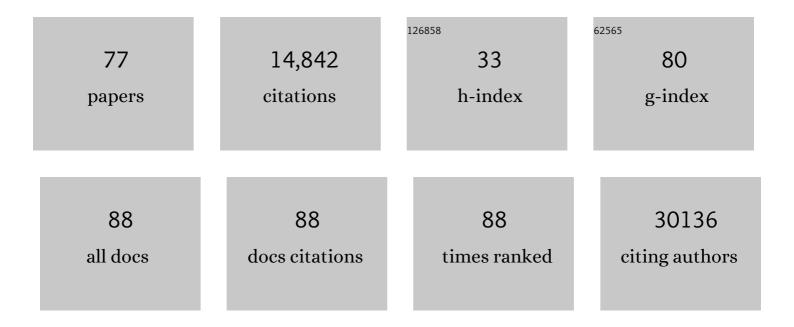
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Virological assessment of hospitalized patients with COVID-2019. Nature, 2020, 581, 465-469.	13.7	5,822
2	Transmission of 2019-nCoV Infection from an Asymptomatic Contact in Germany. New England Journal of Medicine, 2020, 382, 970-971.	13.9	3,343
3	Real-time, portable genome sequencing for Ebola surveillance. Nature, 2016, 530, 228-232.	13.7	1,179
4	Multilevel proteomics reveals host perturbations by SARS-CoV-2 and SARS-CoV. Nature, 2021, 594, 246-252.	13.7	475
5	Investigation of a COVID-19 outbreak in Germany resulting from a single travel-associated primary case: a case series. Lancet Infectious Diseases, The, 2020, 20, 920-928.	4.6	383
6	Experimental Treatment with Favipiravir for Ebola Virus Disease (the JIKI Trial): A Historically Controlled, Single-Arm Proof-of-Concept Trial in Guinea. PLoS Medicine, 2016, 13, e1001967.	3.9	382
7	Virus genomes reveal factors that spread and sustained the Ebola epidemic. Nature, 2017, 544, 309-315.	13.7	346
8	Temporal and spatial analysis of the 2014–2015 Ebola virus outbreak in West Africa. Nature, 2015, 524, 97-101.	13.7	272
9	Unique human immune signature of Ebola virus disease in Guinea. Nature, 2016, 533, 100-104.	13.7	170
10	Virus Detection and Monitoring of Viral Load in Crimean-Congo Hemorrhagic Fever Virus Patients. Emerging Infectious Diseases, 2007, 13, 1097-1100.	2.0	112
11	Rapid outbreak sequencing of Ebola virus in Sierra Leone identifies transmission chains linked to sporadic cases. Virus Evolution, 2016, 2, vew016.	2.2	105
12	Clinical and virological features of first human monkeypox cases in Germany. Infection, 2023, 51, 265-270.	2.3	105
13	Persistence and clearance of Ebola virus RNA from seminal fluid of Ebola virus disease survivors: a longitudinal analysis and modelling study. The Lancet Global Health, 2017, 5, e80-e88.	2.9	100
14	SARS-CoV-2 antibody testing—questions to be asked. Journal of Allergy and Clinical Immunology, 2020, 146, 35-43.	1.5	82
15	A new Puumala hantavirus subtype in rodents associated with an outbreak of Nephropathia epidemica in South-East Germany in 2004. Epidemiology and Infection, 2006, 134, 1333-1344.	1.0	68
16	Molecular identification of adenovirus sequences: A rapid scheme for early typing of human adenoviruses in diagnostic samples of immunocompetent and immunodeficient patients. Journal of Medical Virology, 2006, 78, 1210-1217.	2.5	67
17	Diagnostics of tick-borne rickettsioses in Germany: A modern concept for a neglected disease. International Journal of Medical Microbiology, 2008, 298, 368-374.	1.5	66
18	Diagnostic Assays for Crimean-Congo Hemorrhagic Fever. Emerging Infectious Diseases, 2012, 18, 1958-1965.	2.0	66

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19	Yersinia pestis Lineages in Mongolia. PLoS ONE, 2012, 7, e30624.	1.1	64
20	Typhus and Other Rickettsioses. Deutsches Ärzteblatt International, 2009, 106, 348-54.	0.6	63
21	Multicentre comparison of quantitative PCR-based assays to detect SARS-CoV-2, Germany, March 2020. Eurosurveillance, 2020, 25, .	3.9	60
22	Dilemmas in Managing Pregnant Women With Ebola: 2 Case Reports: Table 1 Clinical Infectious Diseases, 2016, 62, 903-905.	2.9	56
23	In Search of the SARS-CoV-2 Protection Correlate: Head-to-Head Comparison of Two Quantitative S1 Assays in Pre-characterized Oligo-/Asymptomatic Patients. Infectious Diseases and Therapy, 2021, 10, 1505-1518.	1.8	53
24	Detection of a questing Hyalomma marginatum marginatum adult female (Acari, Ixodidae) in southern Germany. Experimental and Applied Acarology, 2007, 43, 227-231.	0.7	52
25	Prevalence and Risk Factors of Infection in the Representative COVID-19 Cohort Munich. International Journal of Environmental Research and Public Health, 2021, 18, 3572.	1.2	47
26	Different features of Vδ2 T and NK cells in fatal and non-fatal human Ebola infections. PLoS Neglected Tropical Diseases, 2017, 11, e0005645.	1.3	46
27	Mobile diagnostics in outbreak response, not only for Ebola: a blueprint for a modular and robust field laboratory. Eurosurveillance, 2015, 20, .	3.9	45
28	Rickettsia raoultii, the predominant Rickettsia found in Mongolian Dermacentor nuttalli. Ticks and Tick-borne Diseases, 2012, 3, 227-231.	1.1	40
