Tony L Goldberg

List of Publications by Year
in descending order

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191 papers	8,270 citations	38660 50 h-index	64668 79 g-index
201 all docs	201 docs citations	201 times ranked	9469 citing authors

#	Article	IF	Citations
1	Primates and the Ecology of their Infectious Diseases: How will Anthropogenic Change Affect Host-Parasite Interactions?. Evolutionary Anthropology, 2005, 14, 134-144.	1.7	277
2	Reorganization and expansion of the nidoviral family Arteriviridae. Archives of Virology, 2016, 161, 755-768.	0.9	254
3	Evolutionary trends in host physiology outweigh dietary niche in structuring primate gut microbiomes. ISME Journal, 2019, 13, 576-587.	4.4	236
4	Culex pipiens (Diptera: Culicidae): A Bridge Vector of West Nile Virus to Humans. Journal of Medical Entomology, 2008, 45, 125-128.	0.9	227
5	Characterization of the Fecal Microbiome from Non-Human Wild Primates Reveals Species Specific Microbial Communities. PLoS ONE, 2010, 5, e13963.	1.1	225
6	Biodiversity Loss Affects Global Disease Ecology. BioScience, 2009, 59, 945-954.	2.2	211
7	Local impact of temperature and precipitation on West Nile virus infection in Culex species mosquitoes in northeast Illinois, USA. Parasites and Vectors, 2010, 3, 19.	1.0	211
8	Gastrointestinal Bacterial Transmission among Humans, Mountain Gorillas, and Livestock in Bwindi Impenetrable National Park, Uganda. Conservation Biology, 2008, 22, 1600-1607.	2.4	183
9	Host selection by Culex pipiens mosquitoes and West Nile virus amplification. American Journal of Tropical Medicine and Hygiene, 2009, 80, 268-78.	0.6	157
10	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	0.9	153
11	Patterns of gastrointestinal bacterial exchange between chimpanzees and humans involved in research and tourism in western Uganda. Biological Conservation, 2007, 135, 511-517.	1.9	152
12	Forest Fragmentation as Cause of Bacterial Transmission among Nonhuman Primates, Humans, and Livestock, Uganda. Emerging Infectious Diseases, 2008, 14, 1375-1382.	2.0	145
13	Effects of recreational angling on the post-release behavior and predation of bonefish (Albula) Tj ETQq1 1 0.784314 rgBT /Overlock 1 and Ecology, 2007, 346, 127-133.	0.7	138
14	Molecular Epidemiology of Cross-Species Giardia duodenalis Transmission in Western Uganda. PLoS Neglected Tropical Diseases, 2010, 4, e683.	1.3	136
15	Genetic correlates of social behaviour in wild chimpanzees: evidence from mitochondrial DNA. Animal Behaviour, 1997, 54, 559-570.	0.8	126
16	A Multicomponent Animal Virus Isolated from Mosquitoes. Cell Host and Microbe, 2016, 20, 357-367.	5.1	123
17	The Fertility of Agricultural and Non-Agricultural Traditional Societies. Population Studies, 1993, 47, 269-281.	1.1	112
18	Culex Flavivirus and West Nile Virus Mosquito Coinfection and Positive Ecological Association in Chicago, United States. Vector-Borne and Zoonotic Diseases, 2011, 11, 1099-1105.	0.6	106

#	Article	IF	Citations
19	Fine-Scale Variation in Vector Host Use and Force of Infection Drive Localized Patterns of West Nile Virus Transmission. PLoS ONE, 2011, 6, e23767.	1.1	106
20	Increased Infectious Disease Susceptibility Resulting from Outbreeding Depression. Conservation Biology, 2005, 19, 455-462.	2.4	98
21	Rapid Amplification of West Nile Virus: The Role of Hatch-Year Birds. Vector-Borne and Zoonotic Diseases, 2008, 8, 57-68.	0.6	97
22	A Novel Hepacivirus with an Unusually Long and Intrinsically Disordered NS5A Protein in a Wild Old World Primate. Journal of Virology, 2013, 87, 8971-8981.	1.5	88
23	Wild Birds and Urban Ecology of Ticks and Tick-borne Pathogens, Chicago, Illinois, USA, 2005â€“2010. Emerging Infectious Diseases, 2012, 18, 1589-1595.	2.0	86
24	Coinfection of Ugandan Red Colobus (Procolobus [Piliocolobus] rufomitratus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 5 2009, 83, 11318-11329.	1.5	82
