

JÃ¼rg BÃ¶ni

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1401450/publications.pdf>

Version: 2024-02-01

123
papers

3,038
citations

186265

28
h-index

197818

49
g-index

135
all docs

135
docs citations

135
times ranked

4121
citing authors

#	ARTICLE	IF	CITATIONS
1	Secondary attack rates from asymptomatic and symptomatic influenza virus shedders in hospitals: Results from the TransFLUas influenza transmission study. <i>Infection Control and Hospital Epidemiology</i> , 2022, 43, 312-318.	1.8	9
2	Increasing Frequency and Transmission of HIV-1 Non-B Subtypes Among Men Who Have Sex With Men in the Swiss HIV Cohort Study. <i>Journal of Infectious Diseases</i> , 2022, 225, 306-316.	4.0	5
3	Identifying and Characterizing Trans Women in the Swiss HIV Cohort Study as an Epidemiologically Distinct Risk Group. <i>Clinical Infectious Diseases</i> , 2022, 74, 1468-1475.	5.8	3
4	Decreasing Incidence and Determinants of Bacterial Pneumonia in People With HIV: The Swiss HIV Cohort Study. <i>Journal of Infectious Diseases</i> , 2022, 225, 1592-1600.	4.0	4
5	Integrase strand transfer inhibitor use and cancer incidence in a large cohort setting. <i>Open Forum Infectious Diseases</i> , 2022, 9, ofac029.	0.9	3
6	Impact of Latent Tuberculosis on Diabetes. <i>Journal of Infectious Diseases</i> , 2022, 225, 2229-2234.	4.0	3
7	Detecting Selection in the HIV-1 Genome during Sexual Transmission Events. <i>Viruses</i> , 2022, 14, 406.	3.3	1
8	Antibody Response in Immunocompromised Patients After the Administration of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Vaccine BNT162b2 or mRNA-1273: A Randomized Controlled Trial. <i>Clinical Infectious Diseases</i> , 2022, 75, e585-e593.	5.8	26
9	Similar but different: Integrated phylogenetic analysis of Austrian and Swiss HIV-1 sequences reveal differences in transmission patterns of the local HIV-1 epidemics. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2022, Publish Ahead of Print, .	2.1	0
10	The Interplay Between Replication Capacity of HIV-1 and Surrogate Markers of Disease. <i>Journal of Infectious Diseases</i> , 2022, 226, 1057-1068.	4.0	2
11	An Approach to Quantifying the Interaction between Behavioral and Transmission Clusters. <i>Viruses</i> , 2022, 14, 784.	3.3	2
12	Sustained Effect on Hepatitis C Elimination Among Men Who Have Sex With Men in the Swiss HIV Cohort Study: A Systematic Re-Screening for Hepatitis C RNA Two Years Following a Nation-Wide Elimination Program. <i>Clinical Infectious Diseases</i> , 2022, 75, 1723-1731.	5.8	14
13	A systematic molecular epidemiology screen reveals numerous HIV-1 superinfections in the Swiss HIV Cohort Study. <i>Journal of Infectious Diseases</i> , 2022, , .	4.0	3
14	Antibodies from convalescent plasma promote SARS-CoV-2 clearance in individuals with and without endogenous antibody response. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	26
15	A Treatment-as-Prevention Trial to Eliminate Hepatitis C Among Men Who Have Sex With Men Living With Human Immunodeficiency Virus (HIV) in the Swiss HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2021, 73, e2194-e2202.	5.8	47
16	The Impact of Surgical Strategy and Rifampin on Treatment Outcome in <i>Cutibacterium</i> Periprosthetic Joint Infections. <i>Clinical Infectious Diseases</i> , 2021, 72, e1064-e1073.	5.8	22
17	Phylogenetic Cluster Analysis Identifies Virological and Behavioral Drivers of Human Immunodeficiency Virus Transmission in Men Who Have Sex With Men. <i>Clinical Infectious Diseases</i> , 2021, 72, 2175-2183.	5.8	10
18	The Role of Human Immunodeficiency Virus (HIV) Asymptomatic Status When Starting Antiretroviral Therapy on Adherence and Treatment Outcomes and Implications for Test and Treat: The Swiss HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2021, 72, 1413-1421.	5.8	2

