Tony L Goldberg

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9078159/publications.pdf

Version: 2024-02-01

191 papers 8,270 citations

50 h-index 79 g-index

201 all docs

201 docs citations

times ranked

201

9469 citing authors

#	Article	IF	CITATIONS
1	Viruses associated with ill health in wild chimpanzees. American Journal of Primatology, 2022, 84, e23358.	0.8	11
2	A Novel Gonadotropic Microsporidian Parasite (Microsporidium clinchi n. sp.) Infecting a Declining Population of Pheasantshell Mussels (Actinonaias pectorosa) (Unioinidae) from the Clinch River, USA. Parasitologia, 2022, 2, 1-12.	0.6	6
3	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
4	Genetics and community-based restoration can guide conservation of forest fragments for endangered primates. Perspectives in Ecology and Conservation, 2022, , .	1.0	0
5	Skin fungal assemblages of bats vary based on susceptibility to white-nose syndrome. ISME Journal, 2021, 15, 909-920.	4.4	16
6	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
7	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
8	A multi-omic investigation of male lower urinary tract symptoms: Potential role for JC virus. PLoS ONE, 2021, 16, e0246266.	1.1	7
9	A Sarcina bacterium linked to lethal disease in sanctuary chimpanzees in Sierra Leone. Nature Communications, 2021, 12, 763.	5.8	17
10	Widespread Seropositivity to Viral Hemorrhagic Septicemia Virus (VHSV) in Four Species of Inland Sport Fishes in Wisconsin. Journal of Aquatic Animal Health, 2021, 33, 53-65.	0.6	2
11	Nephroblastoma in a Common Mudpuppy <i>Necturus maculosus</i> simultaneously Present with a Mollicute Bacterium of the Genus <i>Acholeplasma</i> Journal of Aquatic Animal Health, 2021, 33, 44-52.	0.6	O
12	Development and Characterization of a cDNA-Launch Recombinant Simian Hemorrhagic Fever Virus Expressing Enhanced Green Fluorescent Protein: ORF 2b' Is Not Required for In Vitro Virus Replication. Viruses, 2021, 13, 632.	1.5	5
13	Bat Flies of the Family Streblidae (Diptera: Hippoboscoidea) Host Relatives of Medically and Agriculturally Important "Bat-Associated―Viruses. Viruses, 2021, 13, 860.	1.5	16
14	ICTV Virus Taxonomy Profile: Arteriviridae 2021. Journal of General Virology, 2021, 102, .	1.3	64
15	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
16	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
17	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
18	Typical intracranial myiasis in Nigerian red river hogs (Potamochoerus porcus) caused by an unknown bot fly (Diptera: Oestridae). International Journal for Parasitology: Parasites and Wildlife, 2021, 17, 14-19.	0.6	1

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19	Wisconsin dairy farm worker perceptions and practices related to antibiotic use, resistance, and infection prevention using a systems engineering framework. PLoS ONE, 2021, 16, e0258290.	1.1	2
20	Biodiversity of protists and nematodes in the wild nonhuman primate gut. ISME Journal, 2020, 14, 609-622.	4.4	32
21	Relatives of rubella virus in diverse mammals. Nature, 2020, 586, 424-428.	13.7	58
22	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
23	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
24	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
25	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
26	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
27	Discovery of Lanama Virus, a Distinct Member of Species Kunsagivirus C (Picornavirales:) Tj ETQq1 1 0.784314 r	gBŢ <i>ၙ[</i> Overl	ock 10 Tf 50
28	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
29	Bacterial communities on the gills of bonefish (Albula vulpes) in the Florida Keys and The Bahamas show spatial structure and differential abundance of disease-associated bacteria. Marine Biology, 2020, 167, 1.	0.7	6
30	Assessment of a Serologic Diagnostic Test and Kinetics of Antibody Development in Northern Pike Experimentally Infected with Viral Hemorrhagic Septicemia Virus. Journal of Aquatic Animal Health, 2020, 32, 3-10.	0.6	1
31	Eating Bushmeat Improves Food Security in a Biodiversity and Infectious Disease "Hotspot― EcoHealth, 2020, 17, 125-138.	0.9	40
32	Dynamically evolving novel overlapping gene as a factor in the SARS-CoV-2 pandemic. ELife, 2020, 9, .	2.8	74
33	One Health: Farmworker Perceptions of Antibiotic Resistance and Personal Protective Practices on Wisconsin Dairy Farms. Infection Control and Hospital Epidemiology, 2020, 41, s448-s449.	1.0	0
34	Title is missing!. , 2020, 16, e1008717.		0
35	Title is missing!. , 2020, 16, e1008717.		0
36	Title is missing!. , 2020, 16, e1008717.		0