25	Avian host community structure and prevalence of West Nile virus in Chicago, Illinois. Oecologia, 2009, 159, 415-424.	0.9	82
26	Lethal Respiratory Disease Associated with Human RhinovirusÂC in Wild Chimpanzees, Uganda, 2013. Emerging Infectious Diseases, 2018, 24, 267-274.	2.0	80
27	Drivers of Bushmeat Hunting and Perceptions of Zoonoses in Nigerian Hunting Communities. PLoS Neglected Tropical Diseases, 2015, 9, e0003792.	1.3	79
28	Giardia sp. and Cryptosporidium sp. Infections in Primates in Fragmented and Undisturbed Forest in Western Uganda. Journal of Parasitology, 2007, 93, 439-440.	0.3	77
29	Simultaneous outbreaks of respiratory disease in wild chimpanzees caused by distinct viruses of human origin. Emerging Microbes and Infections, 2019, 8, 139-149.	3.0	77
30	Going, Going, Gone: A 15-Year History of the Decline of Primates in Forest Fragments near Kibale National Park, Uganda. , 2013, , 89-100.		77
31	Dynamically evolving novel overlapping gene as a factor in the SARS-CoV-2 pandemic. ELife, 2020, 9, .	2.8	74
32	Is catch-and-release recreational angling compatible with no-take marine protected areas?. Ocean and Coastal Management, 2006, 49, 342-354.	2.0	73
33	Fecal microbiomes of nonâ€human primates in Western Uganda reveal speciesâ€specific communities largely resistant to habitat perturbation. American Journal of Primatology, 2014, 76, 347-354.	0.8	72
34	Epidemiology and Molecular Relationships of Cryptosporidium spp. in People, Primates, and Livestock from Western Uganda. PLoS Neglected Tropical Diseases, 2012, 6, e1597.	1.3	68
35	Altruism towards panhandlers: Who gives?. Human Nature, 1995, 6, 79-89.	0.8	67
36	Correlation of cell-mediated immunity against porcine reproductive and respiratory syndrome virus with protection against reproductive failure in sows during outbreaks of porcine reproductive and respiratory syndrome in commercial herds. Journal of the American Veterinary Medical Association, 2005, 226, 1707-1711.	0.2	67

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37	Genetic Variation Associated with Mammalian Feeding inCulex pipiensfrom a West Nile Virus Epidemic Region in Chicago, Illinois. Vector-Borne and Zoonotic Diseases, 2009, 9, 637-642.	0.6	65
38	GB Virus C Coinfections in West African Ebola Patients. Journal of Virology, 2015, 89, 2425-2429.	1.5	65
39	Is the Fertility of Agriculturalists Higher Than That of Nonagriculturalists?. Current Anthropology, 1993, 34, 778-785.	0.8	65
40	Fine-scale genetic variation and evolution of West Nile Virus in a transmission â€œhot spotâ€•in suburban Chicago, USA. Virology, 2008, 374, 381-389.	1.1	64
41	Hidden Population Structure and Cross-species Transmission of Whipworms (Trichuris sp.) in Humans and Non-human Primates in Uganda. PLoS Neglected Tropical Diseases, 2014, 8, e3256.	1.3	64
42	ICTV Virus Taxonomy Profile: Arteriviridae 2021. Journal of General Virology, 2021, 102, .	1.3	64
43	Novel, Divergent Simian Hemorrhagic Fever Viruses in a Wild Ugandan Red Colobus Monkey Discovered Using Direct Pyrosequencing. PLoS ONE, 2011, 6, e19056.	1.1	63
44	Nodule Worm Infection in Humans and Wild Primates in Uganda: Cryptic Species in a Newly Identified Region of Human Transmission. PLoS Neglected Tropical Diseases, 2014, 8, e2641.	1.3	63
45	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	0.9	62
46	Exceptional Simian Hemorrhagic Fever Virus Diversity in a Wild African Primate Community. Journal of Virology, 2013, 87, 688-691.	1.5	61
47	An agent-based model of red colobus resources and disease dynamics implicates key resource sites as hot spots of disease transmission. Ecological Modelling, 2010, 221, 2491-2500.	1.2	59
48	Movement patterns of bonefish (Albula vulpes) in tidal creeks and coastal waters of Eleuthera, The Bahamas. Fisheries Research, 2013, 147, 404-412.	0.9	58
