

Ahidjo Ayouba

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4828295/publications.pdf>

Version: 2024-02-01

124
papers

5,265
citations

76326

40
h-index

106344

65
g-index

132
all docs

132
docs citations

132
times ranked

6031
citing authors

#	ARTICLE	IF	CITATIONS
1	Tetherin-Driven Adaptation of Vpu and Nef Function and the Evolution of Pandemic and Nonpandemic HIV-1 Strains. <i>Cell Host and Microbe</i> , 2009, 6, 409-421.	11.0	391
2	Self-reactive antibodies (natural autoantibodies) in healthy individuals. <i>Journal of Immunological Methods</i> , 1998, 216, 117-137.	1.4	299
3	SIVcpz in Wild Chimpanzees. <i>Science</i> , 2002, 295, 465-465.	12.6	207
4	African origin of the malaria parasite <i>Plasmodium vivax</i> . <i>Nature Communications</i> , 2014, 5, 3346.	12.8	167
5	Human herpesvirus 8 primary infection occurs during childhood in Cameroon, Central Africa. , 1999, 81, 189-192.		158
6	<i>env</i> Sequences of Simian Immunodeficiency Viruses from Chimpanzees in Cameroon Are Strongly Related to Those of Human Immunodeficiency Virus Group N from the Same Geographic Area. <i>Journal of Virology</i> , 2000, 74, 529-534.	3.4	152
7	Origin of the HIV-1 group O epidemic in western lowland gorillas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E1343-52.	7.1	136
8	Genomes of cryptic chimpanzee <i>Plasmodium</i> species reveal key evolutionary events leading to human malaria. <i>Nature Communications</i> , 2016, 7, 11078.	12.8	122
9	Foci of Endemic Simian Immunodeficiency Virus Infection in Wild-Living Eastern Chimpanzees (<i>Pan</i>) Tj ETQq1 1 0.784314 rgBT /Over 116	3.4	116
10	Resurgence of Ebola virus in 2021 in Guinea suggests a new paradigm for outbreaks. <i>Nature</i> , 2021, 597, 539-543.	27.8	113
11	Origin and Biology of Simian Immunodeficiency Virus in Wild-Living Western Gorillas. <i>Journal of Virology</i> , 2009, 83, 1635-1648.	3.4	106
12	Extensive survey on the prevalence and genetic diversity of SIVs in primate bushmeat provides insights into risks for potential new cross-species transmissions. <i>Infection, Genetics and Evolution</i> , 2010, 10, 386-396.	2.3	100
13	Multidisciplinary assessment of post-Ebola sequelae in Guinea (Postebogui): an observational cohort study. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 545-552.	9.1	96
14	Extraordinary Heterogeneity of Virological Outcomes in Patients Receiving Highly Antiretroviral Therapy and Monitored With the World Health Organization Public Health Approach in Sub-Saharan Africa and Southeast Asia. <i>Clinical Infectious Diseases</i> , 2014, 58, 99-109.	5.8	83
15	Depressive symptoms among survivors of Ebola virus disease in Conakry (Guinea): preliminary results of the PostEboGui cohort. <i>BMC Psychiatry</i> , 2017, 17, 127.	2.6	75
16	Phylogenetic Analysis of 49 Newly Derived HIV-1 Group O Strains: High Viral Diversity but No Group M-like Subtype Structure. <i>Virology</i> , 2002, 302, 259-273.	2.4	74
17	HIV-1 group N among HIV-1-seropositive individuals in Cameroon. <i>Aids</i> , 2000, 14, 2623-2625.	2.2	72
18	Compartmentalization of HIV-1 between Breast Milk and Blood of HIV-Infected Mothers. <i>Virology</i> , 2002, 300, 109-117.	2.4	71