29	Rickettsia spp. in Ixodes ricinus Ticks in Bavaria, Germany. Annals of the New York Academy of Sciences, 2006, 1078, 509-511.	1.8	39
30	Laboratory Findings, Compassionate Use of Favipiravir, and Outcome in Patients With Ebola Virus Disease, Guinea, 2015—A Retrospective Observational Study. Journal of Infectious Diseases, 2019, 220, 195-202.	1.9	38
31	Phylogenetic analysis of Puumala virus subtype Bavaria, characterization and diagnostic use of its recombinant nucleocapsid protein. Virus Genes, 2011, 43, 177-191.	0.7	35
32	COVID-19 in Patients Receiving CD20-depleting Immunochemotherapy for B-cell Lymphoma. HemaSphere, 2021, 5, e603.	1.2	35
33	Seroepidemiological study in a Puumala virus outbreak area in South-East Germany. Medical Microbiology and Immunology, 2009, 198, 83-91.	2.6	34
34	Low-Density Macroarray for Rapid Detection and Identification of Crimean-Congo Hemorrhagic Fever Virus. Journal of Clinical Microbiology, 2009, 47, 1025-1030.	1.8	32
35	Analysis of Diagnostic Findings From the European Mobile Laboratory in Guéckédou, Guinea, March 2014 Through March 2015. Journal of Infectious Diseases, 2016, 214, S250-S257.	1.9	32
36	Simultaneous Detection of Ricin and Abrin DNA by Real-Time PCR (qPCR). Toxins, 2012, 4, 633-642.	1.5	31

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37	First International External Quality Assessment of Molecular Detection of Crimean-Congo Hemorrhagic Fever Virus. PLoS Neglected Tropical Diseases, 2012, 6, e1706.	1.3	30
38	High prevalence of genetically diverse Borrelia bavariensis-like strains in Ixodes persulcatus from Selenge Aimag, Mongolia. Ticks and Tick-borne Diseases, 2013, 4, 89-92.	1.1	30
39	Deep Sequencing of RNA from Blood and Oral Swab Samples Reveals the Presence of Nucleic Acid from a Number of Pathogens in Patients with Acute Ebola Virus Disease and Is Consistent with Bacterial Translocation across the Gut. MSphere, 2017, 2, .	1.3	30
40	Comparison of two commercial surrogate ELISAs to detect a neutralising antibody response to SARS-CoV-2. Journal of Virological Methods, 2021, 292, 114122.	1.0	30
41	Isolation, preliminary characterization, and full-genome analyses of tick-borne encephalitis virus from Mongolia. Virus Genes, 2012, 45, 413-425.	0.7	29
42	Results of the first pilot external quality assessment (EQA) scheme for anti-SARS-CoV2-antibody testing. Clinical Chemistry and Laboratory Medicine, 2020, 58, 2121-2130.	1.4	26
43	A pair of noncompeting neutralizing human monoclonal antibodies protecting from disease in a SARSâ€CoVâ€2 infection model. European Journal of Immunology, 2022, 52, 770-783.	1.6	24
44	Head-to-head evaluation of seven different seroassays including direct viral neutralisation in a representative cohort for SARS-CoV-2. Journal of General Virology, 2021, 102, .	1.3	21
45	Isolation and characterization of lytic phage TUN1 specific for Klebsiella pneumoniae K64 clinical isolates from Tunisia. BMC Microbiology, 2021, 21, 186.	1.3	19
46	Detection of Coxiella burnetii in heart valve sections by fluorescence in situ hybridization. Journal of Medical Microbiology, 2018, 67, 537-542.	0.7	18
47	Screening for carbapenemases in ertapenem-resistant Enterobacteriaceae collected at a Tunisian hospital between 2014 and 2018. European Journal of Microbiology and Immunology, 2019, 9, 9-13.	1.5	17
48	High seroprevalence for indigenous spotted fever group rickettsiae in forestry workers from the federal state of Brandenburg, Eastern Germany. Ticks and Tick-borne Diseases, 2017, 8, 132-138.	1.1	15
49	Viral Hemorrhagic Fever Cases in the Country of Georgia: Acute Febrile Illness Surveillance Study Results. American Journal of Tropical Medicine and Hygiene, 2014, 91, 246-248.	0.6	14
50	Establishment of a specimen panel for the decentralised technical evaluation of the sensitivity of 31 rapid diagnostic tests for SARS-CoV-2 antigen, Germany, September 2020 to April 2021. Eurosurveillance, 2021, 26, .	3.9	14
51	Ebola: missed opportunities for Europe–Africa research. Lancet Infectious Diseases, The, 2015, 15, 1254-1255.	4.6	13