49	The need for a global health ethic. Lancet, The, 2015, 386, e37-e39.	6.3	58
50	Relatives of rubella virus in diverse mammals. Nature, 2020, 586, 424-428.	13.7	58
51	High Rates of Escherichia coli Transmission between Livestock and Humans in Rural Uganda. Journal of Clinical Microbiology, 2008, 46, 3187-3191.	1.8	56
52	Health and disease in the people, primates, and domestic animals of Kibale National Park: implications for conservation. , 2008, , 75-87.		55
53	Beyond Bushmeat: Animal Contact, Injury, and Zoonotic Disease Risk in Western Uganda. EcoHealth, 2014, 11, 534-543.	0.9	54
54	Strain Variation in an Emerging Iridovirus of Warm-Water Fishes. Journal of Virology, 2003, 77, 8812-8818.	1.5	53

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55	Dispersal of Adult Culex Mosquitoes in an Urban West Nile Virus Hotspot: A Mark-Capture Study Incorporating Stable Isotope Enrichment of Natural Larval Habitats. PLoS Neglected Tropical Diseases, 2014, 8, e2768.	1.3	53
56	Sickness behaviour associated with non-lethal infections in wild primates. Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20151436.	1.2	53
57	Optimization of Analytical Parameters for Inferring Relationships among Escherichia coli Isolates from Repetitive-Element PCR by Maximizing Correspondence with Multilocus Sequence Typing Data. Applied and Environmental Microbiology, 2006, 72, 6049-6052.	1.4	52
58	Social Behaviours and Networks of Vervet Monkeys Are Influenced by Gastrointestinal Parasites. PLoS ONE, 2016, 11, e0161113.	1.1	50
59	Diversity and evolution of West Nile virus in Illinois and the United States, 2002â€“2005. Virology, 2007, 360, 143-149.	1.1	48
60	Zoonotic Potential of Simian Arteriviruses. Journal of Virology, 2016, 90, 630-635.	1.5	48
61	Spatial ecology and residency patterns of adult great barracuda (Sphyraena barracuda) in coastal waters of The Bahamas. Marine Biology, 2011, 158, 2227-2237.	0.7	47
62	Discovery and Characterization of Distinct Simian Pegiviruses in Three Wild African Old World Monkey Species. PLoS ONE, 2014, 9, e98569.	1.1	45
63	Antibiotic-Resistant Escherichia coli and Class 1 Integrons in Humans, Domestic Animals, and Wild Primates in Rural Uganda. Applied and Environmental Microbiology, 2018, 84, .	1.4	41
64	Mass mortality in freshwater mussels (Actinonaias pectorosa) in the Clinch River, USA, linked to a novel densovirus. Scientific Reports, 2020, 10, 14498.	1.6	41
65	Associations between genetics, farm characteristics and clinical disease in field outbreaks of porcine reproductive and respiratory syndrome virus. Preventive Veterinary Medicine, 2000, 43, 293-302.	0.7	40
66	Eating Bushmeat Improves Food Security in a Biodiversity and Infectious Disease â€œHotspotâ€• EcoHealth, 2020, 17, 125-138.	0.9	40
67	Broad Protection against Avian Influenza Virus by Using a Modified Vaccinia Ankara Virus Expressing a Mosaic Hemagglutinin Gene. Journal of Virology, 2014, 88, 13300-13309.	1.5	39
68	Spatial ecology of juvenile lemon sharks (Negaprion brevirostris) in tidal creeks and coastal waters of Eleuthera, The Bahamas. Environmental Biology of Fishes, 2010, 89, 95-104.	0.4	38
69	Simian Hemorrhagic Fever Virus Cell Entry Is Dependent on CD163 and Uses a Clathrin-Mediated Endocytosis-Like Pathway. Journal of Virology, 2015, 89, 844-856.	1.5	38
70	Evaluation of a Stable Isotope Method to Mark Naturally-Breeding Larval Mosquitoes for Adult Dispersal Studies. Journal of Medical Entomology, 2012, 49, 61-70.	0.9	37
71	Discovery and full genome characterization of two highly divergent simian immunodeficiency viruses infecting black-and-white colobus monkeys (Colobus guereza) in Kibale National Park, Uganda. Retrovirology, 2013, 10, 107.	0.9	37
