

# M Ciccozzi

## List of Publications by Year in descending order

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421  
papers

11,706  
citations

47006

47  
h-index

53230

85  
g-index

434  
all docs

434  
docs citations

434  
times ranked

17337  
citing authors

#	ARTICLE	IF	CITATIONS
1	Emerging SARS-CoV-2 mutation hot spots include a novel RNA-dependent-RNA polymerase variant. <i>Journal of Translational Medicine</i> , 2020, 18, 179.	4.4	784
2	The COVID-19 pandemic. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2020, 57, 365-388.	6.1	737
3	The 2019â€new coronavirus epidemic: Evidence for virus evolution. <i>Journal of Medical Virology</i> , 2020, 92, 455-459.	5.0	508
4	Application of the ARIMA model on the COVID-2019 epidemic dataset. <i>Data in Brief</i> , 2020, 29, 105340.	1.0	455
5	COVIDâ€™2019: The role of the nsp2 and nsp3 in its pathogenesis. <i>Journal of Medical Virology</i> , 2020, 92, 584-588.	5.0	308
6	The first two cases of 2019â€™nCoV in Italy: Where they come from?. <i>Journal of Medical Virology</i> , 2020, 92, 518-521.	5.0	263
7	COVID-19 Outbreak: An Overview. <i>Chemotherapy</i> , 2019, 64, 215-223.	1.6	240
8	Evolutionary analysis of SARS-CoV-2: how mutation of Non-Structural Protein 6 (NSP6) could affect viral autophagy. <i>Journal of Infection</i> , 2020, 81, e24-e27.	3.3	211
9	The global spread of 2019-nCoV: a molecular evolutionary analysis. <i>Pathogens and Global Health</i> , 2020, 114, 64-67.	2.3	161
10	Sars-CoV-2 Envelope and Membrane Proteins: Structural Differences Linked to Virus Characteristics?. <i>BioMed Research International</i> , 2020, 2020, 1-6.	1.9	150
11	Fecal and Mucosal Microbiota Profiling in Irritable Bowel Syndrome and Inflammatory Bowel Disease. <i>Frontiers in Microbiology</i> , 2019, 10, 1655.	3.5	146
12	Transmission of HIV Drug Resistance and the Predicted Effect on Current First-line Regimens in Europe. <i>Clinical Infectious Diseases</i> , 2016, 62, 655-663.	5.8	135
13	Evolution patterns of SARS-CoV-2: Snapshot on its genome variants. <i>Biochemical and Biophysical Research Communications</i> , 2021, 538, 88-91.	2.1	121
14	Elevated aspartic proteinase secretion and experimental pathogenicity of <i>Candida albicans</i> isolates from oral cavities of subjects infected with human immunodeficiency virus. <i>Infection and Immunity</i> , 1996, 64, 466-471.	2.2	120
15	The Presence of Antiâ€™at Antibodies Is Predictive of Longâ€™Term Nonprogression to AIDS or Severe Immunodeficiency: Findings in a Cohort of HIVâ€™1 Seroconverters. <i>Journal of Infectious Diseases</i> , 2005, 191, 1321-1324.	4.0	118
16	A novel methodology for large-scale phylogeny partition. <i>Nature Communications</i> , 2011, 2, 321.	12.8	118
17	HIV-1 and HCV sequences from Libyan outbreak. <i>Nature</i> , 2006, 444, 836-837.	27.8	114
18	Mayaro Virus in Child with Acute Febrile Illness, Haiti, 2015. <i>Emerging Infectious Diseases</i> , 2016, 22, 2000-2002.	4.3	113