#	ARTICLE	IF	CITATIONS
19	New Evidence of Long-lasting Persistence of Ebola Virus Genetic Material in Semen of Survivors: Table 1.. Journal of Infectious Diseases, 2016, 214, 1475-1476.	4.0	70
20	No evidence for transmission of SIVwrc from western red colobus monkeys (piliocolobus badius) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 7 hunting. BMC Microbiology, 2011, 11, 24.	3.3	69
21	Multiplex detection and dynamics of IgG antibodies to SARS-CoV2 and the highly pathogenic human coronaviruses SARS-CoV and MERS-CoV. Journal of Clinical Virology, 2020, 129, 104521.	3.1	68
22	Hepatitis C virus infection in cameroon: A cohort-effect. Journal of Medical Virology, 2005, 76, 208-214.	5.0	67
23	Evidence for continuing cross-species transmission of SIVsmm to humans. Aids, 2013, 27, 2488-2491.	2.2	66
24	Survey of Ebola Viruses in Frugivorous and Insectivorous Bats in Guinea, Cameroon, and the Democratic Republic of the Congo, 2015â€“2017. Emerging Infectious Diseases, 2018, 24, 2228-2240.	4.3	66
25	The hepatitis C virus epidemic in Cameroon: Genetic evidence for rapid transmission between 1920 and 1960. Infection, Genetics and Evolution, 2007, 7, 361-367.	2.3	64
26	HIV-1 Group O Infection in Cameroon, 1986 to 1998. Emerging Infectious Diseases, 2001, 7, 466-467.	4.3	62
27	Medical countermeasures during the 2018 Ebola virus disease outbreak in the North Kivu and Ituri Provinces of the Democratic Republic of the Congo: a rapid genomic assessment. Lancet Infectious Diseases, The, 2019, 19, 648-657.	9.1	62
28	Field Evaluation of Dried Blood Spots for Routine HIV-1 Viral Load and Drug Resistance Monitoring in Patients Receiving Antiretroviral Therapy in Africa and Asia. Journal of Clinical Microbiology, 2014, 52, 578-586.	3.9	60
29	Characterization of a new simian immunodeficiency virus strain in a naturally infected Pan troglodytes troglodyteschimpanzee with AIDS related symptoms. Retrovirology, 2011, 8, 4.	2.0	58
30	Low Rate of Mother-to-Child Transmission of HIV-1 After Nevirapine Intervention in a Pilot Public Health Program in Yaounde, Cameroon. Journal of Acquired Immune Deficiency Syndromes (1999), 2003, 34, 274-280.	2.1	56
31	Ocular Complications in Survivors of the Ebola Outbreak in Guinea. American Journal of Ophthalmology, 2017, 175, 114-121.	3.3	55
32	High rate of hepatitis C virus infection and predominance of genotype 4 among elderly inhabitants of a remote village of the rain forest of South Cameroon. Journal of Medical Virology, 2003, 71, 219-225.	5.0	54
33	Hepatitis C virus infection among pregnant women in Yaounde, Cameroon: Prevalence, viremia, and genotypes. Journal of Medical Virology, 2003, 69, 384-390.	5.0	51
34	Distribution and heterogeneity of hepatitis C genotypes in hepatitis patients in Cameroon. Journal of Medical Virology, 2005, 77, 390-398.	5.0	51
35	Novel simian foamy virus infections from multiple monkey species in women from the Democratic Republic of Congo. Retrovirology, 2012, 9, 100.	2.0	51
36	Closer to 90â€“90â€“90. The cascade of care after 10 years of ART scaleâ€“up in rural Malawi: a population study. Journal of the International AIDS Society, 2016, 19, 20673.	3.0	50