72	Predicting West Nile Virus Infection Risk From the Synergistic Effects of Rainfall and Temperature. Journal of Medical Entomology, 2016, 53, 935-944.	0.9	37

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73	Metagenomic assessment of adventitious viruses in commercial bovine sera. Biologicals, 2017, 47, 64-68.	0.5	37
74	High Genetic Diversity and Adaptive Potential of Two Simian Hemorrhagic Fever Viruses in a Wild Primate Population. PLoS ONE, 2014, 9, e90714.	1.1	36
75	Diverse RNA viruses of arthropod origin in the blood of fruit bats suggest a link between bat and arthropod viromes. Virology, 2019, 528, 64-72.	1.1	36
76	Effects of Temperature on the Susceptibility of Largemouth Bass to Largemouth Bass Virus. Journal of Aquatic Animal Health, 2003, 15, 215-220.	0.6	35
77	Molecular phylogenetics and historical biogeography of east African chimpanzees. Biological Journal of the Linnean Society, 1997, 61, 301-324.	0.7	34
78	Impacts of dissolved oxygen on the behavior and physiology of bonefish: Implications for live-release angling tournaments. Journal of Experimental Marine Biology and Ecology, 2011, 402, 19-26.	0.7	34
79	Risk factors for respiratory illness in a community of wild chimpanzees (<i>Pan troglodytes) Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.1	34
80	Novel reovirus associated with epidemic mortality in wild largemouth bass (Micropterus salmoides). Journal of General Virology, 2016, 97, 2482-2487.	1.3	34
81	Multi-year evolutionary dynamics of West Nile virus in suburban Chicago, USA, 2005â€“2007. Philosophical Transactions of the Royal Society B: Biological Sciences, 2010, 365, 1871-1878.	1.8	33
82	Co-infection and cross-species transmission of divergent Hepatocystis lineages in a wild African primate community. International Journal for Parasitology, 2013, 43, 613-619.	1.3	32
83	Kanyawara Virus: A Novel Rhabdovirus Infecting Newly Discovered Nycteribiid Bat Flies Infesting Previously Unknown Pteropodid Bats in Uganda. Scientific Reports, 2017, 7, 5287.	1.6	32
84	Biodiversity of protists and nematodes in the wild nonhuman primate gut. ISME Journal, 2020, 14, 609-622.	4.4	32
85	Application of phylogeny reconstruction and character-evolution analysis to inferring patterns of directional microbial transmission. Preventive Veterinary Medicine, 2003, 61, 59-70.	0.7	31
86	Age Patterning in Wild Chimpanzee Gut Microbiota Diversity Reveals Differences from Humans in Early Life. Current Biology, 2021, 31, 613-620.e3.	1.8	31
87	Two Novel Simian Arteriviruses in Captive and Wild Baboons (Papio spp.). Journal of Virology, 2014, 88, 13231-13239.	1.5	28
88	Detection of Known and Novel Adenoviruses in Cattle Wastes via Broad-Spectrum Primers. Applied and Environmental Microbiology, 2011, 77, 5001-5008.	1.4	27
89	Prevalence of filarioid nematodes and trypanosomes in American robins and house sparrows, Chicago USA. International Journal for Parasitology: Parasites and Wildlife, 2013, 2, 42-49.	0.6	26
90	Arteriviruses, Pegiviruses, and Lentiviruses Are Common among Wild African Monkeys. Journal of Virology, 2016, 90, 6724-6737.	1.5	26

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91	Prevalence and Risk Factors Associated with Hemoparasites in Cattle and Goats at the Edge of Kibale National Park, Western Uganda. Journal of Parasitology, 2017, 103, 69-74.	0.3	26
92	Effects of Factors Related to Water Quality and Population Density on the Sensitivity of Juvenile Largemouth Bass to Mortality Induced by Viral Infection. Journal of Aquatic Animal Health, 2005, 17, 304-314.	0.6	25
93	Uncovering zoonoses awareness in an emerging disease â€˜hotspotâ€™. Social Science and Medicine, 2015, 129, 78-86.	1.8	25
94	Chytrid Fungus in Frogs from an Equatorial African Montane Forest in Western Uganda. Journal of Wildlife Diseases, 2007, 43, 521-524.	0.3	23