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19	Zika Virus Outbreak in Haiti in 2014: Molecular and Clinical Data. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004687.	3.0	106
20	The impact of the hepatitis B mass immunisation campaign on the incidence and risk factors of acute hepatitis B in Italy. <i>Journal of Hepatology</i> , 2000, 33, 980-985.	3.7	101
21	Inactivation of Hepatitis A virus in heat-treated mussels. <i>Journal of Applied Microbiology</i> , 1999, 87, 884-888.	3.1	97
22	Impact of lockdown on Covid-19 case fatality rate and viral mutations spread in 7 countries in Europe and North America. <i>Journal of Translational Medicine</i> , 2020, 18, 338.	4.4	86
23	First detection of SARS-CoV-2 spike protein N501 mutation in Italy in August, 2020. <i>Lancet Infectious Diseases</i> , 2021, 21, e147.	9.1	84
24	Chikungunya virus, epidemiology, clinics and phylogenesis: A review. <i>Asian Pacific Journal of Tropical Medicine</i> , 2014, 7, 925-932.	0.8	82
25	SARS-CoV-2 B.1.617 Indian variants: Are electrostatic potential changes responsible for a higher transmission rate?. <i>Journal of Medical Virology</i> , 2021, 93, 6551-6556.	5.0	79
26	SARS-Cov-2 ORF3a: Mutability and function. <i>International Journal of Biological Macromolecules</i> , 2021, 170, 820-826.	7.5	77
27	Sequence Conservation and Antibody Cross-Recognition of Clade B Human Immunodeficiency Virus (HIV) Type 1 Tat Protein in HIV-1 Infected Italians, Ugandans, and South Africans. <i>Journal of Infectious Diseases</i> , 2003, 188, 1171-1180.	4.0	75
28	Emerging of a SARS-CoV-2 viral strain with a deletion in nsp1. <i>Journal of Translational Medicine</i> , 2020, 18, 329.	4.4	71
29	Enigmatic origin of hepatitis B virus: An ancient travelling companion or a recent encounter?. <i>World Journal of Gastroenterology</i> , 2014, 20, 7622.	3.3	71
30	Long-term protection against SHIV89.6P replication in HIV-1 Tat vaccinated cynomolgus monkeys. <i>Vaccine</i> , 2004, 22, 3258-3269.	3.8	70
31	Gut mucosal-associated microbiota better discloses inflammatory bowel disease differential patterns than faecal microbiota. <i>Digestive and Liver Disease</i> , 2019, 51, 648-656.	0.9	67
32	Emergence of recombinant Mayaro virus strains from the Amazon basin. <i>Scientific Reports</i> , 2017, 7, 8718.	3.3	66
33	Unusual microorganisms and antimicrobial resistances in a group of Syrian migrants: Sentinel surveillance data from an asylum seekers centre in Italy. <i>Travel Medicine and Infectious Disease</i> , 2016, 14, 115-122.	3.0	65
34	Protective efficacy of a multicomponent vector vaccine in cynomolgus monkeys after intrarectal simian immunodeficiency virus challenge. <i>Journal of General Virology</i> , 2004, 85, 1191-1201.	2.9	63
35	Analysis of the ORFK1 hypervariable regions reveal distinct HHV-8 clustering in Kaposi's sarcoma and non-Kaposi's cases. <i>Journal of Experimental and Clinical Cancer Research</i> , 2015, 34, 1.	8.6	62
36	The COVID-19 pandemic: viral variants and vaccine efficacy. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2022, 59, 66-75.	6.1	61