#	ARTICLE	IF	CITATIONS
19	HIV-1 integration sites in CD4+ T-cells during primary, chronic, and late presentation of HIV-1 infection. <i>JCI Insight</i> , 2021, 6, .	5.0	7
20	High Efficacy of Saliva in Detecting SARS-CoV-2 by RT-PCR in Adults and Children. <i>Microorganisms</i> , 2021, 9, 642.	3.6	41
21	Systematic screening of viral and human genetic variation identifies antiretroviral resistance and immune escape link. <i>ELife</i> , 2021, 10, .	6.0	3
22	Differences Between Infectious Disease Events in First Liver Transplant Versus Replantation in the Swiss Transplant Cohort Study. <i>Liver Transplantation</i> , 2021, 27, 1283-1290.	2.4	3
23	Reduced Relative Sensitivity of the Elecsys SARS-CoV-2 Antigen Assay in Saliva Compared to Nasopharyngeal Swabs. <i>Microorganisms</i> , 2021, 9, 1700.	3.6	11
24	Usefulness of the GenMark ePlex RPP assay for the detection of respiratory viruses compared to the FTD21 multiplex RT-PCR. <i>Diagnostic Microbiology and Infectious Disease</i> , 2021, 101, 115424.	1.8	0
25	Absenteeism and presenteeism in healthcare workers due to respiratory illness. <i>Infection Control and Hospital Epidemiology</i> , 2021, 42, 268-273.	1.8	14
26	A trial platform to assess approved SARS-CoV-2 vaccines in immunocompromised patients: first sub-protocol for a pilot trial comparing the mRNA vaccines Comirnaty® and COVID-19 mRNA Vaccine Moderna®. <i>Trials</i> , 2021, 22, 724.	1.6	9
27	Management of Suspected Cases of Feline Immunodeficiency Virus Infection in Eurasian Lynx (<i>Lynx</i>) Tj ETQq1 1 0.784314 rgBT /Over	2.2	0
28	Assessing the drivers of syphilis among men who have sex with men in Switzerland reveals a key impact of screening frequency: A modelling study. <i>PLoS Computational Biology</i> , 2021, 17, e1009529.	3.2	6
29	Determinants for voluntary participation in staff screening during an methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) outbreak on a neonatal ward. <i>Infection Control and Hospital Epidemiology</i> , 2021, 42, 881-884.	1.8	1
30	Multifactorial seroprofiling dissects the contribution of pre-existing human coronaviruses responses to SARS-CoV-2 immunity. <i>Nature Communications</i> , 2021, 12, 6703.	12.8	36
31	Impact of an electronic alert on prescription patterns of meropenem, voriconazole and caspofungin. <i>BMC Infectious Diseases</i> , 2021, 21, 1263.	2.9	0
32	Emergence of Drug Resistance in the Swiss HIV Cohort Study Under Potent Antiretroviral Therapy Is Observed in Socially Disadvantaged Patients. <i>Clinical Infectious Diseases</i> , 2020, 70, 297-303.	5.8	10
33	Self-reported Neurocognitive Impairment in People Living With Human Immunodeficiency Virus (HIV): Characterizing Clusters of Patients With Similar Changes in Self-reported Neurocognitive Impairment, 2013â€“2017, in the Swiss HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2020, 71, 637-644.	5.8	3
34	Vitamin D deficiency is common in kidney transplant recipients, but is not associated with infections after transplantation. <i>Clinical Transplantation</i> , 2020, 34, e13778.	1.6	1
35	Evaluation of the RIDA®GENE RT-PCR assays for detection of sapovirus, astrovirus, adenovirus, and rotavirus in stool samples of adults in Switzerland. <i>Diagnostic Microbiology and Infectious Disease</i> , 2020, 96, 114924.	1.8	5
36	Emergence of Resistance to Integrase Strand Transfer Inhibitors during Dolutegravir Containing Triple-Therapy in a Treatment-Experienced Patient with Pre-Existing M184V/I Mutation. <i>Viruses</i> , 2020, 12, 1330.	3.3	9