#	ARTICLE	IF	CITATIONS
37	Evolutionary history of human <i>Plasmodium vivax</i> revealed by genome-wide analyses of related ape parasites. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E8450-E8459.	7.1	50
38	Development of a Sensitive and Specific Serological Assay Based on Luminex Technology for Detection of Antibodies to Zaire Ebola Virus. <i>Journal of Clinical Microbiology</i> , 2017, 55, 165-176.	3.9	47
39	Simian Immunodeficiency Virus Infection in Wild-Caught Chimpanzees from Cameroon. <i>Journal of Virology</i> , 2005, 79, 1312-1319.	3.4	45
40	The origin and molecular epidemiology of HIV. <i>Expert Review of Anti-Infective Therapy</i> , 2013, 11, 885-896.	4.4	45
41	Forests and emerging infectious diseases: unleashing the beast within. <i>Environmental Research Letters</i> , 2020, 15, 083007.	5.2	42
42	Low Prevalence of HIV Type 1 Drug Resistance Mutations in Untreated, Recently Infected Patients from Burkina Faso, Côte d'Ivoire, Senegal, Thailand, and Vietnam: The ANRS 12134 Study. <i>AIDS Research and Human Retroviruses</i> , 2009, 25, 1193-1196.	1.1	40
43	Rapid Confirmation of the Zaire Ebola Virus in the Outbreak of the Equateur Province in the Democratic Republic of Congo: Implications for Public Health Interventions. <i>Clinical Infectious Diseases</i> , 2019, 68, 330-333.	5.8	39
44	Stability of the gorilla microbiome despite simian immunodeficiency virus infection. <i>Molecular Ecology</i> , 2015, 24, 690-697.	3.9	38
45	Multigenomic Delineation of <i>Plasmodium</i> Species of the <i>Laverania</i> Subgenus Infecting Wild-Living Chimpanzees and Gorillas. <i>Genome Biology and Evolution</i> , 2016, 8, 1929-1939.	2.5	38
46	High Prevalence of Anti-Severe Acute Respiratory Syndrome Coronavirus 2 (Anti-SARS-CoV-2) Antibodies After the First Wave of Coronavirus Disease 2019 (COVID-19) in Kinshasa, Democratic Republic of the Congo: Results of a Cross-sectional Household-Based Survey. <i>Clinical Infectious Diseases</i> , 2022, 74, 882-890.	5.8	38
47	Prevalence of infection among asymptomatic and paucisymptomatic contact persons exposed to Ebola virus in Guinea: a retrospective, cross-sectional observational study. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 308-316.	9.1	36
48	Primate lentiviruses use at least three alternative strategies to suppress NF- κ B-mediated immune activation. <i>PLoS Pathogens</i> , 2017, 13, e1006598.	4.7	34
49	Long-lasting severe immune dysfunction in Ebola virus disease survivors. <i>Nature Communications</i> , 2020, 11, 3730.	12.8	33
50	Molecular Evidence for the Presence of <i>Rickettsia felis</i> in the Feces of Wild-living African Apes. <i>PLoS ONE</i> , 2013, 8, e54679.	2.5	33
51	Interactions of plant lectins with the components of the bacterial cell wall peptidoglycan. <i>Biochemical Systematics and Ecology</i> , 1994, 22, 153-159.	1.3	30
52	In Vitro and In Vivo Human Herpesvirus 8 Infection of Placenta. <i>PLoS ONE</i> , 2008, 3, e4073.	2.5	30
53	Novel Multiplexed HIV/Simian Immunodeficiency Virus Antibody Detection Assay. <i>Emerging Infectious Diseases</i> , 2011, 17, 2277-2286.	4.3	29
54	RegaDB: community-driven data management and analysis for infectious diseases. <i>Bioinformatics</i> , 2013, 29, 1477-1480.	4.1	29

