 850-861.e5	17.6	32
384	Molecular Basis of Binding between Middle East Respiratory Syndrome Coronavirus and CD26 from Seven Bat Species. <i>Journal of Virology</i> , 2020 , 94,	6.6	12
383	Molecular and structural basis of Echovirus 11 infection by using the dual-receptor system of CD55 and FcRn. <i>Chinese Science Bulletin</i> , 2020 , 65, 67-79	2.9	5
382	Structures of the SARS-CoV-2 nucleocapsid and their perspectives for drug design. <i>EMBO Journal</i> , 2020 , 39, e105938	13	74
381	Recent Trends and Challenges with COVID-19 - Africa, April 4, 2020. China CDC Weekly, 2020, 2, 365-369	94	2
380	Diurnal rhythm disruptions induced by chronic unpredictable stress relate to depression-like behaviors in rats. <i>Pharmacology Biochemistry and Behavior</i> , 2020 , 194, 172939	3.9	O
379	A Novel Coronavirus Genome Identified in a Cluster of Pneumonia Cases - Wuhan, China 2019-2020. <i>China CDC Weekly</i> , 2020 , 2, 61-62	4	71
378	Description of the First Strain of 2019-nCoV, C-Tan-nCoV Wuhan Strain - National Pathogen Resource Center, China, 2020. <i>China CDC Weekly</i> , 2020 , 2, 81-82	4	3

377	Wearing Face Masks - the Simple and Effective Way to Block the Infection Source of COVID-19. <i>China CDC Weekly</i> , 2020 , 2, 268-269	4	3
376	Epidemiologic Changes of Scrub Typhus in China, 1952-2016. Emerging Infectious Diseases, 2020 , 26, 10	91d.10	0110
375	The Genome Resequencing of TCR Loci in Revealed Their Distinct Evolutionary Features in Avians. <i>ImmunoHorizons</i> , 2020 , 4, 33-46	2.7	7
374	A Novel Coronavirus Genome Identified in a Cluster of Pneumonia Cases IWuhan, China 2019I020. <i>China CDC Weekly</i> , 2020 , 2, 61-62	4	329
373	Description of the First Strain of 2019-nCoV, C-Tan-nCoV Wuhan Strain [National Pathogen Resource Center, China, 2020. <i>China CDC Weekly</i> , 2020 , 2, 81-82	4	13
372	Wearing Face Masks Ithe Simple and Effective Way to Block the Infection Source of COVID-19. <i>China CDC Weekly</i> , 2020 , 2, 268-269	4	6
371	The Initial Case of COVID-19 - Shulan City, Jilin Province, China, May 8, 2020. <i>China CDC Weekly</i> , 2020 , 2, 458-459	4	6
370	Fighting Against COVID-19 at the Community Level - Wuhan City, Hubei Province, China, 2020. <i>China CDC Weekly</i> , 2020 , 2, 463-466	4	1
369	A Reemergent Case of COVID-19 - Harbin City, Heilongjiang Province, China, April 9, 2020. <i>China CDC Weekly</i> , 2020 , 2, 460-462	4	5
368	Reemergent Cases of COVID-19 - Xinfadi Wholesales Market, Beijing Municipality, China, June 11, 2020. <i>China CDC Weekly</i> , 2020 , 2, 502-504	4	16
367	Interpretation of the Protocol for Prevention and Control of COVID-19 in China (Edition 7). <i>China CDC Weekly</i> , 2020 , 2, 902-905	4	3
366	Epidemiological Model Suggests D614G Spike Protein Mutation Accelerates Transmission of COVID-19 - Worldwide, 2020. <i>China CDC Weekly</i> , 2020 , 2, 946-947	4	4
365	Detection of mobile colistin resistance gene mcr-9 in carbapenem-resistant Klebsiella pneumoniae strains of human origin in Europe. <i>Journal of Infection</i> , 2020 , 80, 578-606	18.9	22
364	Metagenomic analysis reveals the abundance and diversity of ARGs in children's respiratory tract microbiomes. <i>Journal of Infection</i> , 2020 , 80, 232-254	18.9	2
363	Broad host range of SARS-CoV-2 and the molecular basis for SARS-CoV-2 binding to cat ACE2. <i>Cell Discovery</i> , 2020 , 6, 68	22.3	69
362	Inference of person-to-person transmission of COVID-19 reveals hidden super-spreading events during the early outbreak phase. <i>Nature Communications</i> , 2020 , 11, 5006	17.4	49
361	Molecular basis of Coxsackievirus A10 entry using the two-in-one attachment and uncoating receptor KRM1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 18711-18718	11.5	2
360	Single-Cell Sequencing of Peripheral Mononuclear Cells Reveals Distinct Immune Response Landscapes of COVID-19 and Influenza Patients. <i>Immunity</i> , 2020 , 53, 685-696.e3	32.3	148

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359	Dominant subtype switch in avian influenza viruses during 2016-2019 in China. <i>Nature Communications</i> , 2020 , 11, 5909	17.4	35
358	Molecular basis of EphA2 recognition by gHgL from gammaherpesviruses. <i>Nature Communications</i> , 2020 , 11, 5964	17.4	8
357	Lattice-translocation defects in specific crystals of the catalytic head domain of influenza neuraminidase. <i>Acta Crystallographica Section D: Structural Biology</i> , 2020 , 76, 1057-1064	5.5	O
356	Cold-chain transportation in the frozen food industry may have caused a recurrence of COVID-19 cases in destination: Successful isolation of SARS-CoV-2 virus from the imported frozen cod package surface. <i>Biosafety and Health</i> , 2020 , 2, 199-201	4.7	80
355	Strengthening public health at the community-level in China. Lancet Public Health, The, 2020 , 5, e629-e6	5 30 .4	10
354	Structure-Based Modification of an Anti-neuraminidase Human Antibody Restores Protection Efficacy against the Drifted Influenza Virus. <i>MBio</i> , 2020 , 11,	7.8	5
353	Excretion of SARS-CoV-2 through faecal specimens. <i>Emerging Microbes and Infections</i> , 2020 , 9, 2501-250	08 8.9	23
352	Immune suppression in the early stage of COVID-19 disease. <i>Nature Communications</i> , 2020 , 11, 5859	17.4	75
351	A human neutralizing antibody targets the receptor-binding site of SARS-CoV-2. <i>Nature</i> , 2020 , 584, 120	-15 24 4	844
350	Inception of the Modern Public Health System in China and Perspectives for Effective Control of Emerging Infectious Diseases: In Commemoration of the 140th Anniversary of the Birth of the Plague Fighter Dr. Wu Lien-Teh. <i>Virologica Sinica</i> , 2020 , 35, 868-874	6.4	1
349	Antihypertensive drugs are associated with reduced fatal outcomes and improved clinical characteristics in elderly COVID-19 patients. <i>Cell Discovery</i> , 2020 , 6, 77	22.3	32
348	Dynamic PB2-E627K substitution of influenza H7N9 virus indicates the in vivo genetic tuning and rapid host adaptation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 23807-23814	11.5	10
347	Structures of Echovirus 30 in complex with its receptors inform a rational prediction for enterovirus receptor usage. <i>Nature Communications</i> , 2020 , 11, 4421	17.4	5
346	Both Boceprevir and GC376 efficaciously inhibit SARS-CoV-2 by targeting its main protease. <i>Nature Communications</i> , 2020 , 11, 4417	17.4	195
345	Cryo-EM structure of the varicella-zoster virus A-capsid. <i>Nature Communications</i> , 2020 , 11, 4795	17.4	5
344	Emerging HxNy Influenza A Viruses. Cold Spring Harbor Perspectives in Medicine, 2020,	5.4	11
343	Zika virus induces myocardial immune response and myocarditis in mice. <i>Journal of Molecular and Cellular Cardiology</i> , 2020 , 148, 103-105	5.8	3
342	Regulation of Circadian Genes by the MAPK Pathway: Implications for Rapid Antidepressant Action. <i>Neuroscience Bulletin</i> , 2020 , 36, 66-76	4.3	9