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37	High-Resolution Molecular Epidemiology and Evolutionary History of HIV-1 Subtypes in Albania. PLoS ONE, 2008, 3, e1390.	2.5	60
38	The electrostatic potential of the Omicron variant spike is higher than in Delta and Delta plus variants: A hint to higher transmissibility?. Journal of Medical Virology, 2022, 94, 1277-1280.	5.0	60
39	Epidemiological history and phylogeography of West Nile virus lineage 2. Infection, Genetics and Evolution, 2013, 17, 46-50.	2.3	58
40	Spatial and Temporal Dynamics of Hepatitis B Virus D Genotype in Europe and the Mediterranean Basin. PLoS ONE, 2012, 7, e37198.	2.5	55
41	Population Dynamics of HIV-1 Subtype B in a Cohort of Men-Having-Sex-With-Men in Rome, Italy. Journal of Acquired Immune Deficiency Syndromes (1999), 2010, 55, 156-160.	2.1	53
42	Phylogeography and epidemiological history of West Nile virus genotype 1a in Europe and the Mediterranean basin. Infection, Genetics and Evolution, 2011, 11, 646-653.	2.3	53
43	Changing epidemiology of parenterally transmitted viral hepatitis: results from the hepatitis surveillance system in Italy. Digestive and Liver Disease, 2001, 33, 778-784.	0.9	51
44	Identification of Merkel cell polyomavirus in the lower respiratory tract of Italian patients. Journal of Medical Virology, 2010, 82, 505-509.	5.0	51
45	Think globally, act locally: Phylodynamic reconstruction of infectious bronchitis virus (IBV) QX genotype (GI-19 lineage) reveals different population dynamics and spreading patterns when evaluated on different epidemiological scales. PLoS ONE, 2017, 12, e0184401.	2.5	51
46	High Sequence Conservation of Human Immunodeficiency Virus Type 1 Reverse Transcriptase under Drug Pressure despite the Continuous Appearance of Mutations. Journal of Virology, 2005, 79, 10718-10729.	3.4	50
47	Procalcitonin and MR-Proadrenomedullin Combination with SOFA and qSOFA Scores for Sepsis Diagnosis and Prognosis: A Diagnostic Algorithm. Shock, 2018, 50, 44-52.	2.1	48
48	Efficacy of radiofrequency ablation in autonomous functioning thyroid nodules. A systematic review and meta-analysis. Reviews in Endocrine and Metabolic Disorders, 2019, 20, 37-44.	5.7	48
49	A doubt of multiple introduction of SARS-CoV-2 in Italy: A preliminary overview. Journal of Medical Virology, 2020, 92, 1634-1636.	5.0	48
50	A persistently replicating SARS-CoV-2 variant derived from an asymptomatic individual. Journal of Translational Medicine, 2020, 18, 362.	4.4	46
51	SARS-CoV-2 infection affects the lower urinary tract and male genital system: A systematic review. Journal of Medical Virology, 2021, 93, 3133-3142.	5.0	46
52	Co-circulation of Dengue and Chikungunya Viruses, Al Hudaydah, Yemen, 2012. Emerging Infectious Diseases, 2014, 20, 1351-1354.	4.3	45
53	A cluster of the new SARS-CoV-2 B.1.621 lineage in Italy and sensitivity of the viral isolate to the BNT162b2 vaccine. Journal of Medical Virology, 2021, 93, 6468-6470.	5.0	45
54	Increase in transmitted resistance to non-nucleoside reverse transcriptase inhibitors among newly diagnosed HIV-1 infections in Europe. BMC Infectious Diseases, 2014, 14, 407.	2.9	43

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55	Procalcitonin, MR-Proadrenomedullin, and Cytokines Measurement in Sepsis Diagnosis: Advantages from Test Combination. <i>Disease Markers</i> , 2015, 2015, 1-14.	1.3	42
56	The ongoing COVID-19 epidemic in Minas Gerais, Brazil: insights from epidemiological data and SARS-CoV-2 whole genome sequencing. <i>Emerging Microbes and Infections</i> , 2020, 9, 1824-1834.	6.5	42
57	Origin and evolution of Nipah virus. <i>Journal of Medical Virology</i> , 2016, 88, 380-388.	5.0	41
58	Closed-Circuit System for the Depuration of Mussels Experimentally Contaminated with Hepatitis A Virus. <i>Journal of Food Protection</i> , 2001, 64, 877-880.	1.7	40
59	Excretion of the novel polyomaviruses KI and WU in the stool of patients with hematological disorders. <i>Journal of Medical Virology</i> , 2009, 81, 1668-1673.	5.0	40
60	The human polyomaviruses <scp>KI</scp> and <scp>WU</scp>: virological background and clinical implications. <i>Apmis</i> , 2013, 121, 746-754.	2.0	40
61	Update on epidemiology of HCV in Italy: focus on the Calabria Region. <i>BMC Infectious Diseases</i> , 2014, 14, S2.	2.9	40
62	Coinfection With Zika and Dengue-2 Viruses in a Traveler Returning From Haiti, 2016: Clinical Presentation and Genetic Analysis. <i>Clinical Infectious Diseases</i> , 2017, 64, 72-75.	5.8	39
63	SARS-CoV-2 Lineages and Sub-Lineages Circulating Worldwide: A Dynamic Overview. <i>Chemotherapy</i> , 2021, 66, 3-7.	1.6	39
64	Extended Genetic Diversity of Bovine Viral Diarrhea Virus and Frequency of Genotypes and Subtypes in Cattle in Italy between 1995 and 2013. <i>BioMed Research International</i> , 2014, 2014, 1-8.	1.9	38
65	Point-of-Care Screening, Prevalence, and Risk Factors for Hepatitis B Infection Among 3,728 Mainly Undocumented Migrants From Non-EU Countries in Northern Italy. <i>Journal of Travel Medicine</i> , 2015, 22, 78-86.	3.0	38
66	Molecular epidemiology, evolution and phylogeny of Chikungunya virus: An updating review. <i>Infection, Genetics and Evolution</i> , 2016, 41, 270-278.	2.3	38
67	Tick-borne encephalitis in north-east Italy: a 14-year retrospective study, January 2000 to December 2013. <i>Eurosurveillance</i> , 2015, 20, .	7.0	38
68	Human immunodeficiency virus integrase inhibitors efficiently suppress feline immunodeficiency virus replication in vitro and provide a rationale to redesign antiretroviral treatment for feline AIDS. <i>Retrovirology</i> , 2007, 4, 79.	2.0	37
69	Severe Acute Respiratory Syndrome Coronavirus Sequence Characteristics and Evolutionary Rate Estimate from Maximum Likelihood Analysis. <i>Journal of Virology</i> , 2004, 78, 1602-1603.	3.4	35
70	Matrix-assisted laser desorption ionization time-of-flight mass spectrometry in clinical microbiology: An updating review. <i>Infection, Genetics and Evolution</i> , 2019, 76, 104063.	2.3	35
71	The HIV Type 1 Epidemic in Bulgaria Involves Multiple Subtypes and Is Sustained by Continuous Viral Inflow from West and East European Countries. <i>AIDS Research and Human Retroviruses</i> , 2008, 24, 771-779.	1.1	34
72	Identification of the novel KI and WU polyomaviruses in human tonsils. <i>Journal of Clinical Virology</i> , 2009, 46, 75-79.	3.1	34