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37	Title is missing!. , 2020, 16, e1008717.		O
38	Evolutionary trends in host physiology outweigh dietary niche in structuring primate gut microbiomes. ISME Journal, 2019, 13, 576-587.	4.4	236
39	Molecular Analysis of the Complete Genome of a Simian Foamy Virus Infecting Hylobates pileatus (pileated gibbon) Reveals Ancient Co-Evolution with Lesser Apes. Viruses, 2019, 11, 605.	1.5	2
40	Reference intervals for blood-based biochemical analytes of southern Beaufort Sea polar bears., 2019, 7, coz040.		4
41	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
42	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
43	The evolution of a super-swarm of foot-and-mouth disease virus in cattle. PLoS ONE, 2019, 14, e0210847.	1.1	14
44	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
45	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
46	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
47	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	0.9	153
48	Risk factors for respiratory illness in a community of wild chimpanzees (<i>Pan troglodytes) Tj ETQq0 0 0 rgBT</i>	/Overlock]	10 Tf 50 302 ⁻
49	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
50	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
51	Severe neurologic disease and chick mortality in crested screamers (Chauna torquata) infected with a novel Gyrovirus. Virology, 2018, 520, 111-115.	1.1	18
52	Co-circulation of Flanders Virus and West Nile Virus in Culex Mosquitoes (Diptera: Culicidae) from Chicago, Illinois. Journal of Medical Entomology, 2018, 55, 1062-1066.	0.9	3
53	Lethal Respiratory Disease Associated with Human RhinovirusÂC in Wild Chimpanzees, Uganda, 2013. Emerging Infectious Diseases, 2018, 24, 267-274.	2.0	80
54	Spatial configuration becomes more important with increasing habitat loss: a simulation study of environmentally-transmitted parasites. Landscape Ecology, 2018, 33, 1259-1272.	1.9	6

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55	Demography and health of "village dogs―in rural Western Uganda. Preventive Veterinary Medicine, 2017, 137, 24-27.	0.7	12
56	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
57	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
58	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
59	Within-Host Evolution of Simian Arteriviruses in Crab-Eating Macaques. Journal of Virology, 2017, 91, .	1.5	4
60	Metagenomic assessment of adventitious viruses in commercial bovine sera. Biologicals, 2017, 47, 64-68.	0.5	37
61	Spatial Overlap Between People and Non-human Primates in a Fragmented Landscape. EcoHealth, 2017, 14, 88-99.	0.9	11
62	Comparison of gastrointestinal parasite communities in vervet monkeys. Integrative Zoology, 2017, 12, 512-520.	1.3	10
63	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
64	Cis â€regulatory evolution in a wild primate: Infectionâ€associated genetic variation drives differential expression of MHC ―DQA 1 inÂvitro. Molecular Ecology, 2017, 26, 4523-4535.	2.0	3
65	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
66	Divergent Simian Arteriviruses Cause Simian Hemorrhagic Fever of Differing Severities in Macaques. MBio, 2016, 7, e02009-15.	1.8	14
67	Definitive Hosts of <i>Versteria </i> Tapeworms (Cestoda: Taeniidae) Causing Fatal Infection in North America. Emerging Infectious Diseases, 2016, 22, 707-710.	2.0	18
68	Naturally Circulating Hepatitis A Virus in Olive Baboons, Uganda. Emerging Infectious Diseases, 2016, 22, 1308-1310.	2.0	17
69	Assessing Commitment and Reporting Fidelity to a Text Message-Based Participatory Surveillance in Rural Western Uganda. PLoS ONE, 2016, 11, e0155971.	1.1	4
70	Arteriviruses, Pegiviruses, and Lentiviruses Are Common among Wild African Monkeys. Journal of Virology, 2016, 90, 6724-6737.	1.5	26
71	Safeguarding biodiversity: what is perceived as working, according to the conservation community?. Oryx, 2016, 50, 302-307.	0.5	12
72	Spatial patterns of persistence for environmentally transmitted parasites: Effects of regional climate and local landscape. Ecological Modelling, 2016, 338, 78-89.	1.2	4