95	Serologic Evidence for Novel Poxvirus in Endangered Red Colobus Monkeys, Western Uganda. Emerging Infectious Diseases, 2008, 14, 801-803.	2.0	23
96	Physiological and Behavioral Effects of Capture Darting on Red Colobus Monkeys (Procolobus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54 of Primatology, 2013, 34, 1020-1031.	0.9	23
97	Providing health care to improve community perceptions of protected areas. Oryx, 2015, 49, 636-642.	0.5	23
98	Durable sequence stability and bone marrow tropism in a macaque model of human pegivirus infection. Science Translational Medicine, 2015, 7, 305ra144.	5.8	22
99	Primate reinfection with gastrointestinal parasites: behavioural and physiological predictors of parasite acquisition. Animal Behaviour, 2016, 117, 105-113.	0.8	22
100	Territorial song in the Anna's hummingbird, Calypte anna: costs of attraction and benefits of deterrence. Animal Behaviour, 1991, 42, 221-226.	0.8	21
101	Fatal Metacestode Infection in Bornean Orangutan Caused by UnknownVersteriaSpecies. Emerging Infectious Diseases, 2014, 20, 109-113.	2.0	21
102	Historical Outbreaks of Simian Hemorrhagic Fever in Captive Macaques Were Caused by Distinct Arteriviruses. Journal of Virology, 2015, 89, 8082-8087.	1.5	21
103	Decreased Flight Activity inCulex pipiens(Diptera: Culicidae) Naturally Infected With Culex flavivirus. Journal of Medical Entomology, 2016, 53, 233-236.	0.9	21
104	Diversity, Transmission, and Cophylogeny of Ledanteviruses (Rhabdoviridae: Ledantevirus) and Nycteribiid Bat Flies Parasitizing Angolan Soft-Furred Fruit Bats in Bundibugyo District, Uganda. Microorganisms, 2020, 8, 750.	1.6	21
105	Host group formation decreases exposure to vector-borne disease: a field experiment in a â€˜hotspotâ€™ of West Nile virus transmission. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20141586.	1.2	20
106	Evaluation of a Novel Emergence Trap to Study Culex Mosquitoes in Urban Catch Basins. Journal of the American Mosquito Control Association, 2011, 27, 142-147.	0.2	19
107	Genetic and antigenic relationships of vesicular stomatitis viruses from South America. Archives of Virology, 2011, 156, 1961-1968.	0.9	19
108	An outbreak of the 2009 influenza a (H1N1) virus in a childrenâ€™s hospital. Influenza and Other Respiratory Viruses, 2012, 6, 374-379.	1.5	19

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109	Nestling Passerines Are Not Important Hosts for Amplification of West Nile Virus in Chicago, Illinois. Vector-Borne and Zoonotic Diseases, 2009, 9, 13-18.	0.6	18
110	Protozoan parasites in groupâ€living primates: testing the biological island hypothesis. American Journal of Primatology, 2012, 74, 510-517.	0.8	18
111	Definitive Hosts ofVersteriaTapeworms (Cestoda: Taeniidae) Causing Fatal Infection in North America. Emerging Infectious Diseases, 2016, 22, 707-710.	2.0	18
112	Severe neurologic disease and chick mortality in crested screamers (Chauna torquata) infected with a novel Gyrovirus. Virology, 2018, 520, 111-115.	1.1	18
113	Life on the Rainforest Edge: Food Security in the Agricultural-Forest Frontier of Cross River State, Nigeria. Frontiers in Sustainable Food Systems, 2019, 3, .	1.8	18
114	Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in Hepatocystis parasites from naturally infected red colobus monkeys. PLoS Pathogens, 2020, 16, e1008717.	2.1	18
115	Mussel Mass Mortality and the Microbiome: Evidence for Shifts in the Bacterial Microbiome of a Declining Freshwater Bivalve. Microorganisms, 2021, 9, 1976.	1.6	18
116	Title is missing!. International Journal of Primatology, 1998, 19, 237-254.	0.9	17
117	Effects of Practices Related to Catch-and-Release Angling on Mortality and Viral Transmission in Juvenile Largemouth Bass Infected with Largemouth Bass Virus. Journal of Aquatic Animal Health, 2005, 17, 315-322.	0.6	17
118	Is Markhamia luteaâ€™s abundance determined by animal foraging?. Forest Ecology and Management, 2013, 308, 62-66.	1.4	17