#	ARTICLE	IF	CITATIONS
55	LOW RISK OF MOTHER-TO-CHILD TRANSMISSION OF HEPATITIS C VIRUS IN YAOUNDÉ, CAMEROON: THE ANRS 1262 STUDY. <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 73, 460-466.	1.4	28
56	2018 Ebola virus disease outbreak in Équateur Province, Democratic Republic of the Congo: a retrospective genomic characterisation. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 641-647.	9.1	27
57	Noninvasive Follow-Up of Simian Immunodeficiency Virus Infection in Wild-Living Nonhabituated Western Lowland Gorillas in Cameroon. <i>Journal of Virology</i> , 2012, 86, 9760-9772.	3.4	26
58	A 40 months follow-up of Ebola virus disease survivors in Guinea (Postebogui) reveals longterm detection of Ebola viral RNA in semen and breast milk. <i>Open Forum Infectious Diseases</i> , 2019, 6, ofz482.	0.9	26
59	MOTHER-TO-CHILD TRANSMISSION OF HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 IN RELATION TO THE SEASON IN YAOUNDE, CAMEROON. <i>American Journal of Tropical Medicine and Hygiene</i> , 2003, 69, 447-449.	1.4	26
60	Simian T cell leukaemia virus type I subtype B in a wild-caught gorilla (<i>Gorilla gorilla gorilla</i>) and chimpanzee (<i>Pan troglodytes vellerosus</i>) from Cameroon. <i>Journal of General Virology</i> , 2004, 85, 25-29.	2.9	25
61	Reemergence of chloroquine-sensitive pfcr1 K76 <i>Plasmodium falciparum</i> genotype in southeastern Cameroon. <i>Malaria Journal</i> , 2017, 16, 130.	2.3	25
62	SIVcol Nef counteracts SERINC5 by promoting its proteasomal degradation but does not efficiently enhance HIV-1 replication in human CD4+ T cells and lymphoid tissue. <i>PLoS Pathogens</i> , 2018, 14, e1007269.	4.7	25
63	HIV-1 pol Gene Polymorphism and Antiretroviral Resistance Mutations in Drug-Naive Pregnant Women in Yaoundé, Cameroon. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2006, 42, 256-258.	2.1	24
64	Increasing HIV Type 1 Polymorphic Diversity But No Resistance to Antiretroviral Drugs in Untreated Patients from Central African Republic: A 2005 Study. <i>AIDS Research and Human Retroviruses</i> , 2006, 22, 1036-1044.	1.1	24
65	Vpu-mediated CD4 down-regulation and degradation is conserved among highly divergent SIVcpz strains. <i>Virology</i> , 2005, 335, 46-60.	2.4	23
66	Specific stimulation of HIV-1 replication in human placental trophoblasts by an antigen of <i>Plasmodium falciparum</i> . <i>Aids</i> , 2008, 22, 785-787.	2.2	23
67	Identification and Molecular Characterization of New Simian T Cell Lymphotropic Viruses in Nonhuman Primates Bushmeat from the Democratic Republic of Congo. <i>AIDS Research and Human Retroviruses</i> , 2012, 28, 628-635.	1.1	23
68	Comparison of different nucleic acid preparation methods to improve specific HIV-1 RNA isolation for viral load testing on dried blood spots. <i>Journal of Virological Methods</i> , 2018, 251, 75-79.	2.1	23
69	Extensive Serological Survey of Multiple African Nonhuman Primate Species Reveals Low Prevalence of Immunoglobulin G Antibodies to 4 Ebola Virus Species. <i>Journal of Infectious Diseases</i> , 2019, 220, 1599-1608.	4.0	23
70	SIVcpz from a naturally infected Cameroonian chimpanzee: Biological and genetic comparison with HIV-1 N. <i>Journal of Medical Primatology</i> , 2003, 29, 166-172.	0.6	22
71	Assessment of the gorilla gut virome in association with natural simian immunodeficiency virus infection. <i>Retrovirology</i> , 2018, 15, 19.	2.0	21
72	CD4 receptor diversity in chimpanzees protects against SIV infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 3229-3238.	7.1	21