341	Metagenomic data screening reveals the distribution of mobilized resistance genes tet(X), mcr and carbapenemase in animals and humans. <i>Journal of Infection</i> , 2020 , 80, 121-142	18.9	2
340	Genetic and Phylogenetic Characterization of a Chikungunya Virus Imported into Shenzhen, China. <i>Virologica Sinica</i> , 2020 , 35, 115-119	6.4	0
339	Structures of the four Ig-like domain LILRB2 and the four-domain LILRB1 and HLA-G1 complex. <i>Cellular and Molecular Immunology</i> , 2020 , 17, 966-975	15.4	19
338	Phosphorylation Status of Tyrosine 78 Residue Regulates the Nuclear Export and Ubiquitination of Influenza A Virus Nucleoprotein. <i>Frontiers in Microbiology</i> , 2019 , 10, 1816	5.7	5
337	Structural insight into multistage inhibition of CRISPR-Cas12a by AcrVA4. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 18928-18936	11.5	11
336	Influenza Nanovaccines: A Universal Influenza Nanovaccine for Mixing VessellHosts Confers Potential Ability to Block Cross-Species Transmission (Adv. Healthcare Mater. 16/2019). <i>Advanced Healthcare Materials</i> , 2019 , 8, 1970065	10.1	
335	Peptide presentation by bat MHC class I provides new insight into the antiviral immunity of bats. <i>PLoS Biology</i> , 2019 , 17, e3000436	9.7	15
334	Induction of PGRN by influenza virus inhibits the antiviral immune responses through downregulation of type I interferons signaling. <i>PLoS Pathogens</i> , 2019 , 15, e1008062	7.6	17
333	Comparison between human infections caused by highly and low pathogenic H7N9 avian influenza viruses in Wave Five: Clinical and virological findings. <i>Journal of Infection</i> , 2019 , 78, 241-248	18.9	26
332	Diverse biological characteristics and varied virulence of H7N9 from Wave 5. <i>Emerging Microbes and Infections</i> , 2019 , 8, 94-102	18.9	12
331	Pseudorabies virus: a neglected zoonotic pathogen in humans?. <i>Emerging Microbes and Infections</i> , 2019 , 8, 150-154	18.9	44
330	Structural insight into RNA synthesis by influenza D polymerase. <i>Nature Microbiology</i> , 2019 , 4, 1750-17	59 6.6	35
329	Disrupting LILRB4/APOE Interaction by an Efficacious Humanized Antibody Reverses T-cell Suppression and Blocks AML Development. <i>Cancer Immunology Research</i> , 2019 , 7, 1244-1257	12.5	25
328	Salt bridge-forming residues positioned over viral peptides presented by MHC class I impacts T-cell recognition in a binding-dependent manner. <i>Molecular Immunology</i> , 2019 , 112, 274-282	4.3	3
327	Taxonomy of the order Bunyavirales: update 2019. Archives of Virology, 2019, 164, 1949-1965	2.6	148
326	Molecular Basis of Arthritogenic Alphavirus Receptor MXRA8 Binding to Chikungunya Virus Envelope Protein. <i>Cell</i> , 2019 , 177, 1714-1724.e12	56.2	36
325	Naproxen Exhibits Broad Anti-influenza Virus Activity in Mice by Impeding Viral Nucleoprotein Nuclear Export. <i>Cell Reports</i> , 2019 , 27, 1875-1885.e5	10.6	29
324	"Breathing" Hemagglutinin Reveals Cryptic Epitopes for Universal Influenza Vaccine Design. <i>Cell</i> , 2019 , 177, 1086-1088	56.2	7

Human Neonatal Fc Receptor Is the Cellular Uncoating Receptor for Enterovirus B. Cell, 2019, 177, 1553-4665.e46 323 Noc4L-Mediated Ribosome Biogenesis Controls Activation of Regulatory and Conventional T Cells. 10.6 6 322 Cell Reports, 2019, 27, 1205-1220.e4 Influenza virus matrix protein M1 interacts with SLD5 to block host cell cycle. Cellular Microbiology, 321 3.9 5 2019, 21, e13038 A Mycobacterium tuberculosis surface protein recruits ubiquitin to trigger host xenophagy. Nature 320 65 17.4 Communications, 2019, 10, 1973 Light chain modulates heavy chain conformation to change protection profile of monoclonal 8 319 22.3 antibodies against influenza A viruses. Cell Discovery, 2019, 5, 21 HIV prevalence in suspected Ebola cases during the 2014-2016 Ebola epidemic in Sierra Leone. 318 10.4 Infectious Diseases of Poverty, 2019, 8, 15 Comprehensive Clinical and Laboratory Follow-up of a Female Patient With Ebola Virus Disease: 6 1 317 Sierra Leone Ebola Virus Persistence Study. Open Forum Infectious Diseases, 2019, 6, ofz068 The FG Loop of PD-1 Serves as a "Hotspot" for Therapeutic Monoclonal Antibodies in Tumor 316 6.1 15 Immune Checkpoint Therapy. IScience, 2019, 14, 113-124 Intra-host Ebola viral adaption during human infection. Biosafety and Health, 2019, 1, 14-24 315 4.7 7 Antibiotic resistance gene reservoir in live poultry markets. Journal of Infection, 2019, 78, 445-453 18.9 22 314 Neutralization mechanism of human monoclonal antibodies against Rift Valley fever virus. Nature 313 26.6 22 Microbiology, 2019, 4, 1231-1241 E-protein regulatory network links TCR signaling to effector Treg cell differentiation. *Proceedings* 11.5 6 312 of the National Academy of Sciences of the United States of America, 2019, 116, 4471-4480 Strengthening the role of ethnic minority women in science and medicine in China. Lancet, The, 311 40 4 2019, 393, 527-528 Clinical and Immunological Characteristics of Human Infections With H5N6 Avian Influenza Virus. 11.6 310 35 Clinical Infectious Diseases, 2019, 68, 1100-1109 Postfusion structure of human-infecting Bourbon virus envelope glycoprotein. Journal of Structural 309 O 3.4 Biology, 2019, 208, 99-106 308 Public health priorities for China-Africa cooperation. Lancet Public Health, The, 2019, 4, e177-e178 22.4 Clinical and virological characteristics of human infections with H7N9 avian influenza virus in 18.9 307 3 Shenzhen, China, 2013-2017. Journal of Infection, 2019, 79, 389-399 Co-circulation and persistence of multiple A/H3N2 influenza variants in China. Emerging Microbes 18.9 306 11 and Infections, **2019**, 8, 1157-1167

305	Divergent Peptide Presentations of HLA-A30 Alleles Revealed by Structures With Pathogen Peptides. <i>Frontiers in Immunology</i> , 2019 , 10, 1709	8.4	8
304	A Universal Influenza Nanovaccine for "Mixing Vessel" Hosts Confers Potential Ability to Block Cross-Species Transmission. <i>Advanced Healthcare Materials</i> , 2019 , 8, e1900456	10.1	3
303	PARP-1 mediated cell death is directly activated by ZIKV infection. Virology, 2019, 537, 254-262	3.6	10
302	Glycosylation-independent binding of monoclonal antibody toripalimab to FG loop of PD-1 for tumor immune checkpoint therapy. <i>MAbs</i> , 2019 , 11, 681-690	6.6	17
301	Molecular Basis of a Protective/Neutralizing Monoclonal Antibody Targeting Envelope Proteins of both Tick-Borne Encephalitis Virus and Louping Ill Virus. <i>Journal of Virology</i> , 2019 , 93,	6.6	6
300	Avian Influenza A Viruses among Occupationally Exposed Populations, China, 2014-2016. <i>Emerging Infectious Diseases</i> , 2019 , 25, 2215-2225	10.2	19
299	Avian-to-Human Receptor-Binding Adaptation of Avian H7N9 Influenza Virus Hemagglutinin. <i>Cell Reports</i> , 2019 , 29, 2217-2228.e5	10.6	18
298	Crystal Structure of African Swine Fever Virus dUTPase Reveals a Potential Drug Target. <i>MBio</i> , 2019 , 10,	7.8	16
297	Cryo-EM Structure of the African Swine Fever Virus. <i>Cell Host and Microbe</i> , 2019 , 26, 836-843.e3	23.4	56
296	ID2 and ID3 are indispensable for Th1 cell differentiation during influenza virus infection in mice. <i>European Journal of Immunology</i> , 2019 , 49, 476-489	6.1	5
295	Binding mode of the side-by-side two-IgV molecule CD226/DNAM-1 to its ligand CD155/Necl-5. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 988-996	11.5	14
294	Equine-Origin Immunoglobulin Fragments Protect Nonhuman Primates from Ebola Virus Disease. <i>Journal of Virology</i> , 2019 , 93,	6.6	12
293	Structure-function analysis of neutralizing antibodies to H7N9 influenza from naturally infected humans. <i>Nature Microbiology</i> , 2019 , 4, 306-315	26.6	30
292	Low Pathogenic Avian Influenza A (H5N7) Virus Isolated from a Domestic Duck in Dongting Lake Wetland of China, 2016. <i>Virologica Sinica</i> , 2019 , 34, 97-101	6.4	1
291	Double Lock of a Human Neutralizing and Protective Monoclonal Antibody Targeting the Yellow Fever Virus Envelope. <i>Cell Reports</i> , 2019 , 26, 438-446.e5	10.6	26
290	Naturally Occurring Single Mutations in Ebola Virus Observably Impact Infectivity. <i>Journal of Virology</i> , 2019 , 93,	6.6	18
289	Isolation and Identification of a Highly Divergent Kaeng Khoi Virus from Bat Flies (Eucampsipoda sundaica) in China. <i>Vector-Borne and Zoonotic Diseases</i> , 2019 , 19, 73-80	2.4	7
288	Continued reassortment of avian H6 influenza viruses from Southern China, 2014-2016. Transboundary and Emerging Diseases, 2019 , 66, 592-598	4.2	11

(2018-2019)