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73	Phyldynamic applications in 21st century global infectious disease research. <i>Global Health Research and Policy</i> , 2017, 2, 13.	3.6	34
74	Mediterranean diet and Psoriatic Arthritis activity: a multicenter cross-sectional study. <i>Rheumatology International</i> , 2020, 40, 951-958.	3.0	34
75	Origin, evolution, and phylogeography of recent epidemic CHIKV strains. <i>Infection, Genetics and Evolution</i> , 2012, 12, 392-398.	2.3	33
76	Molecular Diversity of HIV in Albania. <i>Journal of Infectious Diseases</i> , 2005, 192, 475-479.	4.0	32
77	Identification of the novel KI Polyomavirus in paranasal and lung tissues. <i>Journal of Medical Virology</i> , 2009, 81, 558-561.	5.0	32
78	Could nasal irrigation and oral rinse reduce the risk for COVID-19 infection?. <i>International Journal of Immunopathology and Pharmacology</i> , 2020, 34, 205873842094175.	2.1	32
79	Non-toxigenic environmental <i>Vibrio cholerae</i> O1 strain from Haiti provides evidence of pre-pandemic cholera in Hispaniola. <i>Scientific Reports</i> , 2016, 6, 36115.	3.3	31
80	Susceptibility to measles in migrant population: implication for policy makers. <i>Journal of Travel Medicine</i> , 2018, 25, .	3.0	31
81	Reconstruction of the epidemic history of hepatitis B virus genotype D in Albania. <i>Infection, Genetics and Evolution</i> , 2012, 12, 291-298.	2.3	30
82	Phylogeography and phylodynamics of European genotype 3 hepatitis E virus. <i>Infection, Genetics and Evolution</i> , 2014, 25, 138-143.	2.3	30
83	Two Distinct Hepatitis C Virus Genotype 1a Clades Have Different Geographical Distribution and Association With Natural Resistance to NS3 Protease Inhibitors. <i>Open Forum Infectious Diseases</i> , 2015, 2, ofv043.	0.9	30
84	Hepatitis C virus genotype 4d in Southern Italy: Reconstruction of its origin and spread by a phyldynamic analysis. <i>Journal of Medical Virology</i> , 2012, 84, 1613-1619.	5.0	29
85	Phylogenetic analysis provides evidence of interactions between Italian heterosexual and South American homosexual males as the main source of national HIV subtype C epidemics. <i>Journal of Medical Virology</i> , 2014, 86, 729-736.	5.0	29
86	Inverse correlation between average monthly high temperatures and COVID-19-related death rates in different geographical areas. <i>Journal of Translational Medicine</i> , 2020, 18, 251.	4.4	29
87	High value of mid-regional proadrenomedullin in COVID-19: A marker of widespread endothelial damage, disease severity, and mortality. <i>Journal of Medical Virology</i> , 2021, 93, 2820-2827.	5.0	29
88	Chikungunya and Dengue Viruses in Travelers. <i>Emerging Infectious Diseases</i> , 2008, 14, 177-178.	4.3	28
89	Molecular epidemiology and phylogeny of Nipah virus infection: A mini review. <i>Asian Pacific Journal of Tropical Medicine</i> , 2016, 9, 630-634.	0.8	28
90	Performances of five risk algorithms in predicting cardiovascular events in patients with Psoriatic Arthritis: An Italian bicentric study. <i>PLoS ONE</i> , 2018, 13, e0205506.	2.5	28