#	ARTICLE	IF	CITATIONS
73	Wide Diversity of Coronaviruses in Frugivorous and Insectivorous Bat Species: A Pilot Study in Guinea, West Africa. <i>Viruses</i> , 2020, 12, 855.	3.3	20
74	Tumour necrosis factor-alpha stimulates HIV-1 replication in single-cycle infection of human term placental villi fragments in a time, viral dose and envelope dependent manner. <i>Retrovirology</i> , 2006, 3, 36.	2.0	19
75	Role of Wildlife in Emergence of Ebola Virus in Kaigbono (Likati), Democratic Republic of the Congo, 2017. <i>Emerging Infectious Diseases</i> , 2020, 26, 2205-2209.	4.3	19
76	Quantitative analysis of multiple V-region interactions among normal human IgG. <i>European Journal of Immunology</i> , 1996, 26, 710-716.	2.9	18
77	Natural polymorphisms of HIV-1 CRF01_AE integrase coding region in ARV-naïve individuals in Cambodia, Thailand and Vietnam: An ANRS AC12 working group study. <i>Infection, Genetics and Evolution</i> , 2011, 11, 38-43.	2.3	18
78	Ubiquitous Hepatocystis infections, but no evidence of Plasmodium falciparum-like malaria parasites in wild greater spot-nosed monkeys (<i>Cercopithecus nictitans</i>). <i>International Journal for Parasitology</i> , 2012, 42, 709-713.	3.1	18
79	Nonhuman primate retroviruses from Cambodia: High simian foamy virus prevalence, identification of divergent STLV-1 strains and no evidence of SIV infection. <i>Infection, Genetics and Evolution</i> , 2013, 18, 325-334.	2.3	18
80	Multiplex detection of antibodies to Chikungunya, Oâ€™nyong-nyong, Zika, Dengue, West Nile and Usutu viruses in diverse non-human primate species from Cameroon and the Democratic Republic of Congo. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009028.	3.0	18
81	Low Prevalence of Drug Resistance Transmitted Virus in HIV Type 1-Infected ARV-Naive Patients in Cambodia. <i>AIDS Research and Human Retroviruses</i> , 2009, 25, 543-545.	1.1	17
82	Antiretroviral treatment outcome in HIV-1 infected patients routinely followed up in capital cities and remote areas of Senegal, Mali and Guineaâ€™Conakry. <i>Journal of the International AIDS Society</i> , 2014, 17, 19315.	3.0	17
83	Legume lectins interact with muramic acid and N-acetylmuramic acid. <i>FEBS Letters</i> , 1991, 289, 102-104.	2.8	16
84	Rapid Increase of Community SARS-CoV-2 Seroprevalence during Second Wave of COVID-19, YaoundÃ©, Cameroon. <i>Emerging Infectious Diseases</i> , 2022, 28, .	4.3	16
85	Prevalence of pretreatment HIV drug resistance in West African and Southeast Asian countries. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 462-467.	3.0	15
86	Distinguishable Patterns of Connectivity in Serum Immunoglobulins from SLE Patients and Healthy Individuals. <i>Scandinavian Journal of Immunology</i> , 1997, 45, 408-416.	2.7	14
87	Plasmodium falciparum Infection Significantly Impairs Placental Cytokine Profile in HIV Infected Cameroonian Women. <i>PLoS ONE</i> , 2009, 4, e8114.	2.5	13
88	Full-length genome sequence of a simian immunodeficiency virus (SIV) infecting a captive agile mangabey (<i>Cercocebus agilis</i>) is closely related to SIVrcm infecting wild red-capped mangabeys (<i>Cercocebus torquatus</i>) in Cameroon. <i>Journal of General Virology</i> , 2010, 91, 2959-2964.	2.9	13
89	Investigating the Circulation of Ebola Viruses in Bats during the Ebola Virus Disease Outbreaks in the Equateur and North Kivu Provinces of the Democratic Republic of Congo from 2018. <i>Pathogens</i> , 2021, 10, 557.	2.8	13
90	Dried blood spots for HIV-1 drug resistance genotyping in decentralized settings in Senegal. <i>Journal of Medical Virology</i> , 2014, 86, 45-51.	5.0	12