#	ARTICLE	IF	CITATIONS
37	Host Genomics of the HIV-1 Reservoir Size and Its Decay Rate During Suppressive Antiretroviral Treatment. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2020, 85, 517-524.	2.1	7
38	Implementation and evaluation of a care bundle for prevention of non-ventilator-associated hospital-acquired pneumonia (nvHAP) – a mixed-methods study protocol for a hybrid type 2 effectiveness-implementation trial. <i>BMC Infectious Diseases</i> , 2020, 20, 603.	2.9	6
39	Does respiratory co-infection facilitate dispersal of SARS-CoV-2? investigation of a super-spreading event in an open-space office. <i>Antimicrobial Resistance and Infection Control</i> , 2020, 9, 191.	4.1	19
40	HCV Genetic Diversity Can Be Used to Infer Infection Recency and Time since Infection. <i>Viruses</i> , 2020, 12, 1241.	3.3	3
41	Heritability of the HIV-1 reservoir size and decay under long-term suppressive ART. <i>Nature Communications</i> , 2020, 11, 5542.	12.8	5
42	Diagnosis of latent tuberculosis infection is associated with reduced HIV viral load and lower risk for opportunistic infections in people living with HIV. <i>PLoS Biology</i> , 2020, 18, e3000963.	5.6	6
43	Impact of scaling up dolutegravir on antiretroviral resistance in South Africa: A modeling study. <i>PLoS Medicine</i> , 2020, 17, e1003397.	8.4	7
44	Differences in Social and Mental Well-Being of Long-Term Survivors among People who Inject Drugs and Other Participants in the Swiss HIV Cohort Study: 1980–2018. <i>Antiviral Therapy</i> , 2020, 25, 43-54.	1.0	2
45	Title is missing!. , 2020, 18, e3000963.		0
46	Title is missing!. , 2020, 18, e3000963.		0
47	Title is missing!. , 2020, 18, e3000963.		0
48	Title is missing!. , 2020, 18, e3000963.		0
49	Title is missing!. , 2020, 18, e3000963.		0
50	Title is missing!. , 2020, 18, e3000963.		0
51	Impact of scaling up dolutegravir on antiretroviral resistance in South Africa: A modeling study. , 2020, 17, e1003397.		0
52	Impact of scaling up dolutegravir on antiretroviral resistance in South Africa: A modeling study. , 2020, 17, e1003397.		0
53	Impact of scaling up dolutegravir on antiretroviral resistance in South Africa: A modeling study. , 2020, 17, e1003397.		0
54	Impact of scaling up dolutegravir on antiretroviral resistance in South Africa: A modeling study. , 2020, 17, e1003397.		0