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73	A Multicomponent Animal Virus Isolated from Mosquitoes. Cell Host and Microbe, 2016, 20, 357-367.	5.1	123
74	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
75	Changes in physiological stress and behaviour in semi-free-ranging red-capped mangabeys () Tj ETQq1 1 0.784314 Biological Sciences, 2016, 283, 20161201.	rgBT /Ove 1.2	erlock 10 Tf 13
76	Drivers for emerging issues in animal and plant health. EFSA Journal, 2016, 14, e00512.	0.9	17
77	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
78	Primate reinfection with gastrointestinal parasites: behavioural and physiological predictors of parasite acquisition. Animal Behaviour, 2016, 117, 105-113.	0.8	22
79	Zoonotic Potential of Simian Arteriviruses. Journal of Virology, 2016, 90, 630-635.	1.5	48
80	Rapid identification of major histocompatibility complex class I haplotypes using deep sequencing in an endangered Old World monkey. Conservation Genetics Resources, 2016, 8, 23-26.	0.4	4
81	Reorganization and expansion of the nidoviral family Arteriviridae. Archives of Virology, 2016, 161, 755-768.	0.9	254
82	Decreased Flight Activity in <i>Culex pipiens </i> (Diptera: Culicidae) Naturally Infected With Culex flavivirus. Journal of Medical Entomology, 2016, 53, 233-236.	0.9	21
83	Novel reovirus associated with epidemic mortality in wild largemouth bass (Micropterus salmoides). Journal of General Virology, 2016, 97, 2482-2487.	1.3	34
84	Specific Detection of Two Divergent Simian Arteriviruses Using RNAscope In Situ Hybridization. PLoS ONE, 2016, 11, e0151313.	1.1	7
85	Social Behaviours and Networks of Vervet Monkeys Are Influenced by Gastrointestinal Parasites. PLoS ONE, 2016, 11, e0161113.	1.1	50
86	Providing health care to improve community perceptions of protected areas. Oryx, 2015, 49, 636-642.	0.5	23
87	GB Virus C Coinfections in West African Ebola Patients. Journal of Virology, 2015, 89, 2425-2429.	1.5	65
88	Durable sequence stability and bone marrow tropism in a macaque model of human pegivirus infection. Science Translational Medicine, 2015, 7, 305ra144.	5.8	22
89	Temporal Variation in Viral Hemorrhagic Septicemia Virus Antibodies in Freshwater Drum (Aplodinotus grunniens) Indicates Cyclic Transmission in Lake Winnebago, Wisconsin. Journal of Clinical Microbiology, 2015, 53, 2889-2894.	1.8	4
90	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

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91	The need for a global health ethic. Lancet, The, 2015, 386, e37-e39.	6.3	58
92	Drivers of Bushmeat Hunting and Perceptions of Zoonoses in Nigerian Hunting Communities. PLoS Neglected Tropical Diseases, 2015, 9, e0003792.	1.3	79
93	Genomic Resources Notes Accepted 1 December 2014 - 31 January 2015. Molecular Ecology Resources, 2015, 15, 684-684.	2.2	2
94	Identification of Avian and Hemoparasite DNA in Blood-Engorged Abdomens of Culex pipiens (Diptera;) Tj ETQq0 (Entomology, 2015, 52, 461-468.	0 0 rgBT / 0.9	Overlock 10 11
95	Sickness behaviour associated with non-lethal infections in wild primates. Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20151436.	1.2	53
96	Historical Outbreaks of Simian Hemorrhagic Fever in Captive Macaques Were Caused by Distinct Arteriviruses. Journal of Virology, 2015, 89, 8082-8087.	1.5	21
97	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
98	WEST NILE VIRUS ANTIBODY DECAY RATE IN FREE-RANGING BIRDS. Journal of Wildlife Diseases, 2015, 51, 601.	0.3	12
99	Uncovering zoonoses awareness in an emerging disease †hotspot'. Social Science and Medicine, 2015, 129, 78-86.	1.8	25
100	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
101	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
102	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
103	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
104	Genome Sequences of Simian Hemorrhagic Fever Virus Variant NIH LVR42-0/M6941 Isolates (Arteriviridae: Arterivirus). Genome Announcements, 2014, 2, .	0.8	9
105	Fatal Metacestode Infection in Bornean Orangutan Caused by Unknown <i>Versteria</i> Species. Emerging Infectious Diseases, 2014, 20, 109-113.	2.0	21
106	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
107	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
108	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

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109	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
110	Two Novel Simian Arteriviruses in Captive and Wild Baboons (Papio spp.). Journal of Virology, 2014, 88, 13231-13239.	1.5	28
111	Host group formation decreases exposure to vector-borne disease: a field experiment in a †hotspot†of West Nile virus transmission. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20141586.	1.2	20
112	Beyond Bushmeat: Animal Contact, Injury, and Zoonotic Disease Risk in Western Uganda. EcoHealth, 2014, 11, 534-543.	0.9	54
113	Discovery and Characterization of Distinct Simian Pegiviruses in Three Wild African Old World Monkey Species. PLoS ONE, 2014, 9, e98569.	1.1	45
114	Deep sequencing identifies two genotypes and high viral genetic diversity of human pegivirus (GB virus) Tj ETQq0	0 <u>,0</u> rgBT /	Overlock 10
115	Is Markhamia lutea's abundance determined by animal foraging?. Forest Ecology and Management, 2013, 308, 62-66.	1.4	17
116	Physiological and Behavioral Effects of Capture Darting on Red Colobus Monkeys (Procolobus) Tj ETQq0 0 0 rgBT of Primatology, 2013, 34, 1020-1031.	/Overlock 0.9	10 Tf 50 46 23
117	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
118	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
119	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
120	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
121	Exceptional Simian Hemorrhagic Fever Virus Diversity in a Wild African Primate Community. Journal of Virology, 2013, 87, 688-691.	1.5	61
122	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
123	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
124	Going, Going, Gone: A 15-Year History of the Decline of Primates in Forest Fragments near Kibale National Park, Uganda., 2013,, 89-100.		77
125	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
126	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