#	ARTICLE	IF	CITATIONS
91	Extraordinary long-term and fluctuating persistence of Ebola virus RNA in semen of survivors in Guinea: implications for public health. <i>Clinical Microbiology and Infection</i> , 2017, 23, 412-413.	6.0	12
92	Understanding Long-term Evolution and Predictors of Sequelae of Ebola Virus Disease Survivors in Guinea: A 48-Month Prospective, Longitudinal Cohort Study (PostEboGui). <i>Clinical Infectious Diseases</i> , 2021, 73, 2166-2174.	5.8	12
93	Recognition of muramic acid and N-acetylmuramic acid by Leguminosae lectins: possible role in plant-bacteria interactions. <i>FEMS Microbiology Letters</i> , 1992, 92, 41-46.	1.8	10
94	A natural CCL5/RANTES variant antagonist for CCR1 and CCR3. <i>Immunogenetics</i> , 2006, 58, 533-541.	2.4	10
95	A cost-effective algorithm for the diagnosis of Hepatitis C virus infection and prediction of HCV viremia in Cameroon. <i>Journal of Virological Methods</i> , 2006, 133, 223-226.	2.1	10
96	Distinct efficacy of HIV-1 entry inhibitors to prevent cell-to-cell transfer of R5 and X4 viruses across a human placental trophoblast barrier in a reconstitution model in vitro. <i>Retrovirology</i> , 2008, 5, 31.	2.0	10
97	High Rate of Simian Immunodeficiency Virus (SIV) Infections in Wild Chimpanzees in Northeastern Gabon. <i>Viruses</i> , 2015, 7, 4997-5015.	3.3	10
98	Temporal evolution of the humoral antibody response after Ebola virus disease in Guinea: a 60-month observational prospective cohort study. <i>Lancet Microbe</i> , The, 2021, 2, e676-e684.	7.3	10
99	Molecular characterization of a new mosaic Simian Immunodeficiency Virus in a naturally infected tanzania monkey (<i>Chlorocebus tantalus</i>) from Cameroon: A challenge to the virus-host co-evolution of SIVagm in African green monkeys. <i>Infection, Genetics and Evolution</i> , 2015, 30, 65-73.	2.3	9
100	CD4 receptor diversity represents an ancient protection mechanism against primate lentiviruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	9
101	Mother-to-child transmission of human immunodeficiency virus type 1 in relation to the season in Yaounde, Cameroon. <i>American Journal of Tropical Medicine and Hygiene</i> , 2003, 69, 447-9.	1.4	9
102	Zoonotic origin of the human malaria parasite <i>Plasmodium malariae</i> from African apes. <i>Nature Communications</i> , 2022, 13, 1868.	12.8	9
103	Genetic diversity of STLV-2 and interspecies transmission of STLV-3 in wild-living bonobos. <i>Virus Evolution</i> , 2016, 2, vew011.	4.9	8
104	High and Rapid Increase in Seroprevalence for SARS-CoV-2 in Conakry, Guinea: Results From 3 Successive Cross-Sectional Surveys (ANRS COV16-ARIACOV). <i>Open Forum Infectious Diseases</i> , 2022, 9, ofac152.	0.9	8
105	Serological Evidence of Zika Virus Circulation in Burkina Faso. <i>Pathogens</i> , 2022, 11, 741.	2.8	8
106	Dynamics of Antibodies to Ebolaviruses in an Eidolon helvum Bat Colony in Cameroon. <i>Viruses</i> , 2022, 14, 560.	3.3	7
107	Fine sugar specificity of the <i>Butea frondosa</i> seed lectin. <i>Glycoconjugate Journal</i> , 1992, 9, 141-147.	2.7	6
108	Genetic diversity of simian lentivirus in wild De Brazza's monkeys (<i>Cercopithecus neglectus</i>) in Equatorial Africa. <i>Journal of General Virology</i> , 2010, 91, 1810-1816.	2.9	6