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91	Procalcitonin and MR-proAdrenomedullin combination in the etiological diagnosis and prognosis of sepsis and septic shock. <i>Microbial Pathogenesis</i> , 2019, 137, 103763.	2.9	28
92	Evidence for mutations in SARS-CoV-2 Italian isolates potentially affecting virus transmission. <i>Journal of Medical Virology</i> , 2020, 92, 2232-2237.	5.0	28
93	Molecular Epidemiology of HIV-1 in African Countries: A Comprehensive Overview. <i>Pathogens</i> , 2020, 9, 1072.	2.8	28
94	Epidemiological network analysis in HIV-1 B infected patients diagnosed in Italy between 2000 and 2008. <i>Infection, Genetics and Evolution</i> , 2011, 11, 624-632.	2.3	27
95	Spatial and temporal reconstruction of bovine viral diarrhoea virus genotype 1 dispersion in Italy. <i>Infection, Genetics and Evolution</i> , 2012, 12, 324-331.	2.3	27
96	Impact of spatial dispersion, evolution and selection on Ebola Zaire Virus epidemic waves. <i>Scientific Reports</i> , 2015, 5, 10170.	3.3	27
97	The role of procalcitonin in the diagnosis of bacterial infection after major abdominal surgery. <i>Medicine (United States)</i> , 2018, 97, e9496.	1.0	27
98	Role of the Specialized Proresolving Mediator Resolvin D1 in Systemic Lupus Erythematosus: Preliminary Results. <i>Journal of Immunology Research</i> , 2018, 2018, 1-6.	2.2	27
99	The variants question: What is the problem?. <i>Journal of Medical Virology</i> , 2021, 93, 6479-6485.	5.0	26
100	Covid-19 sequelae in working age patients: A systematic review. <i>Journal of Medical Virology</i> , 2022, 94, 858-868.	5.0	26
101	Mechanisms that Trigger a Good Health-Care Response to Intimate Partner Violence in Spain. Combining Realist Evaluation and Qualitative Comparative Analysis Approaches. <i>PLoS ONE</i> , 2015, 10, e0135167.	2.5	25
102	HIV-1 A1 Subtype Epidemic in Italy Originated from Africa and Eastern Europe and Shows a High Frequency of Transmission Chains Involving Intravenous Drug Users. <i>PLoS ONE</i> , 2016, 11, e0146097.	2.5	25
103	Screening of HIV-1 Isolates by Reverse Heteroduplex Mobility Assay and Identification of Non-B Subtypes in Italy. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2004, 37, 1295-1306.	2.1	24
104	Molecular and phylogenetic analysis of HIV-1 variants circulating among injecting drug users in Mashhad-Iran. <i>Infectious Agents and Cancer</i> , 2006, 1, 4.	2.6	24
105	Performance evaluation of the COBAS/TaqMan HIV-1 v2.0 in HIV-1 positive patients with low viral load: A comparative study. <i>Journal of Virological Methods</i> , 2011, 173, 399-402.	2.1	24
106	Reliable timescale inference of HBV genotype A origin and phylodynamics. <i>Infection, Genetics and Evolution</i> , 2015, 32, 361-369.	2.3	24
107	Zika virus complete genome from Salvador, Bahia, Brazil. <i>Infection, Genetics and Evolution</i> , 2016, 41, 142-145.	2.3	24
108	Best diagnostic accuracy of sepsis combining SIRS criteria or qSOFA score with Procalcitonin and Mid-Regional pro-Adrenomedullin outside ICU. <i>Scientific Reports</i> , 2020, 10, 16605.	3.3	24