287	Mycobacterium tuberculosis Mce2E suppresses the macrophage innate immune response and promotes epithelial cell proliferation. <i>Cellular and Molecular Immunology</i> , 2019 , 16, 380-391	15.4	14
286	Crystal Structure of the Capsid Protein from Zika Virus. <i>Journal of Molecular Biology</i> , 2018 , 430, 948-962	26.5	70
285	From "A"IV to "Z"IKV: Attacks from Emerging and Re-emerging Pathogens. <i>Cell</i> , 2018 , 172, 1157-1159	56.2	149
284	Bunyavirales ribonucleoproteins: the viral replication and transcription machinery. <i>Critical Reviews in Microbiology</i> , 2018 , 44, 522-540	7.8	27
283	Vertical Transmission of the Zika Virus Causes Neurological Disorders in Mouse Offspring. <i>Scientific Reports</i> , 2018 , 8, 3541	4.9	25
282	Bat-Origin Coronaviruses Expand Their Host Range to Pigs. <i>Trends in Microbiology</i> , 2018 , 26, 466-470	12.4	34
281	Did ZIKV evolve to be more dangerous? A new clue towards neurovirulence. <i>National Science Review</i> , 2018 , 5, 121-122	10.8	1
2 80	A Sulfonozanamivir Analogue Has Potent Anti-influenza Virus Activity. <i>ChemMedChem</i> , 2018 , 13, 785-78	33 .7	8
279	Endogenous Cellular MicroRNAs Mediate Antiviral Defense against Influenza A Virus. <i>Molecular Therapy - Nucleic Acids</i> , 2018 , 10, 361-375	10.7	45
278	Recombinant Chimpanzee Adenovirus Vaccine AdC7-M/E Protects against Zika Virus Infection and Testis Damage. <i>Journal of Virology</i> , 2018 , 92,	6.6	55
277	Novel cyclo-peptides inhibit Ebola pseudotyped virus entry by targeting primed GP protein. <i>Antiviral Research</i> , 2018 , 155, 1-11	10.8	8
276	Glycan Binding Specificity and Mechanism of Human and Porcine P[6]/P[19] Rotavirus VP8*s. <i>Journal of Virology</i> , 2018 , 92,	6.6	20
275	New Threats from H7N9 Influenza Virus: Spread and Evolution of High- and Low-Pathogenicity Variants with High Genomic Diversity in Wave Five. <i>Journal of Virology</i> , 2018 , 92,	6.6	67
274	Molecular basis of binding between the global post-transcriptional regulator CsrA and the T3SS chaperone CesT. <i>Nature Communications</i> , 2018 , 9, 1196	17.4	17
273	Distinct PD-L1 binding characteristics of therapeutic monoclonal antibody durvalumab. <i>Protein and Cell</i> , 2018 , 9, 135-139	7.2	66
272	The Postfusion Structure of the Heartland Virus Gc Glycoprotein Supports Taxonomic Separation of the Bunyaviral Families Phenuiviridae and Hantaviridae. <i>Journal of Virology</i> , 2018 , 92,	6.6	16
271	Emergence and Adaptation of a Novel Highly Pathogenic H7N9 Influenza Virus in Birds and Humans from a 2013 Human-Infecting Low-Pathogenic Ancestor. <i>Journal of Virology</i> , 2018 , 92,	6.6	72
270	Testing Experimental Therapies in a Guinea Pig Model for Hemorrhagic Fever. <i>Methods in Molecular Biology</i> , 2018 , 1604, 269-278	1.4	1

269	A new threat to human reproduction system posed by Zika virus (ZIKV): From clinical investigations to experimental studies. <i>Virus Research</i> , 2018 , 254, 10-14	6.4	5
268	Serological Investigation of Laboratory-Confirmed and Suspected Ebola Virus Disease Patients During the Late Phase of the Ebola Outbreak in Sierra Leone. <i>Virologica Sinica</i> , 2018 , 33, 323-334	6.4	6
267	Proteomic Analysis of Zika Virus Infected Primary Human Fetal Neural Progenitors Suggests a Role for Doublecortin in the Pathological Consequences of Infection in the Cortex. <i>Frontiers in Microbiology</i> , 2018 , 9, 1067	5.7	22
266	Genomic characterizations of H4 subtype avian influenza viruses from live poultry markets in Sichuan province of China, 2014-2015. <i>Science China Life Sciences</i> , 2018 , 61, 1123-1126	8.5	6
265	Heterosubtypic Protections against Human-Infecting Avian Influenza Viruses Correlate to Biased Cross-T-Cell Responses. <i>MBio</i> , 2018 , 9,	7.8	17
264	Development of a quadruple qRT-PCR assay for simultaneous identification of highly and low pathogenic H7N9 avian influenza viruses and characterization against oseltamivir resistance. <i>BMC Infectious Diseases</i> , 2018 , 18, 406	4	11
263	Zika Virus Envelope Protein and Antibody Complexes. Sub-Cellular Biochemistry, 2018, 88, 147-168	5.5	5
262	Middle East respiratory syndrome coronavirus and bat coronavirus HKU9 both can utilize GRP78 for attachment onto host cells. <i>Journal of Biological Chemistry</i> , 2018 , 293, 11709-11726	5.4	114
261	Building a global atlas of zoonotic viruses. Bulletin of the World Health Organization, 2018, 96, 292-294	8.2	26
260	Transcripts of antibacterial peptides in chicken erythrocytes infected with Marek's disease virus. BMC Veterinary Research, 2018 , 14, 363	2.7	7
259	Evaluation of Zika Virus-specific T-cell Responses in Immunoprivileged Organs of Infected Ifnar1-/-Mice. <i>Journal of Visualized Experiments</i> , 2018 ,	1.6	5
258	On the Centenary of the Spanish Flu: Being Prepared for the Next Pandemic. <i>Virologica Sinica</i> , 2018 , 33, 463-466	6.4	12
257	Clinical, immunological and bacteriological characteristics of H7N9 patients nosocomially co-infected by Acinetobacter Baumannii: a case control study. <i>BMC Infectious Diseases</i> , 2018 , 18, 664	4	5
256	Limited Cross-Linking of 4-1BB by 4-1BB Ligand and the Agonist Monoclonal Antibody Utomilumab. <i>Cell Reports</i> , 2018 , 25, 909-920.e4	10.6	14
255	An Invariant Arginine in Common with MHC Class II Allows Extension at the C-Terminal End of Peptides Bound to Chicken MHC Class I. <i>Journal of Immunology</i> , 2018 , 201, 3084-3095	5.3	11
254	Phylogenomic analysis unravels evolution of yellow fever virus within hosts. <i>PLoS Neglected Tropical Diseases</i> , 2018 , 12, e0006738	4.8	16
253	Structural and functional insights into MCR-2 mediated colistin resistance. <i>Science China Life Sciences</i> , 2018 , 61, 1432-1436	8.5	5
252	Structures of Zika Virus E & NS1: Relations with Virus Infection and Host Immune Responses. <i>Advances in Experimental Medicine and Biology</i> , 2018 , 1062, 77-87	3.6	9

251	Standardized assays for determining the catalytic activity and kinetics of peroxidase-like nanozymes. <i>Nature Protocols</i> , 2018 , 13, 1506-1520	18.8	336
250	CD8 T-Cell Response-Associated Evolution of Hepatitis B Virus Core Protein and Disease Progress. <i>Journal of Virology</i> , 2018 , 92,	6.6	11
249	The first imported case of Rift Valley fever in China reveals a genetic reassortment of different viral lineages. <i>Emerging Microbes and Infections</i> , 2017 , 6, e4	18.9	30
248	An unexpected N-terminal loop in PD-1 dominates binding by nivolumab. <i>Nature Communications</i> , 2017 , 8, 14369	17.4	128
247	The crystal structure of Zika virus NS5 reveals conserved drug targets. <i>EMBO Journal</i> , 2017 , 36, 919-933	3 13	72
246	Human infections with recently-emerging highly pathogenic H7N9 avian influenza virus in China. <i>Journal of Infection</i> , 2017 , 75, 71-75	18.9	115
245	Cryo-EM structures of MERS-CoV and SARS-CoV spike glycoproteins reveal the dynamic receptor binding domains. <i>Nature Communications</i> , 2017 , 8, 15092	17.4	484
244	Virus genomes reveal factors that spread and sustained the Ebola epidemic. <i>Nature</i> , 2017 , 544, 309-315	50.4	238
243	Preliminary Epidemiologic Assessment of Human Infections With Highly Pathogenic Avian Influenza A(H5N6) Virus, China. <i>Clinical Infectious Diseases</i> , 2017 , 65, 383-388	11.6	52
242	An mRNA-based vaccine strategy against Zika. <i>Cell Research</i> , 2017 , 27, 1077-1078	24.7	4
241	Structure of the S1 subunit C-terminal domain from bat-derived coronavirus HKU5 spike protein. <i>Virology</i> , 2017 , 507, 101-109	3.6	9
240	The antibiotic resistome: gene flow in environments, animals and human beings. <i>Frontiers of Medicine</i> , 2017 , 11, 161-168	12	45
239	Protective T Cell Responses Featured by Concordant Recognition of Middle East Respiratory Syndrome Coronavirus-Derived CD8+ T Cell Epitopes and Host MHC. <i>Journal of Immunology</i> , 2017 , 198, 873-882	5.3	32
238	Conserved VII Binding Geometry in a Setting of Locus-Disparate pHLA Recognition by IAT Cell Receptors (TCRs): Insight into Recognition of HIV Peptides by TCRs. <i>Journal of Virology</i> , 2017 , 91,	6.6	8
237	Development of a reverse transcription quantitative polymerase chain reaction-based assay for broad coverage detection of African and Asian Zika virus lineages. <i>Virologica Sinica</i> , 2017 , 32, 199-206	6.4	10
236	Epidemiology of avian influenza A H7N9 virus in human beings across five epidemics in mainland China, 2013-17: an epidemiological study of laboratory-confirmed case series. <i>Lancet Infectious Diseases, The</i> , 2017 , 17, 822-832	25.5	194
235	Alternate binding modes of anti-CRISPR viral suppressors AcrF1/2 to Csy surveillance complex revealed by cryo-EM structures. <i>Cell Research</i> , 2017 , 27, 853-864	24.7	42
234	Structural Biology of the Zika Virus. <i>Trends in Biochemical Sciences</i> , 2017 , 42, 443-456	10.3	75