#	ARTICLE	IF	CITATIONS
55	High Cure Rates With Grazoprevir-Elbasvir With or Without Ribavirin Guided by Genotypic Resistance Testing Among Human Immunodeficiency Virus/Hepatitis C Virus“coinfected Men Who Have Sex With Men. <i>Clinical Infectious Diseases</i> , 2019, 68, 569-576.	5.8	30
56	Determinants of HIV-1 reservoir size and long-term dynamics during suppressive ART. <i>Nature Communications</i> , 2019, 10, 3193.	12.8	112
57	Bridging the gap between HIV epidemiology and antiretroviral resistance evolution: Modelling the spread of resistance in South Africa. <i>PLoS Computational Biology</i> , 2019, 15, e1007083.	3.2	11
58	The TransFLUas influenza transmission study in acute healthcare - recruitment rates and protocol adherence in healthcare workers and inpatients. <i>BMC Infectious Diseases</i> , 2019, 19, 446.	2.9	6
59	Widespread B cell perturbations in HIV-1 infection afflict naive and marginal zone B cells. <i>Journal of Experimental Medicine</i> , 2019, 216, 2071-2090.	8.5	22
60	A Systematic Phylogenetic Approach to Study the Interaction of HIV-1 With Coinfections, Noncommunicable Diseases, and Opportunistic Diseases. <i>Journal of Infectious Diseases</i> , 2019, 220, 244-253.	4.0	6
61	Viral Diversity Based on Next-Generation Sequencing of HIV-1 Provides Precise Estimates of Infection Recency and Time Since Infection. <i>Journal of Infectious Diseases</i> , 2019, 220, 254-265.	4.0	27
62	Clusters of Sexual Behavior in Human Immunodeficiency Virus“positive Men Who Have Sex With Men Reveal Highly Dissimilar Time Trends. <i>Clinical Infectious Diseases</i> , 2019, 70, 416-424.	5.8	9
63	Changing Trends in International Versus Domestic HCV Transmission in HIV-Positive Men Who Have Sex With Men: A Perspective for the Direct-Acting Antiviral Scale-Up Era. <i>Journal of Infectious Diseases</i> , 2019, 220, 91-99.	4.0	24
64	HIV Transmission Chains Exhibit Greater HLA-B Homogeneity Than Randomly Expected. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2019, 81, 508-515.	2.1	0
65	Noninferiority of Simplified Dolutegravir Monotherapy Compared to Continued Combination Antiretroviral Therapy That Was Initiated During Primary Human Immunodeficiency Virus Infection: A Randomized, Controlled, Multisite, Open-label, Noninferiority Trial. <i>Clinical Infectious Diseases</i> , 2019, 69, 1489-1497.	5.8	19
66	Importance of routine viral load monitoring: higher levels of resistance at ART failure in Uganda and Lesotho compared with Switzerland. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 468-472.	3.0	9
67	Metagenomic Virome Sequencing in Living Donor and Recipient Kidney Transplant Pairs Revealed JC Polyomavirus Transmission. <i>Clinical Infectious Diseases</i> , 2019, 69, 987-994.	5.8	13
68	OUP accepted manuscript. <i>Clinical Infectious Diseases</i> , 2019, 68, 561-568.	5.8	13
69	The rate of mother-to-child transmission of antiretroviral drug-resistant HIV strains is low in the Swiss Mother and Child HIV Cohort Study. <i>Swiss Medical Weekly</i> , 2019, 149, w20059.	1.6	4
70	Inferring the age difference in HIV transmission pairs by applying phylogenetic methods on the HIV transmission network of the Swiss HIV Cohort Study. <i>Virus Evolution</i> , 2018, 4, vey024.	4.9	17
71	Dissecting HIV Virulence: Heritability of Setpoint Viral Load, CD4+ T-Cell Decline, and Per-Parasite Pathogenicity. <i>Molecular Biology and Evolution</i> , 2018, 35, 27-37.	8.9	37
72	Tracing HIV-1 strains that imprint broadly neutralizing antibody responses. <i>Nature</i> , 2018, 561, 406-410.	27.8	47

#	ARTICLE	IF	CITATIONS
73	The Cumulative Impact of Harm Reduction on the Swiss HIV Epidemic: Cohort Study, Mathematical Model, and Phylogenetic Analysis. <i>Open Forum Infectious Diseases</i> , 2018, 5, ofy078.	0.9	8
74	Distinct, IgG1-driven antibody response landscapes demarcate individuals with broadly HIV-1 neutralizing activity. <i>Journal of Experimental Medicine</i> , 2018, 215, 1589-1608.	8.5	29
75	Persistent mammalian orthoreovirus, coxsackievirus and adenovirus co-infection in a child with a primary immunodeficiency detected by metagenomic sequencing: a case report. <i>BMC Infectious Diseases</i> , 2018, 18, 33.	2.9	16
76	Quantifying the fitness cost of HIV-1 drug resistance mutations through phylodynamics. <i>PLoS Pathogens</i> , 2018, 14, e1006895.	4.7	53
77	Influence of time to diagnosis of severe influenza on antibiotic use, length of stay, isolation precautions, and mortality: a retrospective study. <i>Influenza and Other Respiratory Viruses</i> , 2017, 11, 337-344.	3.4	19
78	MinVar: A rapid and versatile tool for HIV-1 drug resistance genotyping by deep sequencing. <i>Journal of Virological Methods</i> , 2017, 240, 7-13.	2.1	49
79	Parent-offspring regression to estimate the heritability of an HIV-1 trait in a realistic setup. <i>Retrovirology</i> , 2017, 14, 33.	2.0	16
80	Therapeutic Immune Recovery and Reduction of CXCR4-Tropic HIV-1. <i>Clinical Infectious Diseases</i> , 2017, 64, 295-300.	5.8	14
81	CD4 cell count response to first-line combination ART in HIV-2+ patients compared with HIV-1+ patients: a multinational, multicohort European study. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 2869-2878.	3.0	17
82	Assessing the danger of self-sustained HIV epidemics in heterosexuals by population based phylogenetic cluster analysis. <i>ELife</i> , 2017, 6, .	6.0	16
83	Optimization and validation of sample preparation for metagenomic sequencing of viruses in clinical samples. <i>Microbiome</i> , 2017, 5, 94.	11.1	59
84	Metagenomic sequencing complements routine diagnostics in identifying viral pathogens in lung transplant recipients with unknown etiology of respiratory infection. <i>PLoS ONE</i> , 2017, 12, e0177340.	2.5	56
85	Importance of an Early HIV Antibody Differentiation Immunoassay for Detection of Dual Infection with HIV-1 and HIV-2. <i>PLoS ONE</i> , 2016, 11, e0157690.	2.5	4
86	Genotypic Resistance Tests Sequences Reveal the Role of Marginalized Populations in HIV-1 Transmission in Switzerland. <i>Scientific Reports</i> , 2016, 6, 27580.	3.3	15
87	Successful Prevention of Transmission of Integrase Resistance in the Swiss HIV Cohort Study. <i>Journal of Infectious Diseases</i> , 2016, 214, 399-402.	4.0	47
88	Determinants of HIV-1 broadly neutralizing antibody induction. <i>Nature Medicine</i> , 2016, 22, 1260-1267.	30.7	133
89	Large-scale inference of conjunctive Bayesian networks. <i>Bioinformatics</i> , 2016, 32, i727-i735.	4.1	21
90	A Direct Comparison of Two Densely Sampled HIV Epidemics: The UK and Switzerland. <i>Scientific Reports</i> , 2016, 6, 32251.	3.3	17