119	Naturally Circulating Hepatitis A Virus in Olive Baboons, Uganda. Emerging Infectious Diseases, 2016, 22, 1308-1310.	2.0	17
120	Drivers for emerging issues in animal and plant health. EFSA Journal, 2016, 14, e00512.	0.9	17
121	Combining Footwear with Public Health Iconography to Prevent Soil-Transmitted Helminth Infections. American Journal of Tropical Medicine and Hygiene, 2017, 96, 205-213.	0.6	17
122	A Sarcina bacterium linked to lethal disease in sanctuary chimpanzees in Sierra Leone. Nature Communications, 2021, 12, 763.	5.8	17
123	Skin fungal assemblages of bats vary based on susceptibility to white-nose syndrome. ISME Journal, 2021, 15, 909-920.	4.4	16
124	Bat Flies of the Family Streblidae (Diptera: Hippoboscoidea) Host Relatives of Medically and Agriculturally Important â€œBat-Associatedâ€•Viruses. Viruses, 2021, 13, 860.	1.5	16
125	Deep sequencing identifies two genotypes and high viral genetic diversity of human pegivirus (GB virus) Tj ETQq1 1 0.784314 rgBT /Ov	1.3	15
126	Culex Flavivirus During West Nile Virus Epidemic and Interepidemic Years in Chicago, United States. Vector-Borne and Zoonotic Diseases, 2017, 17, 567-575.	0.6	15

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127	Demography, life-history trade-offs, and the gastrointestinal virome of wild chimpanzees. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190613.	1.8	15
128	Faecal parasites increase with age but not reproductive effort in wild female chimpanzees. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190614.	1.8	15
129	Opportunities for respiratory disease transmission from people to chimpanzees at an East African tourism site. American Journal of Primatology, 2021, 83, e23228.	0.8	15
130	Development and Evaluation of a Blocking Enzyme-Linked Immunosorbent Assay and Virus Neutralization Assay To Detect Antibodies to Viral Hemorrhagic Septicemia Virus. Vaccine Journal, 2014, 21, 435-442.	3.2	14
131	Discovery and full genome characterization of a new SIV lineage infecting red-tailed guenons (Cercopithecus ascanius schmidti) in Kibale National Park, Uganda. Retrovirology, 2014, 11, 55.	0.9	14
132	Divergent Simian Arteriviruses Cause Simian Hemorrhagic Fever of Differing Severities in Macaques. MBio, 2016, 7, e02009-15.	1.8	14
133	Overlap in the Seasonal Infection Patterns of Avian Malaria Parasites and West Nile Virus in Vectors and Hosts. American Journal of Tropical Medicine and Hygiene, 2016, 95, 1121-1129.	0.6	14
134	The evolution of a super-swarm of foot-and-mouth disease virus in cattle. PLoS ONE, 2019, 14, e0210847.	1.1	14
135	Adaptation and Limitations of Established Hemagglutination Inhibition Assays for the Detection of Porcine Antiâ€"Swine Influenza Virus H1N2 Antibodies. Journal of Veterinary Diagnostic Investigation, 2004, 16, 264-270.	0.5	13
136	Immunologic responses and reproductive outcomes following exposure to wild-type or attenuated porcine reproductive and respiratory syndrome virus in swine under field conditions. Journal of the American Veterinary Medical Association, 2006, 228, 1082-1088.	0.2	13
137	West Nile virus may have hitched a ride across the Western United States onCulex tarsalismosquitoes. Molecular Ecology, 2010, 19, 1518-1519.	2.0	13
138	FIELD INVESTIGATION OF INNATE IMMUNITY IN PASSERINE BIRDS IN SUBURBAN CHICAGO, ILLINOIS, USA. Journal of Wildlife Diseases, 2011, 47, 603-611.	0.3	13
139	Changes in physiological stress and behaviour in semi-free-ranging red-capped mangabeys () Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf Biological Sciences, 2016, 283, 20161201.	1.2	13
140	Selective constraint and adaptive potential of West Nile virus within and among naturally infected avian hosts and mosquito vectors. Virus Evolution, 2018, 4, vey013.	2.2	13