233	The recombinant N-terminal domain of spike proteins is a potential vaccine against Middle East respiratory syndrome coronavirus (MERS-CoV) infection. <i>Vaccine</i> , 2017 , 35, 10-18	4.1	64
232	Targeted disruption of Noc4l leads to preimplantation embryonic lethality in mice. <i>Protein and Cell</i> , 2017 , 8, 230-235	7.2	3
231	Detection and differentiation of influenza viruses with glycan-functionalized gold nanoparticles. <i>Biosensors and Bioelectronics</i> , 2017 , 91, 46-52	11.8	39
230	Infectious disease trends in China since the SARS outbreak. <i>Lancet Infectious Diseases, The</i> , 2017 , 17, 1113-1115	25.5	11
229	Structures of human-infecting fusogens support a common ancestor with insect baculovirus. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8905-E8912	211.5	17
228	Bacterial effector NleL promotes enterohemorrhagic E. coli-induced attaching and effacing lesions by ubiquitylating and inactivating JNK. <i>PLoS Pathogens</i> , 2017 , 13, e1006534	7.6	20
227	Two classes of protective antibodies against Pseudorabies virus variant glycoprotein B: Implications for vaccine design. <i>PLoS Pathogens</i> , 2017 , 13, e1006777	7.6	22
226	Structural basis of nectin-1 recognition by pseudorabies virus glycoprotein D. <i>PLoS Pathogens</i> , 2017 , 13, e1006314	7.6	26
225	A47 Origin and possible genetic recombination of the middle east respiratory syndrome coronavirus from the first imported case in china: phylogenetics and coalescence analysis. <i>Virus Evolution</i> , 2017 , 3,	3.7	2
224	Remarkably similar CTLA-4 binding properties of therapeutic ipilimumab and tremelimumab antibodies. <i>Oncotarget</i> , 2017 , 8, 67129-67139	3.3	48
223	CTL immunogenicity of Rv3615c antigen and diagnostic performances of an ESAT-6/CFP-10/Rv3615c antigen cocktail for Mycobacterium tuberculosis infection. <i>Tuberculosis</i> , 2017 , 107, 5-12	2.6	3
222	CD8 T Cell Immune Response in Immunocompetent Mice during Zika Virus Infection. <i>Journal of Virology</i> , 2017 , 91,	6.6	79
221	Clinical Evaluation of Ebola Virus Disease Therapeutics. <i>Trends in Molecular Medicine</i> , 2017 , 23, 820-830	11.5	14
220	Structures of phlebovirus glycoprotein Gn and identification of a neutralizing antibody epitope. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E7564-E7573	3 ^{11.5}	58
219	Hemagglutinin-specific CD4 T-cell responses following 2009-pH1N1 inactivated split-vaccine inoculation in humans. <i>Vaccine</i> , 2017 , 35, 5644-5652	4.1	6
218	China in action: national strategies to combat against emerging infectious diseases. <i>Science China Life Sciences</i> , 2017 , 60, 1383-1385	8.5	10
217	The mycobacterial phosphatase PtpA regulates the expression of host genes and promotes cell proliferation. <i>Nature Communications</i> , 2017 , 8, 244	17.4	35
216	Synthesis of Sulfo-Sialic Acid Analogues: Potent Neuraminidase Inhibitors in Regards to Anomeric Functionality. <i>Scientific Reports</i> , 2017 , 7, 8239	4.9	10

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215	Epidemiology, Evolution, and Pathogenesis of H7N9 Influenza Viruses in Five Epidemic Waves since 2013 in China. <i>Trends in Microbiology</i> , 2017 , 25, 713-728	12.4	151
214	Avian-to-Human Receptor-Binding Adaptation by Influenza A Virus Hemagglutinin H4. <i>Cell Reports</i> , 2017 , 20, 1201-1214	10.6	34
213	Conserved peptides enhance immune efficiency of inactive vaccines against emerging avian influenza viruses in chicken. <i>Science China Life Sciences</i> , 2017 , 60, 1340-1347	8.5	6
212	Crystal Structure of the Marburg Virus Nucleoprotein Core Domain Chaperoned by a VP35 Peptide Reveals a Conserved Drug Target for Filovirus. <i>Journal of Virology</i> , 2017 , 91,	6.6	16
211	Differential nucleocytoplasmic shuttling of the nucleoprotein of influenza a viruses and association with host tropism. <i>Cellular Microbiology</i> , 2017 , 19, e12692	3.9	8
210	Structural basis of anti-PD-L1 monoclonal antibody avelumab for tumor therapy. <i>Cell Research</i> , 2017 , 27, 151-153	24.7	72
209	Threonine 80 phosphorylation of non-structural protein 1 regulates the replication of influenza A virus by reducing the binding affinity with RIG-I. <i>Cellular Microbiology</i> , 2017 , 19, e12643	3.9	17
208	T-cell immunity of SARS-CoV: Implications for vaccine development against MERS-CoV. <i>Antiviral Research</i> , 2017 , 137, 82-92	10.8	249
207	Crystal structure of the C-terminal fragment of NS1 protein from yellow fever virus. <i>Science China Life Sciences</i> , 2017 , 60, 1403-1406	8.5	9
206	The persistent prevalence and evolution of cross-family recombinant coronavirus GCCDC1 among a bat population: a two-year follow-up. <i>Science China Life Sciences</i> , 2017 , 60, 1357-1363	8.5	17
205	Human T-cell immunity against the emerging and re-emerging viruses. <i>Science China Life Sciences</i> , 2017 , 60, 1307-1316	8.5	13
204	Induction of Mkp-1 and Nuclear Translocation of Nrf2 by Limonoids from C.DC Protect L-02 Hepatocytes against Acetaminophen-Induced Hepatotoxicity. <i>Frontiers in Pharmacology</i> , 2017 , 8, 653	5.6	11
203	Novel avian influenza A (H5N6) viruses isolated in migratory waterfowl before the first human case reported in China, 2014. <i>Scientific Reports</i> , 2016 , 6, 29888	4.9	46
202	Zika Virus Causes Testis Damage and Leads to Male Infertility in Mice. <i>Cell</i> , 2016 , 167, 1511-1524.e10	56.2	251
201	Treatment with hyperimmune equine immunoglobulin or immunoglobulin fragments completely protects rodents from Ebola virus infection. <i>Scientific Reports</i> , 2016 , 6, 24179	4.9	28
200	Pathogenicity Comparison Between the Kikwit and Makona Ebola Virus Variants in Rhesus Macaques. <i>Journal of Infectious Diseases</i> , 2016 , 214, S281-S289	7	23
199	Effects of a Single Escape Mutation on T Cell and HIV-1 Co-adaptation. <i>Cell Reports</i> , 2016 , 15, 2279-229	110.6	14
198	Ribavirin is effective against drug-resistant H7N9 influenza virus infections. <i>Protein and Cell</i> , 2016 , 7, 611-4	7.2	8