#	ARTICLE	IF	CITATIONS
91	Tracing HIV-1 transmission: envelope traits of HIV-1 transmitter and recipient pairs. <i>Retrovirology</i> , 2016, 13, 62.	2.0	45
92	Emergence of Acquired HIV-1 Drug Resistance Almost Stopped in Switzerland: A 15-Year Prospective Cohort Analysis. <i>Clinical Infectious Diseases</i> , 2016, 62, 1310-1317.	5.8	52
93	HIV-1 Transmission During Recent Infection and During Treatment Interruptions as Major Drivers of New Infections in the Swiss HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2016, 62, 115-122.	5.8	60
94	Resolution of plasma sample mix-ups through comparison of patient antibody patterns to <i>E. coli</i> . <i>Journal of Immunological Methods</i> , 2015, 427, 130-133.	1.4	0
95	A Lead-In with Silibinin Prior to Triple-Therapy Translates into Favorable Treatment Outcomes in Difficult-To-Treat HIV/Hepatitis C Coinfected Patients. <i>PLoS ONE</i> , 2015, 10, e0133028.	2.5	18
96	Gammaretrovirus-Specific Antibodies in Free-Ranging and Captive Namibian Cheetahs. <i>Vaccine Journal</i> , 2015, 22, 611-617.	3.1	5
97	Assessing the Paradox Between Transmitted and Acquired HIV Type 1 Drug Resistance Mutations in the Swiss HIV Cohort Study From 1998 to 2012. <i>Journal of Infectious Diseases</i> , 2015, 212, 28-38.	4.0	61
98	Persistence of Transmitted HIV-1 Drug Resistance Mutations Associated with Fitness Costs and Viral Genetic Backgrounds. <i>PLoS Pathogens</i> , 2015, 11, e1004722.	4.7	68
99	Unbiased metagenomic sequencing complements specific routine diagnostic methods and increases chances to detect rare viral strains. <i>Diagnostic Microbiology and Infectious Disease</i> , 2015, 83, 133-138.	1.8	29
100	Assessing efficacy of different nucleos(t)ide backbones in NNRTI-containing regimens in the Swiss HIV Cohort Study. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, dkv257.	3.0	6
101	Partial rescue of V1V2 mutant infectivity by HIV-1 cell-cell transmission supports the domain's exceptional capacity for sequence variation. <i>Retrovirology</i> , 2014, 11, 75.	2.0	16
102	Comparative Performances of HIV-1 RNA Load Assays at Low Viral Load Levels: Results of an International Collaboration. <i>Journal of Clinical Microbiology</i> , 2014, 52, 517-523.	3.9	47
103	Use of reverse-transcriptase-based HIV-1 viral load assessment to confirm low viral loads in newly diagnosed patients in Switzerland. <i>BMC Infectious Diseases</i> , 2014, 14, 84.	2.9	2
104	Limited clinical benefit of minority K103N and Y181C-variant detection in addition to routine genotypic resistance testing in antiretroviral therapy-naïve patients. <i>Aids</i> , 2014, 28, 2231-2239.	2.2	20
105	Treatment-Naïve Individuals Are the Major Source of Transmitted HIV-1 Drug Resistance in Men Who Have Sex With Men in the Swiss HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2014, 58, 285-294.	5.8	75
106	Clustering of HCV coinfections on HIV phylogeny indicates domestic and sexual transmission of HCV. <i>International Journal of Epidemiology</i> , 2014, 43, 887-896.	1.9	36
107	Higher Risk of Incident Hepatitis C Virus Coinfection Among Men Who Have Sex With Men, in Whom the HIV Genetic Bottleneck at Transmission Was Wide. <i>Journal of Infectious Diseases</i> , 2014, 210, 1555-1561.	4.0	16
108	Social Meets Molecular: Combining Phylogenetic and Latent Class Analyses to Understand HIV-1 Transmission in Switzerland. <i>American Journal of Epidemiology</i> , 2014, 179, 1514-1525.	3.4	25