141	Multidecade Mortality and a Homolog of Hepatitis C Virus in Bald Eagles (Haliaeetus leucocephalus), the National Bird of the USA. Scientific Reports, 2019, 9, 14953.	1.6	13
142	Coincident Tick Infestations in the Nostrils of Wild Chimpanzees and a Human in Uganda. American Journal of Tropical Medicine and Hygiene, 2013, 89, 924-927.	0.6	12
143	WEST NILE VIRUS ANTIBODY DECAY RATE IN FREE-RANGING BIRDS. Journal of Wildlife Diseases, 2015, 51, 601.	0.3	12
144	Safeguarding biodiversity: what is perceived as working, according to the conservation community?. Oryx, 2016, 50, 302-307.	0.5	12

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145	Demography and health of â€œvillage dogsâ€•in rural Western Uganda. Preventive Veterinary Medicine, 2017, 137, 24-27.	0.7	12
146	Identification of Avian and Hemoparasite DNA in Blood-Engorged Abdomens of Culex pipiens (Diptera;) Tj ETQq0 0 0 rgBT /Overlock 10 T Entomology, 2015, 52, 461-468.	0.9	11
147	Spatial Overlap Between People and Non-human Primates in a Fragmented Landscape. EcoHealth, 2017, 14, 88-99.	0.9	11
148	Viruses associated with ill health in wild chimpanzees. American Journal of Primatology, 2022, 84, e23358.	0.8	11
149	Evaluation of contact exposure as a method for acclimatizing growing pigs to porcine reproductive and respiratory syndrome virus. Journal of the American Veterinary Medical Association, 2008, 232, 1530-1535.	0.2	10
150	Comparison of gastrointestinal parasite communities in vervet monkeys. Integrative Zoology, 2017, 12, 512-520.	1.3	10
151	Genome-Wide Patterns of Gene Expression in a Wild Primate Indicate Species-Specific Mechanisms Associated with Tolerance to Natural Simian Immunodeficiency Virus Infection. Genome Biology and Evolution, 2019, 11, 1630-1643.	1.1	10
152	Mycobiome Traits Associated with Disease Tolerance Predict Many Western North American Bat Species Will Be Susceptible to White-Nose Syndrome. Microbiology Spectrum, 2021, 9, e0025421.	1.2	10
153	Killing of a pearl-spotted owlet (Glaucidium perlatum) by male red colobus monkeys (Procolobus) Tj ETQq1 1 0.784314 rgBT /Overloc Primatology, 2006, 68, 1007-1011.	0.8	9
154	Genome Sequences of Simian Hemorrhagic Fever Virus Variant NIH LVR42-0/M6941 Isolates (Arteriviridae: Arterivirus). Genome Announcements, 2014, 2, .	0.8	9
155	Lung fluke (Paragonimus africanus) infects Nigerian red-capped mangabeys and causes respiratory disease. International Journal for Parasitology: Parasites and Wildlife, 2015, 4, 329-332.	0.6	9
156	Inferring the Geographic Origins of "Refugee" Chimpanzees in Uganda from Mitochondrial DNA Sequences. Conservation Biology, 1997, 11, 1441-1446.	2.4	8
157	Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat (Lissonycteris angolensis) in Uganda Dramatically Expands the Global Distribution of an Emerging Bat-Borne Respiratory Virus. Viruses, 2020, 12, 740.	1.5	8
158	Diversity ofWolbachia pipientisStrainwPip in a Genetically Admixtured, Above-GroundCulex pipiens(Diptera: Culicidae) Population: Association With Form Molestus Ancestry and Host Selection Patterns. Journal of Medical Entomology, 2012, 49, 474-481.	0.9	7
159	A multi-omic investigation of male lower urinary tract symptoms: Potential role for JC virus. PLoS ONE, 2021, 16, e0246266.	1.1	7
160	Specific Detection of Two Divergent Simian Arteriviruses Using RNAscope In Situ Hybridization. PLoS ONE, 2016, 11, e0151313.	1.1	7
161	Zootherapy as a potential pathway for zoonotic spillover: a mixed-methods study of the use of animal products in medicinal and cultural practices in Nigeria. One Health Outlook, 2022, 4, 5.	1.4	7
162	Identification by next-generation sequencing of Aichivirus B in a calf with enterocolitis and neurologic signs. Journal of Veterinary Diagnostic Investigation, 2017, 29, 208-211.	0.5	6

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