197	Highly pathogenic avian influenza H5N1 Clade 2.3.2.1c virus in migratory birds, 2014-2015. <i>Virologica Sinica</i> , 2016 , 31, 300-5	6.4	28
196	Intra-host dynamics of Ebola virus during 2014. <i>Nature Microbiology</i> , 2016 , 1, 16151	26.6	54
195	More Challenges From Ebola: Infection of the Central Nervous System. <i>Journal of Infectious Diseases</i> , 2016 , 214, S294-S296	7	14
194	Cross-immunity Against Avian Influenza A(H7N9) Virus in the Healthy Population Is Affected by Antigenicity-Dependent Substitutions. <i>Journal of Infectious Diseases</i> , 2016 , 214, 1937-1946	7	18
193	Nosocomial transmission of Clostridium difficile ribotype 027 in a Chinese hospital, 2012-2014, traced by whole genome sequencing. <i>BMC Genomics</i> , 2016 , 17, 405	4.5	26
192	Potent neutralizing monoclonal antibodies against Ebola virus infection. <i>Scientific Reports</i> , 2016 , 6, 258	5.6 .9	36
191	Structural and functional analysis of an anchorless fibronectin-binding protein FBPS from Gram-positive bacterium Streptococcus suis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 13869-13874	11.5	21
190	Crystal clear: visualizing the intervention mechanism of the PD-1/PD-L1 interaction by two cancer therapeutic monoclonal antibodies. <i>Protein and Cell</i> , 2016 , 7, 866-877	7.2	26
189	Structure-Based Tetravalent Zanamivir with Potent Inhibitory Activity against Drug-Resistant Influenza Viruses. <i>Journal of Medicinal Chemistry</i> , 2016 , 59, 6303-12	8.3	20
188	Can Ebola virus become endemic in the human population?. <i>Protein and Cell</i> , 2016 , 7, 4-6	7.2	6
187	Highly diversified Zika viruses imported to China, 2016. <i>Protein and Cell</i> , 2016 , 7, 461-4	7.2	46
186	Swift and Strong NK Cell Responses Protect 129 Mice against High-Dose Influenza Virus Infection. Journal of Immunology, 2016 , 196, 1842-54	5.3	21
185	Ebola Viral Glycoprotein Bound to Its Endosomal Receptor Niemann-Pick C1. <i>Cell</i> , 2016 , 164, 258-268	56.2	165
184	Epidemiology, Genetic Recombination, and Pathogenesis of Coronaviruses. <i>Trends in Microbiology</i> , 2016 , 24, 490-502	12.4	1599
183	Structural Illumination of Equine MHC Class I Molecules Highlights Unconventional Epitope Presentation Manner That Is Evolved in Equine Leukocyte Antigen Alleles. <i>Journal of Immunology</i> , 2016 , 196, 1943-54	5.3	14
182	Changes in the Length of the Neuraminidase Stalk Region Impact H7N9 Virulence in Mice. <i>Journal of Virology</i> , 2016 , 90, 2142-9	6.6	23
181	Dissemination of the mcr-1 colistin resistance gene. <i>Lancet Infectious Diseases, The</i> , 2016 , 16, 146-7	25.5	128
180	Structural basis of collagen recognition by human osteoclast-associated receptor and design of osteoclastogenesis inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 1038-43	11.5	17

(2016-2016)

179	Characterization of clade 2.3.4.4 highly pathogenic H5 avian influenza viruses in ducks and chickens. <i>Veterinary Microbiology</i> , 2016 , 182, 116-22	3.3	51
178	Low Protective Efficacy of the Current Japanese Encephalitis Vaccine against the Emerging Genotype 5 Japanese Encephalitis Virus. <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0004686	4.8	63
177	Rapid HIV-1 Disease Progression in Individuals Infected with a Virus Adapted to Its Host Population. <i>PLoS ONE</i> , 2016 , 11, e0150397	3.7	11
176	An Open Receptor-Binding Cavity of Hemagglutinin-Esterase-Fusion Glycoprotein from Newly-Identified Influenza D Virus: Basis for Its Broad Cell Tropism. <i>PLoS Pathogens</i> , 2016 , 12, e100541	1 ^{7.6}	55
175	A Bat-Derived Putative Cross-Family Recombinant Coronavirus with a Reovirus Gene. <i>PLoS Pathogens</i> , 2016 , 12, e1005883	7.6	70
174	The NS1 gene from bat-derived influenza-like virus H17N10 can be rescued in influenza A PR8 backbone. <i>Journal of General Virology</i> , 2016 , 97, 1797-1806	4.9	11
173	Nosocomial Co-Transmission of Avian Influenza A(H7N9) and A(H1N1)pdm09 Viruses between 2 Patients with Hematologic Disorders. <i>Emerging Infectious Diseases</i> , 2016 , 22, 598-607	10.2	19
172	Melatonin alleviates acute lung injury through inhibiting the NLRP3 inflammasome. <i>Journal of Pineal Research</i> , 2016 , 60, 405-14	10.4	146
171	Cyclophilin A protects mice against infection by influenza A virus. Scientific Reports, 2016, 6, 28978	4.9	14
170	Genesis, Evolution and Prevalence of H5N6 Avian Influenza Viruses in China. <i>Cell Host and Microbe</i> , 2016 , 20, 810-821	23.4	187
169	Two-mAb cocktail protects macaques against the Makona variant of Ebola virus. <i>Science Translational Medicine</i> , 2016 , 8, 329ra33	17.5	62
168	Molecular determinants of human neutralizing antibodies isolated from a patient infected with Zika virus. <i>Science Translational Medicine</i> , 2016 , 8, 369ra179	17.5	152
167	Contribution of intertwined loop to membrane association revealed by Zika virus full-length NS1 structure. <i>EMBO Journal</i> , 2016 , 35, 2170-2178	13	99
166	Structural Insights into the Niemann-Pick C1 (NPC1)-Mediated Cholesterol Transfer and Ebola Infection. <i>Cell</i> , 2016 , 165, 1467-1478	56.2	204
165	Enfuvirtide-PEG conjugate: A potent HIV fusion inhibitor with improved pharmacokinetic properties. <i>European Journal of Medicinal Chemistry</i> , 2016 , 121, 232-237	6.8	17
164	Global epidemiology of avian influenza A H5N1 virus infection in humans, 1997-2015: a systematic review of individual case data. <i>Lancet Infectious Diseases, The</i> , 2016 , 16, e108-e118	25.5	152
163	Diversified mcr-1-Harbouring Plasmid Reservoirs Confer Resistance to Colistin in Human Gut Microbiota. <i>MBio</i> , 2016 , 7, e00177	7.8	105
162	Zika virus NS1 structure reveals diversity of electrostatic surfaces among flaviviruses. <i>Nature Structural and Molecular Biology</i> , 2016 , 23, 456-8	17.6	134

161	Ebola virus encodes a miR-155 analog to regulate importin- expression. <i>Cellular and Molecular Life Sciences</i> , 2016 , 73, 3733-44	10.3	20
160	The Emerging Duck Flavivirus Is Not Pathogenic for Primates and Is Highly Sensitive to Mammalian Interferon Antiviral Signaling. <i>Journal of Virology</i> , 2016 , 90, 6538-6548	6.6	30
159	Structures of the Zika Virus Envelope Protein and Its Complex with a Flavivirus Broadly Protective Antibody. <i>Cell Host and Microbe</i> , 2016 , 19, 696-704	23.4	321
158	Robust Lys63-Linked Ubiquitination of RIG-I Promotes Cytokine Eruption in Early Influenza B Virus Infection. <i>Journal of Virology</i> , 2016 , 90, 6263-6275	6.6	19
157	The Bacterial Mobile Resistome Transfer Network Connecting the Animal and Human Microbiomes. <i>Applied and Environmental Microbiology</i> , 2016 , 82, 6672-6681	4.8	145
156	In vitro assembly of Ebola virus nucleocapsid-like complex expressed in E. coli. <i>Protein and Cell</i> , 2016 , 7, 888-898	7.2	7
155	Characterization of a 2016 Clinical Isolate of Zika Virus in Non-human Primates. <i>EBioMedicine</i> , 2016 , 12, 170-177	8.8	102
154	Resistance to Mutant Group 2 Influenza Virus Neuraminidases of an Oseltamivir-Zanamivir Hybrid Inhibitor. <i>Journal of Virology</i> , 2016 , 90, 10693-10700	6.6	14
153	Carcinoembryonic Antigen-Related Cell Adhesion Molecule 5 Is an Important Surface Attachment Factor That Facilitates Entry of Middle East Respiratory Syndrome Coronavirus. <i>Journal of Virology</i> , 2016 , 90, 9114-27	6.6	56
152	Diversified Anchoring Features the Peptide Presentation of DLA-88*50801: First Structural Insight into Domestic Dog MHC Class I. <i>Journal of Immunology</i> , 2016 , 197, 2306-15	5.3	17
151	MERS-CoV spike protein: Targets for vaccines and therapeutics. <i>Antiviral Research</i> , 2016 , 133, 165-77	10.8	72
150	Two novel reassortants of avian influenza A (H5N6) virus in China. <i>Journal of General Virology</i> , 2015 , 96, 975-981	4.9	79
149	Structural basis for preferential avian receptor binding by the human-infecting H10N8 avian influenza virus. <i>Nature Communications</i> , 2015 , 6, 5600	17.4	27
148	Epidemiology, Evolution, and Recent Outbreaks of Avian Influenza Virus in China. <i>Journal of Virology</i> , 2015 , 89, 8671-6	6.6	177
147	Bat-to-human: spike features determining 'host jump' of coronaviruses SARS-CoV, MERS-CoV, and beyond. <i>Trends in Microbiology</i> , 2015 , 23, 468-78	12.4	363
146	A potent broad-spectrum protective human monoclonal antibody crosslinking two haemagglutinin monomers of influenza A virus. <i>Nature Communications</i> , 2015 , 6, 7708	17.4	101
145	Adaptation of avian influenza A (H6N1) virus from avian to human receptor-binding preference. <i>EMBO Journal</i> , 2015 , 34, 1661-73	13	34
144	MERS in South Korea and China: a potential outbreak threat?. <i>Lancet, The</i> , 2015 , 385, 2349-50	40	62

(2015-2015)

