Yingying Li

List of Publications by Year in descending order

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35 papers	2,571 citations	304743 22 h-index	345221 36 g-index
36 all docs	36 docs citations	36 times ranked	3776 citing authors

#	Article	IF	Citations
1	Zoonotic origin of the human malaria parasite Plasmodium malariae from African apes. Nature Communications, 2022, 13, 1868.	12.8	9
2	CD4 receptor diversity represents an ancient protection mechanism against primate lentiviruses. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	9
3	Heightened resistance to host type 1 interferons characterizes HIV-1 at transmission and after antiretroviral therapy interruption. Science Translational Medicine, 2021, 13, .	12.4	54
4	Urine as a highâ€quality source of host genomic DNA from wild populations. Molecular Ecology Resources, 2021, 21, 170-182.	4.8	5
5	CD4 receptor diversity in chimpanzees protects against SIV infection. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3229-3238.	7.1	21
6	Contribution of proteasome-catalyzed peptidecis-splicing to viral targeting by CD8⁺T cells in HIV-1 infection. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 24748-24759.	7.1	48
7	Investigating zoonotic infection barriers to ape Plasmodium parasites using faecal DNA analysis. International Journal for Parasitology, 2018, 48, 531-542.	3.1	9
8	Allometry and Ecology of the Bilaterian Gut Microbiome. MBio, 2018, 9, .	4.1	29
9	Oesophagostomiasis in nonâ€human primates of Gombe National Park, Tanzania. American Journal of Primatology, 2018, 80, e22572.	1.7	20
10	Destabilization of the gut microbiome marks the endâ€stage of simian immunodeficiency virus infection in wild chimpanzees. American Journal of Primatology, 2018, 80, e22515.	1.7	27
11	Socioecological correlates of clinical signs in two communities of wild chimpanzees (Pan) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf	1.7	18
12	Completeness of HIV-1 Envelope Glycan Shield at Transmission Determines Neutralization Breadth. Cell Reports, 2018, 25, 893-908.e7.	6.4	91
13	Adaptive Evolution of RH5 in Ape Plasmodium species of the Laverania Subgenus. MBio, 2018, 9, .	4.1	13
14	Reply to Forni et al., â€œMultiple Selected Changes May Modulate the Molecular Interaction between Laverania RH5 and Primate Basiginâ€. MBio, 2018, 9, .	4.1	1
15	Evolutionary history of humanPlasmodium vivaxrevealed by genome-wide analyses of related ape parasites. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E8450-E8459.	7.1	50
16	Out of Africa: origins and evolution of the human malaria parasites Plasmodium falciparum and Plasmodium vivax. International Journal for Parasitology, 2017, 47, 87-97.	3.1	163
17	Resistance to type 1 interferons is a major determinant of HIV-1 transmission fitness. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E590-E599.	7.1	137
18	Chimpanzees breed with genetically dissimilar mates. Royal Society Open Science, 2017, 4, 160422.	2.4	28

#	ARTICLE	IF	CITATIONS
19	Bonobos Maintain Immune System Diversity with Three Functional Types of MHC-B. Journal of Immunology, 2017, 198, 3480-3493.	0.8	19
20	Wild bonobos host geographically restricted malaria parasites including a putative new Laverania species. Nature Communications, 2017, 8, 1635.	12.8	45
21	Effective treatment of SIVcpz-induced immunodeficiency in a captive western chimpanzee. Retrovirology, 2017, 14, 35.	2.0	12
22	Multigenomic Delineation ofPlasmodiumSpecies of theLaveraniaSubgenus Infecting Wild-Living Chimpanzees and Gorillas. Genome Biology and Evolution, 2016, 8, 1929-1939.	2.5	38
23	Genomes of cryptic chimpanzee Plasmodium species reveal key evolutionary events leading to human malaria. Nature Communications, 2016, 7, 11078.	12.8	122
24	Genetic diversity of STLV-2 and interspecies transmission of STLV-3 in wild-living bonobos. Virus Evolution, 2016, 2, vew011.	4.9	8
25	Longitudinal Antigenic Sequences and Sites from Intra-Host Evolution (LASSIE) Identifies Immune-Selected HIV Variants. Viruses, 2015, 7, 5443-5475.	3.3	26
26	Signature Patterns of MHC Diversity in Three Gombe Communities of Wild Chimpanzees Reflect Fitness in Reproduction and Immune Defense against SIVcpz. PLoS Biology, 2015, 13, e1002144.	5.6	31
27	Rapid changes in the gut microbiome during human evolution. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 16431-16435.	7.1	287
28	African origin of the malaria parasite Plasmodium vivax. Nature Communications, 2014, 5, 3346.	12.8	167
29	Evidence for continuing cross-species transmission of SIVsmm to humans. Aids, 2013, 27, 2488-2491.	2.2	66
30	Eastern Chimpanzees, but Not Bonobos, Represent a Simian Immunodeficiency Virus Reservoir. Journal of Virology, 2012, 86, 10776-10791.	3.4	73
31	Impact of Simian Immunodeficiency Virus Infection on Chimpanzee Population Dynamics. PLoS Pathogens, 2010, 6, e1001116.	4.7	91
32	Origin and Biology of Simian Immunodeficiency Virus in Wild-Living Western Gorillas. Journal of Virology, 2009, 83, 1635-1648.	3.4	106
33	Increased mortality and AIDS-like immunopathology in wild chimpanzees infected with SIVcpz. Nature, 2009, 460, 515-519.	27.8	315
34	SIV infection in wild gorillas. Nature, 2006, 444, 164-164.	27.8	315
35	Foci of Endemic Simian Immunodeficiency Virus Infection in Wild-Living Eastern Chimpanzees (Pan) Tj ETQq1 1 0.784314 rgBT /Overlo	3.4	116