#	ARTICLE	IF	CITATIONS
109	Serological Evidence of Ebola Virus Infection in Rural Guinea before the 2014 West African Epidemic Outbreak. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 99, 425-427.	1.4	6
110	Longitudinal Survey of Coronavirus Circulation and Diversity in Insectivorous Bat Colonies in Zimbabwe. <i>Viruses</i> , 2022, 14, 781.	3.3	6
111	Evidence of STLV 2 and STLV 3 infections in wild living bonobos (<i>P. paniscus</i>) from the Democratic Republic of Congo. <i>Retrovirology</i> , 2011, 8, .	2.0	5
112	Added Value of an Anti-Ebola Serology for the Management of Clinically Suspected Ebola Virus Disease Patients Discharged as Negative in an Epidemic Context. <i>Journal of Infectious Diseases</i> , 2022, 226, 352-356.	4.0	5
113	Simian Immunodeficiency Viruses and the Origin of HIVs. , 2002, , 104-120.		4
114	Field evaluation of an open and polyvalent universal HIV-1/SIVcpz/SIVgor quantitative RT-PCR assay for HIV-1 viral load monitoring in comparison to Abbott RealTime HIV-1 in Cameroon. <i>Journal of Virological Methods</i> , 2016, 237, 121-126.	2.1	3
115	Full-Genome Characterization of Simian T-Cell Leukemia Virus Type 1 Subtype b from a Wild-Born Captive <i>Gorilla gorilla gorilla</i> with T-Cell Lymphoma. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
116	Identification of a Novel Simian Immunodeficiency Virus-Infected African Green Monkey (<i>Chlorocebus tantalus</i>) Confirms that Tantalus Monkeys in Cameroon Are Infected with a Mosaic SIVagm Lineage. <i>AIDS Research and Human Retroviruses</i> , 2020, 36, 167-170.	1.1	2
117	Quantification de la Charge Virale et tests de résistance du VIH-1 aux ARV à partir de échantillons DBS (Dried Blood Spots) chez des patients Guinéens sous traitement antirétroviral. <i>African Journal of Laboratory Medicine</i> , 2015, 4, .	0.6	2
118	Seroprevalence of IgG Antibodies Against Multiple Arboviruses in Bats from Cameroon, Guinea, and the Democratic Republic of Congo. <i>Vector-Borne and Zoonotic Diseases</i> , 2022, , .	1.5	2
119	Full Genome Characterization of a New Simian Immune Deficiency Virus Lineage in a Naturally Infected <i>Cercopithecus ascanius whitesidei</i> in the Democratic Republic of Congo Reveals High Genetic Diversity Among Red-Tailed Monkeys in Central and Eastern Africa. <i>AIDS Research and Human Retroviruses</i> . 2017, 33, 735-739.	1.1	1
120	Reply to Zhang et al. <i>Journal of Infectious Diseases</i> , 2020, 222, 1065-1066.	4.0	1
121	High HIV burden and recent transmission chains in rural forest areas in southern Cameroon, where ancestors of HIV-1 have been identified in ape populations. <i>Infection, Genetics and Evolution</i> , 2020, 84, 104358.	2.3	1
122	Evolution of simian retroviruses. , 0, , 117-149.		0
123	Noninvasive western lowland gorilla's health monitoring: A decade of simian immunodeficiency virus surveillance in southern Cameroon. <i>Ecology and Evolution</i> , 2018, 8, 10698-10710.	1.9	0
124	Recognition of muramic acid and N-acetylmuramic acid by Leguminosae lectins: possible role in plant-bacteria interactions. <i>FEMS Microbiology Letters</i> , 1992, 92, 41-46.	1.8	0