143	Newly Emergent Highly Pathogenic H5N9 Subtype Avian Influenza A Virus. <i>Journal of Virology</i> , 2015 , 89, 8806-15	6.6	11
142	Poultry farms as a source of avian influenza A (H7N9) virus reassortment and human infection. <i>Scientific Reports</i> , 2015 , 5, 7630	4.9	37
141	Nanozyme-strip for rapid local diagnosis of Ebola. <i>Biosensors and Bioelectronics</i> , 2015 , 74, 134-41	11.8	237
140	Genetic diversity and evolutionary dynamics of Ebola virus in Sierra Leone. <i>Nature</i> , 2015 , 524, 93-6	50.4	121
139	Phosphorylation controls the nuclear-cytoplasmic shuttling of influenza A virus nucleoprotein. Journal of Virology, 2015 , 89, 5822-34	6.6	44
138	A humanized neutralizing antibody against MERS-CoV targeting the receptor-binding domain of the spike protein. <i>Cell Research</i> , 2015 , 25, 1237-49	24.7	116
137	Robust expression of vault RNAs induced by influenza A virus plays a critical role in suppression of PKR-mediated innate immunity. <i>Nucleic Acids Research</i> , 2015 , 43, 10321-37	20.1	52
136	MERS, SARS, and Ebola: The Role of Super-Spreaders in Infectious Disease. <i>Cell Host and Microbe</i> , 2015 , 18, 398-401	23.4	224
135	Origin and Possible Genetic Recombination of the Middle East Respiratory Syndrome Coronavirus from the First Imported Case in China: Phylogenetics and Coalescence Analysis. <i>MBio</i> , 2015 , 6, e01280-1	₹ .8	70
134	TCRklass: a new K-string-based algorithm for human and mouse TCR repertoire characterization. <i>Journal of Immunology</i> , 2015 , 194, 446-54	5.3	29
133	Identification of climate factors related to human infection with avian influenza A H7N9 and H5N1 viruses in China. <i>Scientific Reports</i> , 2015 , 5, 18094	4.9	26
132	Recombinant Receptor Binding Domain Protein Induces Partial Protective Immunity in Rhesus Macaques Against Middle East Respiratory Syndrome Coronavirus Challenge. <i>EBioMedicine</i> , 2015 , 2, 143	8 8 -46	87
131	Highly Pathogenic Avian Influenza A(H5N1) Virus Struck Migratory Birds in China in 2015. <i>Scientific Reports</i> , 2015 , 5, 12986	4.9	38
130	Atypical group 1 neuraminidase pH1N1-N1 bound to a group 1 inhibitor. <i>Protein and Cell</i> , 2015 , 6, 771-3	7.2	7
129	The Effects of Socioeconomic and Environmental Factors on the Incidence of Dengue Fever in the Pearl River Delta, China, 2013. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0004159	4.8	53
128	The Serum Profile of Hypercytokinemia Factors Identified in H7N9-Infected Patients can Predict Fatal Outcomes. <i>Scientific Reports</i> , 2015 , 5, 10942	4.9	74
127	Differences in the Epidemiology of Human Cases of Avian Influenza A(H7N9) and A(H5N1) Viruses Infection. <i>Clinical Infectious Diseases</i> , 2015 , 61, 563-71	11.6	52
126	Distribution of sialic acid receptors and experimental infections with different subtypes of influenza A viruses in Qinghai-Tibet plateau wild pika. <i>Virology Journal</i> , 2015 , 12, 63	6.1	8

125	Assessment of the internal genes of influenza A (H7N9) virus contributing to high pathogenicity in mice. <i>Journal of Virology</i> , 2015 , 89, 2-13	6.6	60
124	Identification of the source of A (H10N8) virus causing human infection. <i>Infection, Genetics and Evolution</i> , 2015 , 30, 159-163	4.5	15
123	Influenza viral neuraminidase primes bacterial coinfection through TGF-Emediated expression of host cell receptors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 238-43	11.5	83
122	Mycobacterium tuberculosis suppresses innate immunity by coopting the host ubiquitin system. <i>Nature Immunology</i> , 2015 , 16, 237-45	19.1	98
121	Emergence of Lamivudine-Resistant HBV during Antiretroviral Therapy Including Lamivudine for Patients Coinfected with HIV and HBV in China. <i>PLoS ONE</i> , 2015 , 10, e0134539	3.7	16
120	NK-cells are involved in thymic atrophy induced by influenza A virus infection. <i>Journal of General Virology</i> , 2015 , 96, 3223-3235	4.9	14
119	Molecular characterization of the monoclonal antibodies composing ZMAb: a protective cocktail against Ebola virus. <i>Scientific Reports</i> , 2014 , 4, 6881	4.9	74
118	Genomic and transcriptomic analysis of NDM-1 Klebsiella pneumoniae in spaceflight reveal mechanisms underlying environmental adaptability. <i>Scientific Reports</i> , 2014 , 4, 6216	4.9	18
117	H7N9: a low pathogenic avian influenza A virus infecting humans. <i>Current Opinion in Virology</i> , 2014 , 5, 91-7	7.5	56
116	Clinical and epidemiological characteristics of a fatal case of avian influenza A H10N8 virus	40	434
110	infection: a descriptive study. <i>Lancet, The</i> , 2014 , 383, 714-21	40	TJT
115	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43	3.4	22
	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum		
115	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43 Implementing immunization program efficiently, and establishing national database of adverse	3.4	
115	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43 Implementing immunization program efficiently, and establishing national database of adverse reactions. <i>Science China Life Sciences</i> , 2014 , 57, 553-4	3.4	22
115 114 113	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43 Implementing immunization program efficiently, and establishing national database of adverse reactions. <i>Science China Life Sciences</i> , 2014 , 57, 553-4 Bat-derived influenza-like viruses H17N10 and H18N11. <i>Trends in Microbiology</i> , 2014 , 22, 183-91 Angiotensin II plasma levels are linked to disease severity and predict fatal outcomes in	3·4 8.5	22 217
115 114 113	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43 Implementing immunization program efficiently, and establishing national database of adverse reactions. <i>Science China Life Sciences</i> , 2014 , 57, 553-4 Bat-derived influenza-like viruses H17N10 and H18N11. <i>Trends in Microbiology</i> , 2014 , 22, 183-91 Angiotensin II plasma levels are linked to disease severity and predict fatal outcomes in H7N9-infected patients. <i>Nature Communications</i> , 2014 , 5, 3595 Dynamic reassortments and genetic heterogeneity of the human-infecting influenza A (H7N9) virus.	3·4 8.5 12.4	22 217 119 120
115 114 113 112	Benzophenone C-glucosides and gallotannins from mango tree stem bark with broad-spectrum anti-viral activity. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 2236-43 Implementing immunization program efficiently, and establishing national database of adverse reactions. <i>Science China Life Sciences</i> , 2014 , 57, 553-4 Bat-derived influenza-like viruses H17N10 and H18N11. <i>Trends in Microbiology</i> , 2014 , 22, 183-91 Angiotensin II plasma levels are linked to disease severity and predict fatal outcomes in H7N9-infected patients. <i>Nature Communications</i> , 2014 , 5, 3595 Dynamic reassortments and genetic heterogeneity of the human-infecting influenza A (H7N9) virus. <i>Nature Communications</i> , 2014 , 5, 3142 Enabling the 'host jump': structural determinants of receptor-binding specificity in influenza A	3.4 8.5 12.4 17.4	22 217 119 120

(2014-2014)

107	Bat origins of MERS-CoV supported by bat coronavirus HKU4 usage of human receptor CD26. <i>Cell Host and Microbe</i> , 2014 , 16, 328-37	23.4	198
106	Continuous reassortments with local chicken H9N2 virus underlie the human-infecting influenza A (H7N9) virus in the new influenza season, Guangdong, China. <i>Protein and Cell</i> , 2014 , 5, 878-82	7.2	12
105	Crystal structure of herpes simplex virus 2 gD bound to nectin-1 reveals a conserved mode of receptor recognition. <i>Journal of Virology</i> , 2014 , 88, 13678-88	6.6	29
104	Characteristics of nucleocytoplasmic transport of H1N1 influenza A virus nuclear export protein. <i>Journal of Virology</i> , 2014 , 88, 7455-63	6.6	12
103	The new emerging H7N9 influenza virus indicates poultry as new mixing vessels. <i>Science China Life Sciences</i> , 2014 , 57, 731-2	8.5	7
102	Phylogenetics of varied subtypes of avian influenza viruses in China: potential threat to humans. <i>Protein and Cell</i> , 2014 , 5, 253-7	7.2	27
101	A new unconventional HLA-A2-restricted epitope from HBV core protein elicits antiviral cytotoxic T lymphocytes. <i>Protein and Cell</i> , 2014 , 5, 317-27	7.2	15
100	Assessment of the pathogenesis of Streptococcus suis type 2 infection in piglets for understanding streptococcal toxic shock-like syndrome, meningitis, and sequelae. <i>Veterinary Microbiology</i> , 2014 , 173, 299-309	3.3	15
99	Poultry carrying H9N2 act as incubators for novel human avian influenza viruses. <i>Lancet, The</i> , 2014 , 383, 869	40	80
98	Heat shock protein gp96 adjuvant induces T cell responses and cross-protection to a split influenza vaccine. <i>Vaccine</i> , 2014 , 32, 2703-11	4.1	14
97	Characterization of human #CR repertoire and discovery of D-D fusion in TCRIzhains. <i>Protein and Cell</i> , 2014 , 5, 603-15	7.2	18
96	Characterization of the nucleocytoplasmic shuttle of the matrix protein of influenza B virus. <i>Journal of Virology</i> , 2014 , 88, 7464-73	6.6	8
95	Structure of influenza virus N7: the last piece of the neuraminidase "jigsaw" puzzle. <i>Journal of Virology</i> , 2014 , 88, 9197-207	6.6	33
94	Insights into battles between Mycobacterium tuberculosis and macrophages. <i>Protein and Cell</i> , 2014 , 5, 728-36	7.2	62
93	A motif in LILRB2 critical for Angptl2 binding and activation. <i>Blood</i> , 2014 , 124, 924-35	2.2	57
92	Tailoring subunit vaccine immunity with adjuvant combinations and delivery routes using the Middle East respiratory coronavirus (MERS-CoV) receptor-binding domain as an antigen. <i>PLoS ONE</i> , 2014 , 9, e112602	3.7	64
91	Nterminal acetylation for T cell recognition: molecular basis of MHC class I-restricted ntectylpeptide presentation. <i>Journal of Immunology</i> , 2014 , 192, 5509-19	5.3	9
90	Suppression of interferon lambda signaling by SOCS-1 results in their excessive production during influenza virus infection. <i>PLoS Pathogens</i> , 2014 , 10, e1003845	7.6	76