#	ARTICLE	IF	CITATIONS
109	Generation of a Recombinant Gag Virus-Like-Particle Panel for the Evaluation of p24 Antigen Detection by Diagnostic HIV Tests. PLoS ONE, 2014, 9, e111552.	2.5	11
110	The Individualized Genetic Barrier Predicts Treatment Response in a Large Cohort of HIV-1 Infected Patients. PLoS Computational Biology, 2013, 9, e1003203.	3.2	19
111	Origin of Minority Drug-Resistant HIV-1 Variants in Primary HIV-1 Infection. Journal of Infectious Diseases, 2013, 208, 1102-1112.	4.0	35
112	Adherence as a Predictor of the Development of Class-Specific Resistance Mutations: The Swiss HIV Cohort Study. PLoS ONE, 2013, 8, e77691.	2.5	49
113	Estimating the Basic Reproductive Number from Viral Sequence Data. Molecular Biology and Evolution, 2012, 29, 347-357.	8.9	206
114	Minor Protease Inhibitor Mutations at Baseline Do Not Increase the Risk for a Virological Failure in HIV-1 Subtype B Infected Patients. PLoS ONE, 2012, 7, e37983.	2.5	15
115	Long-Lasting Protection of Activity of Nucleoside Reverse Transcriptase Inhibitors and Protease Inhibitors (PIs) by Boosted PI Containing Regimens. PLoS ONE, 2012, 7, e50307.	2.5	16
116	Comparison of HIV-1 viral load based on RNA or reverse transcriptase activity in patients with suspected viral load underestimation. Retrovirology, 2012, 9, .	2.0	0
117	Characterization of Human Immunodeficiency Virus Type 1 (HIV-1) Diversity and Tropism in 145 Patients With Primary HIV-1 Infection. Clinical Infectious Diseases, 2011, 53, 1271-1279.	5.8	84
118	Ambiguous Nucleotide Calls From Population-based Sequencing of HIV-1 are a Marker for Viral Diversity and the Age of Infection. Clinical Infectious Diseases, 2011, 52, 532-539.	5.8	127
119	HIV-1 transmission after cessation of early antiretroviral therapy among men having sex with men. Aids, 2010, 24, 1177-1183.	2.2	62
120	Molecular Epidemiology Reveals Long-Term Changes in HIV Type 1 Subtype B Transmission in Switzerland. Journal of Infectious Diseases, 2010, 201, 1488-1497.	4.0	172
121	Phylogenetic Approach Reveals That Virus Genotype Largely Determines HIV Set-Point Viral Load. PLoS Pathogens, 2010, 6, e1001123.	4.7	108
122	African descent is associated with slower CD4 cell count decline in treatment-naive patients of the Swiss HIV Cohort Study. Aids, 2009, 23, 1269-1276.	2.2	28
123	Transmission of HIV-1 drug resistance in Switzerland: a 10-year molecular epidemiology survey. Aids, 2007, 21, 2223-2229.	2.2	117