52	Seroprevalence of tick-borne and mosquito-borne arboviruses in European brown hares in Northern and Western Germany. International Journal of Medical Microbiology, 2006, 296, 80-83.	1.5	12
53	Lassa and Crimean-Congo Hemorrhagic Fever Viruses, Mali. Emerging Infectious Diseases, 2019, 25, 999-1002.	2.0	12
54	Evaluation of sampling technique and transport media for the diagnostics of adenoviral eye infections. Graefe's Archive for Clinical and Experimental Ophthalmology, 2006, 244, 1497-1504.	1.0	11

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55	Phylogenetic Placement of Isolates Within the Trans-Eurasian Clade A.Br.008/009 of Bacillus anthracis. Microorganisms, 2019, 7, 689.	1.6	11
56	In vitro evaluation of the effect of mutations in primer binding sites on detection of SARS-CoV-2 by RT-qPCR. Journal of Virological Methods, 2022, 299, 114352.	1.0	11
57	Evaluation of Two Rapid Lateral Flow Tests and Two Surrogate ELISAs for the Detection of SARS-CoV-2 Specific Neutralizing Antibodies. Frontiers in Medicine, 2022, 9, 820151.	1.2	11
58	Whole genome sequencing and phylogenetic classification of Tunisian SARS-CoV-2 strains from patients of the Military Hospital in Tunis. Virus Genes, 2020, 56, 767-771.	0.7	10
59	The interplay of viral loads, clinical presentation, and serological responses in SARS-CoV-2 – Results from a prospective cohort of outpatient COVID-19 cases. Virology, 2022, 569, 37-43.	1.1	9
60	Assessing the Quality of Serological Testing in the COVID-19 Pandemic: Results of a European External Quality Assessment (EQA) Scheme for Anti-SARS-CoV-2 Antibody Detection. Journal of Clinical Microbiology, 2021, 59, e0055921.	1.8	8
61	Emerging SARS-CoV-2 variant B.1.1.7 reduces neutralisation activity of antibodies against wild-type SARS-CoV-2. Journal of Clinical Virology, 2021, 142, 104912.	1.6	8
62	Development of a versatile and stable internal control system for RT-qPCR assays. Journal of Virological Methods, 2014, 208, 33-40.	1.0	7
63	First Phylogenetic Analysis of Malian SARS-CoV-2 Sequences Provides Molecular Insights into the Genomic Diversity of the Sahel Region. Viruses, 2020, 12, 1251.	1.5	7
64	Isolation and whole genome analysis of endospore-forming bacteria from heroin. Forensic Science International: Genetics, 2018, 32, 1-6.	1.6	6
65	The identification of novel single nucleotide polymorphisms to assist in mapping the spread of Bacillus anthracis across the Southern Caucasus. Scientific Reports, 2018, 8, 11254.	1.6	6
66	Prevalence of Middle East Respiratory Syndrome Coronavirus in Dromedary Camels, Tunisia. Emerging Infectious Diseases, 2021, 27, 1964-1968.	2.0	6
67	Prevalence and Risk Factors of Infection in the Representative COVID-19 Cohort Munich. SSRN Electronic Journal, 0, , .	0.4	6
68	EvaGreen based real-time RT-PCR assay for broad-range detection of hantaviruses in the field. Journal of Clinical Virology, 2013, 58, 334-335.	1.6	5
69	Non-randomised Ebola trials—lessons for optimal outbreak research. Lancet Infectious Diseases, The, 2016, 16, 407-408.	4.6	5
70	NOTIFy (non-toxic lyophilized field)-FISH for the identification of biological agents by Fluorescence in situ Hybridization. PLoS ONE, 2020, 15, e0230057.	1.1	4
71	Rapid detection of SARS-CoV-2 by pulse-controlled amplification (PCA). Journal of Virological Methods, 2021, 290, 114083.	1.0	4
72	Technical Note: Simple, scalable, and sensitive protocol for retrieving Bacillus anthracis (and other) Tj ETQq0 0	0 rgBT /Ov	erlogk 10 Tf 50

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73	Assessment of SARS-CoV-2 rapid antigen tests. Journal of Laboratory Medicine, 2021, 45, 143-148.	1.1	3
74	Sensitivity of two SARS-CoV-2 variants with spike protein mutations to neutralising antibodies. Virus Genes, 2021, 57, 502-509.	0.7	3
75	Genome Sequence of Historical Bacillus anthracis Strain Tyrol 4675 Isolated from a Bovine Anthrax Case in Austria. Genome Announcements, 2017, 5, .	0.8	2
76	PCR based prevalence study of <i>Francisella tularensis</i> in Kharkiv, Dnipropetrovsk, and Mykolaiv oblasts during 2015–2018. Journal of Veterinary Research (Poland), 2020, 64, 63-71.	0.3	1
77	P1038 Seroprevalence of IgG antibodies againstfiaviviruses in German soldiers. International Journal of Antimicrobial Agents, 2007, 29, S278-S279.	1.1	0