89	Calculating the burden of disease of avian-origin H7N9 infections in China. BMJ Open, 2014, 4, e004189	3	25
88	Poultry market closures and human infection with influenza A(H7N9) virus, China, 2013-14. <i>Emerging Infectious Diseases</i> , 2014 , 20, 1891-4	10.2	44
87	Human infection with influenza virus A(H10N8) from live poultry markets, China, 2014. <i>Emerging Infectious Diseases</i> , 2014 , 20, 2076-9	10.2	75
86	PILR and PILR have a siglec fold and provide the basis of binding to sialic acid. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 8221-6	11.5	19
85	On the ground in Sierra Leone. <i>Science</i> , 2014 , 346, 666	33.3	19
84	Peptide-dependent conformational fluctuation determines the stability of the human leukocyte antigen class I complex. <i>Journal of Biological Chemistry</i> , 2014 , 289, 24680-90	5.4	27
83	Development and customization of a color-coded microbeads-based assay for drug resistance in HIV-1 reverse transcriptase. <i>PLoS ONE</i> , 2014 , 9, e109823	3.7	
82	Biological features of novel avian influenza A (H7N9) virus. <i>Nature</i> , 2013 , 499, 500-3	50.4	289
81	Molecular basis of binding between novel human coronavirus MERS-CoV and its receptor CD26. <i>Nature</i> , 2013 , 500, 227-31	50.4	466
80	Metagenome-wide analysis of antibiotic resistance genes in a large cohort of human gut microbiota. <i>Nature Communications</i> , 2013 , 4, 2151	17.4	436
79	Bat-derived influenza hemagglutinin H17 does not bind canonical avian or human receptors and most likely uses a unique entry mechanism. <i>Cell Reports</i> , 2013 , 3, 769-78	10.6	79
78	Structures and receptor binding of hemagglutinins from human-infecting H7N9 influenza viruses. <i>Science</i> , 2013 , 342, 243-7	33.3	206
77	Crystal structures of the two membrane-proximal Ig-like domains (D3D4) of LILRB1/B2: alternative models for their involvement in peptide-HLA binding. <i>Protein and Cell</i> , 2013 , 4, 761-70	7.2	10
76	Structure of measles virus hemagglutinin bound to its epithelial receptor nectin-4. <i>Nature Structural and Molecular Biology</i> , 2013 , 20, 67-72	17.6	66
75	Conserved epitopes dominate cross-CD8+ T-cell responses against influenza A H1N1 virus among Asian populations. <i>European Journal of Immunology</i> , 2013 , 43, 2055-69	6.1	24
74	Competition of cell adhesion and immune recognition: insights into the interaction between CRTAM and nectin-like 2. <i>Structure</i> , 2013 , 21, 1430-9	5.2	7
73	Structural basis for the differential classification of HLA-A*6802 and HLA-A*6801 into the A2 and A3 supertypes. <i>Molecular Immunology</i> , 2013 , 55, 381-92	4.3	13
72	Human infection with a novel avian-origin influenza A (H7N9) virus. <i>New England Journal of Medicine</i> , 2013 , 368, 1888-97	59.2	1819

(2011-2013)

71	Origin and diversity of novel avian influenza A H7N9 viruses causing human infection: phylogenetic, structural, and coalescent analyses. <i>Lancet, The</i> , 2013 , 381, 1926-32	40	436
70	An airborne transmissible avian influenza H5 hemagglutinin seen at the atomic level. <i>Science</i> , 2013 , 340, 1463-7	33.3	92
69	Lessons learnt from the human infections of avian-origin influenza A H7N9 virus: live free markets and human health. <i>Science China Life Sciences</i> , 2013 , 56, 493-4	8.5	36
68	Structure and receptor-binding properties of an airborne transmissible avian influenza A virus hemagglutinin H5 (VN1203mut). <i>Protein and Cell</i> , 2013 , 4, 502-11	7.2	29
67	Duck egg drop syndrome virus: an emerging Tembusu-related flavivirus in China. <i>Science China Life Sciences</i> , 2013 , 56, 701-10	8.5	44
66	Structure of the fusion core and inhibition of fusion by a heptad repeat peptide derived from the S protein of Middle East respiratory syndrome coronavirus. <i>Journal of Virology</i> , 2013 , 87, 13134-40	6.6	118
65	Severe fever with thrombocytopenia syndrome virus expands its borders. <i>Emerging Microbes and Infections</i> , 2013 , 2, e36	18.9	10
64	Characterization of two distinct neuraminidases from avian-origin human-infecting H7N9 influenza viruses. <i>Cell Research</i> , 2013 , 23, 1347-55	24.7	77
63	Live-animal markets and influenza A (H7N9) virus infection. <i>New England Journal of Medicine</i> , 2013 , 368, 2337-9	59.2	108
62	A replicating modified vaccinia tiantan strain expressing an avian-derived influenza H5N1 hemagglutinin induce broadly neutralizing antibodies and cross-clade protective immunity in mice. <i>PLoS ONE</i> , 2013 , 8, e83274	3.7	8
61	Transport of influenza virus neuraminidase (NA) to host cell surface is regulated by ARHGAP21 and Cdc42 proteins. <i>Journal of Biological Chemistry</i> , 2012 , 287, 9804-9816	5.4	62
60	Guidelines for the use and interpretation of assays for monitoring autophagy. <i>Autophagy</i> , 2012 , 8, 445-	5 44 .2	2783
59	Genomic and antigenic characterization of the newly emerging Chinese duck egg-drop syndrome flavivirus: genomic comparison with Tembusu and Sitiawan viruses. <i>Journal of General Virology</i> , 2012 , 93, 2158-2170	4.9	78
58	Crystal structure of cell adhesion molecule nectin-2/CD112 and its binding to immune receptor DNAM-1/CD226. <i>Journal of Immunology</i> , 2012 , 188, 5511-20	5.3	44
57	Cross-allele cytotoxic T lymphocyte responses against 2009 pandemic H1N1 influenza A virus among HLA-A24 and HLA-A3 supertype-positive individuals. <i>Journal of Virology</i> , 2012 , 86, 13281-94	6.6	33
56	Narrow groove and restricted anchors of MHC class I molecule BF2*0401 plus peptide transporter restriction can explain disease susceptibility of B4 chickens. <i>Journal of Immunology</i> , 2012 , 189, 4478-87	5.3	42
55	Major Histocompatibility Complex: Interaction with Peptides 2011,		2
54	Binding of herpes simplex virus glycoprotein D to nectin-1 exploits host cell adhesion. <i>Nature Communications</i> , 2011 , 2, 577	17.4	66