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109	Risk of hepatitis A infection following travel. <i>Journal of Viral Hepatitis</i> , 2002, 9, 460-465.	2.0	23
110	H5N1 influenza virus evolution: a comparison of different epidemics in birds and humans (1997-2004). <i>Journal of General Virology</i> , 2006, 87, 955-960.	2.9	23
111	Characterization of the patterns of drug-resistance mutations in newly diagnosed HIV-1 infected patients naïve to the antiretroviral drugs. <i>BMC Infectious Diseases</i> , 2009, 9, 111.	2.9	23
112	Molecular Epidemiology of HIV Type 1 CRF02_AG in Cameroon and African Patients Living in Italy. <i>AIDS Research and Human Retroviruses</i> , 2011, 27, 1173-1182.	1.1	23
113	Treatment-associated polymorphisms in protease are significantly associated with higher viral load and lower CD4 count in newly diagnosed drug-naïve HIV-1 infected patients. <i>Retrovirology</i> , 2012, 9, 81.	2.0	23
114	Critical review of sham surgery clinical trials: Confounding factors analysis. <i>Annals of Medicine and Surgery</i> , 2016, 12, 21-26.	1.1	23
115	MALDI-TOF MS Identification and Clustering Applied to Enterobacter Species in Nosocomial Setting. <i>Frontiers in Microbiology</i> , 2018, 9, 1885.	3.5	23
116	The impact of the SARS-CoV-2 infection, with special reference to the hematological setting. <i>Journal of Medical Virology</i> , 2021, 93, 223-233.	5.0	23
117	SARS-CoV-2 shifting transmission dynamics and hidden reservoirs potentially limit efficacy of public health interventions in Italy. <i>Communications Biology</i> , 2021, 4, 489.	4.4	23
118	HIV-1 Subtype F1 Epidemiological Networks among Italian Heterosexual Males Are Associated with Introduction Events from South America. <i>PLoS ONE</i> , 2012, 7, e42223.	2.5	22
119	Reconstruction of the evolutionary dynamics of hepatitis C virus subtypes in Montenegro and the Balkan region. <i>Infection, Genetics and Evolution</i> , 2013, 17, 223-230.	2.3	22
120	Multi-drug resistant <i>Klebsiella pneumoniae</i> strains circulating in hospital setting: whole-genome sequencing and Bayesian phylogenetic analysis for outbreak investigations. <i>Scientific Reports</i> , 2017, 7, 3534.	3.3	22
121	Hepatitis E virus genotypes and subgenotypes causing acute hepatitis, Bulgaria, 2013-2015. <i>PLoS ONE</i> , 2018, 13, e0198045.	2.5	22
122	Phylogenetic analysis of porcine circovirus type 2 in Sardinia, Italy, shows genotype 2d circulation among domestic pigs and wild boars. <i>Infection, Genetics and Evolution</i> , 2019, 71, 189-196.	2.3	22
123	Preservation of microvascular barrier function requires CD31 receptor-induced metabolic reprogramming. <i>Nature Communications</i> , 2020, 11, 3595.	12.8	22
124	Targeting Microbiome: An Alternative Strategy for Fighting SARS-CoV-2 Infection. <i>Chemotherapy</i> , 2021, 66, 24-32.	1.6	22
125	An outbreak of Q fever in Bulgaria. <i>Annali Dell'Istituto Superiore Di Sanita</i> , 2009, 45, 83-6.	0.4	22
126	An outbreak of HIV-1 subtype G among Italian injecting drug users. <i>Aids</i> , 2007, 21, 1213-1215.	2.2	21