53	Structural basis of cross-allele presentation by HLA-A*0301 and HLA-A*1101 revealed by two HIV-derived peptide complexes. <i>Molecular Immunology</i> , 2011 , 49, 395-401	4.3	21
52	A friend to man, IDr. Feifan Tang: a story of causative agent of trachoma, from IIang virus Ito Chlamydia trachomatis, to Phylum Chlamydiae (Protein and Cell, 2011, 2, 349-350)	7.2	3
51	Special features of the 2009 pandemic swine-origin influenza A H1N1 hemagglutinin and neuraminidase. <i>Science Bulletin</i> , 2011 , 56, 1747-1752		13
50	An infectious clone of the highly pathogenic porcine reproductive and respiratory syndrome virus: Topology of glycoprotein 3 (GP3) addressing the intrachain disulfide bonds. <i>Science Bulletin</i> , 2011 , 56, 2785-2793		1
49	A strategy to produce monoclonal antibodies against gp96 by prime-boost regimen using endogenous protein and E. coli heterologously-expressed fragment. <i>Central South University</i> , 2011 , 18, 1857-1864		5
48	Two distinct conformations of a rinderpest virus epitope presented by bovine major histocompatibility complex class I N*01801: a host strategy to present featured peptides. <i>Journal of Virology</i> , 2011 , 85, 6038-48	6.6	35
47	Diverse peptide presentation of rhesus macaque major histocompatibility complex class I Mamu-A 02 revealed by two peptide complex structures and insights into immune escape of simian immunodeficiency virus. <i>Journal of Virology</i> , 2011 , 85, 7372-83	6.6	17
46	Enterovirus 71 and coxsackievirus A16 3C proteases: binding to rupintrivir and their substrates and anti-hand, foot, and mouth disease virus drug design. <i>Journal of Virology</i> , 2011 , 85, 10319-31	6.6	70
45	Revival of the identification of cytotoxic T-lymphocyte epitopes for immunological diagnosis, therapy and vaccine development. <i>Experimental Biology and Medicine</i> , 2011 , 236, 253-67	3.7	32
44	Influenza A virus N5 neuraminidase has an extended 150-cavity. <i>Journal of Virology</i> , 2011 , 85, 8431-5	6.6	51
43	Crystal structure of swine major histocompatibility complex class I SLA-1 0401 and identification of 2009 pandemic swine-origin influenza A H1N1 virus cytotoxic T lymphocyte epitope peptides. <i>Journal of Virology</i> , 2011 , 85, 11709-24	6.6	54
42	Crystal structure of leukocyte Ig-like receptor LILRB4 (ILT3/LIR-5/CD85k): a myeloid inhibitory receptor involved in immune tolerance. <i>Journal of Biological Chemistry</i> , 2011 , 286, 18013-25	5.4	33
41	Duck egg-drop syndrome caused by BYD virus, a new Tembusu-related flavivirus. <i>PLoS ONE</i> , 2011 , 6, e18106	3.7	231
40	The 2009 pandemic H1N1 neuraminidase N1 lacks the 150-cavity in its active site. <i>Nature Structural and Molecular Biology</i> , 2010 , 17, 1266-8	17.6	141
39	The membrane protein of severe acute respiratory syndrome coronavirus acts as a dominant immunogen revealed by a clustering region of novel functionally and structurally defined cytotoxic T-lymphocyte epitopes. <i>Journal of Infectious Diseases</i> , 2010 , 202, 1171-80	7	86
38	Novel immunodominant peptide presentation strategy: a featured HLA-A*2402-restricted cytotoxic T-lymphocyte epitope stabilized by intrachain hydrogen bonds from severe acute respiratory syndrome coronavirus nucleocapsid protein. <i>Journal of Virology</i> , 2010 , 84, 11849-57	6.6	94
37	Serological surveillance of influenza A virus infection in swine populations in Fujian province, China: no evidence of naturally occurring H5N1 infection in pigs. <i>Zoonoses and Public Health</i> , 2010 , 57, 291-8	2.9	26
36	A dimeric structure of PD-L1: functional units or evolutionary relics?. <i>Protein and Cell</i> , 2010 , 1, 153-60	7.2	50

(2004-2010)

35	Crystal structure of the swine-origin A (H1N1)-2009 influenza A virus hemagglutinin (HA) reveals similar antigenicity to that of the 1918 pandemic virus. <i>Protein and Cell</i> , 2010 , 1, 459-67	7.2	85
34	It is not just AIV: from avian to swine-origin influenza virus. Science China Life Sciences, 2010, 53, 151-1.	53 8.5	21
33	Structural immunology and crystallography help immunologists see the immune system in action: how T and NK cells touch their ligands. <i>IUBMB Life</i> , 2009 , 61, 579-90	4.7	23
32	Interspecies transmission and host restriction of avian H5N1 influenza virus. <i>Science in China Series C: Life Sciences</i> , 2009 , 52, 428-38		27
31	Website for avian flu information and bioinformatics. <i>Science in China Series C: Life Sciences</i> , 2009 , 52, 470-3		6
30	Crystal structure of myeloid cell activating receptor leukocyte Ig-like receptor A2 (LILRA2/ILT1/LIR-7) domain swapped dimer: molecular basis for its non-binding to MHC complexes. <i>Journal of Molecular Biology</i> , 2009 , 386, 841-53	6.5	17
29	1SA2-08 Avian influenza virus: from migratory bird infection to peptide inhibitor discovery(1SA2 BSJ&ABA Joint Symposium, "Prion and Virus Infections",The 47th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2009 , 49, S4	О	
28	H5N1 avian influenza re-emergence of Lake Qinghai: phylogenetic and antigenic analyses of the newly isolated viruses and roles of migratory birds in virus circulation. <i>Journal of General Virology</i> , 2008 , 89, 697-702	4.9	87
27	Towards our understanding of SARS-CoV, an emerging and devastating but quickly conquered virus. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2007 , 30, 309-27	2.6	27
26	First glimpse of the peptide presentation by rhesus macaque MHC class I: crystal structures of Mamu-A*01 complexed with two immunogenic SIV epitopes and insights into CTL escape. <i>Journal of Immunology</i> , 2007 , 178, 944-52	5.3	30
25	Screening and identification of severe acute respiratory syndrome-associated coronavirus-specific CTL epitopes. <i>Journal of Immunology</i> , 2006 , 177, 2138-45	5.3	85
24	Highly pathogenic H5N1 influenza virus infection in migratory birds. <i>Science</i> , 2005 , 309, 1206	33.3	540
23	Filling the hole: evidence of a small molecule binding to the fusion core pocket in human respiratory syncytial virus. <i>Expert Opinion on Investigational Drugs</i> , 2005 , 14, 195-7	5.9	2
22	Generation of murine CTL by a hepatitis B virus-specific peptide and evaluation of the adjuvant effect of heat shock protein glycoprotein 96 and its terminal fragments. <i>Journal of Immunology</i> , 2005 , 174, 195-204	5.3	72
21	Design and Characterization of Human Respiratory Syncytial Virus Entry Inhibitors. <i>Antiviral Therapy</i> , 2005 , 10, 833-840	1.6	7
20	Crystal structure of severe acute respiratory syndrome coronavirus spike protein fusion core. <i>Journal of Biological Chemistry</i> , 2004 , 279, 49414-9	5.4	159
19	Complex assembly, crystallization and preliminary X-ray crystallographic studies of MHC H-2Kd complexed with an HBV-core nonapeptide. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 1473-5		24
18	Following the rule: formation of the 6-helix bundle of the fusion core from severe acute respiratory syndrome coronavirus spike protein and identification of potent peptide inhibitors. <i>Biochemical and Biophysical Research Communications</i> , 2004 , 319, 283-8	3.4	83

17	The crystal structures of severe acute respiratory syndrome virus main protease and its complex with an inhibitor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 13190-5	11.5	712
16	Molecular coordination of alphabeta T-cell receptors and coreceptors CD8 and CD4 in their recognition of peptide-MHC ligands. <i>Trends in Immunology</i> , 2002 , 23, 408-13	14.4	96
15	Molecular interactions of coreceptor CD8 and MHC class I: the molecular basis for functional coordination with the T-cell receptor. <i>Trends in Immunology</i> , 2000 , 21, 630-6		119
14	Production of soluble alphabeta T-cell receptor heterodimers suitable for biophysical analysis of ligand binding. <i>Protein Science</i> , 1999 , 8, 2418-23	6.3	40
13	Antagonism of cytotoxic T-lymphocyte activation by soluble CD8. <i>Nature Medicine</i> , 1999 , 5, 399-404	50.5	34
12	Assembly and crystallization of the complex between the human T cell coreceptor CD8alpha homodimer and HLA-A2. <i>Protein Science</i> , 1998 , 7, 1245-9	6.3	25
11	Crystal structure of the complex between human CD8alpha(alpha) and HLA-A2. <i>Nature</i> , 1997 , 387, 630-4	4 50.4	388
10	Reduced sera neutralization to Omicron SARS-CoV-2 by both inactivated and protein subunit vaccines and the convalescents		7
9	Structural and biochemical characterization of nsp12-nsp7-nsp8 core polymerase complex from COVID-19 virus		1
8	A non-competing pair of human neutralizing antibodies block COVID-19 virus binding to its receptor ACE2		13
7	Safety and immunogenicity of a recombinant tandem-repeat dimeric RBD protein vaccine against COVID-19 in adults: pooled analysis of two randomized, double-blind, placebo-controlled, phase 1 and 2 trials		13
6	Landscapes and dynamic diversifications of B-cell receptor repertoires in COVID-19 patients		1
5	A tandem-repeat dimeric RBD protein-based COVID-19 vaccine ZF2001 protects mice and nonhuman primates		5
4	Comparative evaluation of the transmissibility of SARS-CoV-2 variants of concern		1
3	Neutralization of SARS-CoV-2 VOC 501Y.V2 by human antisera elicited by both inactivated BBIBP-CorV and recombinant dimeric RBD ZF2001 vaccines		37
2	Neutralization of recombinant RBD-subunit vaccine ZF2001-elicited antisera to SARS-CoV-2 variants including Delta		6
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