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127	Genetic diversity of human papillomavirus type 16 E6, E7, and L1 genes in Italian women with different grades of cervical lesions. <i>Journal of Medical Virology</i> , 2009, 81, 1627-1634.	5.0	21
128	Back to the origin of HCV 2c subtype and spreading to the Calabria region (Southern Italy) over the last two centuries: A phylogenetic study. <i>Infection, Genetics and Evolution</i> , 2014, 26, 352-358.	2.3	21
129	Pro-inflammatory adipokine profile in psoriatic arthritis: results from a cross-sectional study comparing PsA subset with evident cutaneous involvement and subset "seronegative psoriasis". <i>Clinical Rheumatology</i> , 2019, 38, 2547-2552.	2.2	21
130	Influenza B virus infection complicated by life-threatening pericarditis: a unique case-report and literature review. <i>BMC Infectious Diseases</i> , 2019, 19, 40.	2.9	21
131	Cardiovascular Risk Prediction in Ankylosing Spondylitis: From Traditional Scores to Machine Learning Assessment. <i>Rheumatology and Therapy</i> , 2020, 7, 867-882.	2.3	21
132	A close relationship between HIF-1 $\alpha$ expression and bone metastases in advanced NSCLC, a retrospective analysis. <i>Oncotarget</i> , 2019, 10, 7071-7079.	1.8	21
133	Lymphomatoid papulosis. <i>Minerva Medica</i> , 2020, 111, 166-172.	0.9	21
134	Competition for dominance within replicating quasispecies during prolonged SARS-CoV-2 infection in an immunocompromised host. <i>Virus Evolution</i> , 2022, 8, .	4.9	21
135	Contamination of mussels by hepatitis A virus: a public-health problem in southern Italy. <i>Food Control</i> , 2003, 14, 559-563.	5.5	20
136	Detection of Toscana virus central nervous system infections in Sardinia Island, Italy. <i>Journal of Clinical Virology</i> , 2007, 40, 90-91.	3.1	20
137	Identification of the novel KI polyomavirus in the respiratory tract of an Italian patient. <i>Journal of Medical Virology</i> , 2008, 80, 2012-2014.	5.0	20
138	Molecular analysis of hepatitis C virus infection in Bulgarian injecting drug users. <i>Journal of Medical Virology</i> , 2011, 83, 1565-1570.	5.0	20
139	Molecular epidemiology and genetic diversity of human rhinovirus affecting hospitalized children in Rome. <i>Medical Microbiology and Immunology</i> , 2013, 202, 303-311.	4.8	20
140	Evolutionary dynamics of HBV $\alpha$ 1 genotype epidemic in Turkey. <i>Journal of Medical Virology</i> , 2014, 86, 109-116.	5.0	20
141	Molecular epidemiology of hepatitis B virus genotypes circulating in acute hepatitis B patients in the Campania region. <i>Journal of Medical Virology</i> , 2014, 86, 1683-1693.	5.0	20
142	Genetic diversity of the haemagglutinin (HA) of human influenza A (H1N1) virus in Montenegro: Focus on its origin and evolution. <i>Journal of Medical Virology</i> , 2016, 88, 1905-1913.	5.0	20
143	Bayesian Phylogeography of Crimean-Congo Hemorrhagic Fever Virus in Europe. <i>PLoS ONE</i> , 2013, 8, e79663.	2.5	20
144	T-cell-mediated protective efficacy of a systemic vaccine approach in cynomolgus monkeys after SIV mucosal challenge. <i>Journal of Medical Primatology</i> , 2004, 33, 251-261.	0.6	19

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145	Phylogeny of Dengue and Chikungunya viruses in Al Hudayda governorate, Yemen. <i>Infection, Genetics and Evolution</i> , 2014, 27, 395-401.	2.3	19
146	Multi-drug resistant <i>Pseudomonas aeruginosa</i> nosocomial strains: Molecular epidemiology and evolution. <i>Microbial Pathogenesis</i> , 2018, 123, 233-241.	2.9	19
147	Within-Host Dynamics of the Hepatitis C Virus Quasispecies Population in HIV-1/HCV Coinfected Patients. <i>PLoS ONE</i> , 2011, 6, e16551.	2.5	19
148	The unresolved question on COVID-19 virus origin: The three cards game?. <i>Journal of Medical Virology</i> , 2022, 94, 1257-1260.	5.0	19
149	Non-B HIV Type 1 Subtypes among Men Who Have Sex with Men in Rome, Italy. <i>AIDS Research and Human Retroviruses</i> , 2009, 25, 157-164.	1.1	18
150	HIV-1 subtype C transmission network: The phylogenetic reconstruction strongly supports the epidemiological data. <i>Journal of Clinical Virology</i> , 2010, 48, 212-214.	3.1	18
151	Intratype variations of HPV 31 and 58 in Italian women with abnormal cervical cytology. <i>Journal of Medical Virology</i> , 2011, 83, 1752-1761.	5.0	18
152	Reconstruction of the Evolutionary Dynamics of the A(H1N1)pdm09 Influenza Virus in Italy during the Pandemic and Post-Pandemic Phases. <i>PLoS ONE</i> , 2012, 7, e47517.	2.5	18
153	Procalcitonin and MR-proAdrenomedullin combined score in the diagnosis and prognosis of systemic and localized bacterial infections. <i>Journal of Infection</i> , 2016, 72, 395-398.	3.3	18
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