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127	Diversity of <i>Wolbachia pipientis </i> Strain <i>Woli>Pip in a Genetically Admixtured, Above-Ground <i>Culex pipiens </i>(Diptera: Culicidae) Population: Association With Form Molestus Ancestry and Host Selection Patterns. Journal of Medical Entomology, 2012, 49, 474-481.</i>	0.9	7
128	Wild Birds and Urban Ecology of Ticks and Tick-borne Pathogens, Chicago, Illinois, USA, 2005–2010. Emerging Infectious Diseases, 2012, 18, 1589-1595.	2.0	86
129	Protozoan parasites in groupâ€living primates: testing the biological island hypothesis. American Journal of Primatology, 2012, 74, 510-517.	0.8	18
130	An outbreak of the 2009 influenza a (H1N1) virus in a children's hospital. Influenza and Other Respiratory Viruses, 2012, 6, 374-379.	1.5	19
131	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
132	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
133	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
134	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
135	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
136	Genetic and antigenic relationships of vesicular stomatitis viruses from South America. Archives of Virology, 2011, 156, 1961-1968.	0.9	19
137	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
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	Detection of Known and Novel Adenoviruses in Cattle Wastes via Broad-Spectrum Primers. Applied and Environmental Microbiology, 2011, 77, 5001-5008.	1.4	27
140	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
141	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
142	West Nile virus may have hitched a ride across the Western United States on <i>Culex tarsalis </i>	2.0	13
143	Characterization of the Fecal Microbiome from Non-Human Wild Primates Reveals Species Specific Microbial Communities. PLoS ONE, 2010, 5, e13963.	1.1	225
144	Multi-year evolutionary dynamics of West Nile virus in suburban Chicago, USA, 2005–2007. Philosophical Transactions of the Royal Society B: Biological Sciences, 2010, 365, 1871-1878.	1.8	33

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145	Molecular Epidemiology of Cross-Species Giardia duodenalis Transmission in Western Uganda. PLoS Neglected Tropical Diseases, 2010, 4, e683.	1.3	136
146	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
147	Coinfection of Ugandan Red Colobus (<i>Procolobus </i> [<i>Piliocolobus </i>] <i>rufomitratus) Tj ETQq1 1 0.78</i>	84314 rgE 1.5	BT /Overlock 82
148	Avian host community structure and prevalence of West Nile virus in Chicago, Illinois. Oecologia, 2009, 159, 415-424.	0.9	82
149	Biodiversity Loss Affects Global Disease Ecology. BioScience, 2009, 59, 945-954.	2.2	211
150	Genetic Variation Associated with Mammalian Feeding in <i>Culex pipiens</i> from a West Nile Virus Epidemic Region in Chicago, Illinois. Vector-Borne and Zoonotic Diseases, 2009, 9, 637-642.	0.6	65
151	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
152	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
153	Commentary on "Pandemic Human Viruses Cause Decline of Endangered Great Apes,―by Köndgen et al., 2008, Current Biology 18: 260–264. American Journal of Primatology, 2008, 70, 716-718.	0.8	2
154	Gastrointestinal Bacterial Transmission among Humans, Mountain Gorillas, and Livestock in Bwindi Impenetrable National Park, Uganda. Conservation Biology, 2008, 22, 1600-1607.	2.4	183
155	Fine-scale genetic variation and evolution of West Nile Virus in a transmission "hot spot―in suburban Chicago, USA. Virology, 2008, 374, 381-389.	1.1	64
156	Rapid Amplification of West Nile Virus: The Role of Hatch-Year Birds. Vector-Borne and Zoonotic Diseases, 2008, 8, 57-68.	0.6	97
157	High Rates of <i>Escherichia coli</i> Journal of Clinical Microbiology, 2008, 46, 3187-3191.	1.8	56
158	<i>Culex pipiens</i> (Diptera: Culicidae): A Bridge Vector of West Nile Virus to Humans. Journal of Medical Entomology, 2008, 45, 125-128.	0.9	227
159	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
160	Health and disease in the people, primates, and domestic animals of Kibale National Park: implications for conservation., 2008,, 75-87.		55
161	Forest Fragmentation as Cause of Bacterial Transmission among Nonhuman Primates, Humans, and Livestock, Uganda. Emerging Infectious Diseases, 2008, 14, 1375-1382.	2.0	145
162	Serologic Evidence for Novel Poxvirus in Endangered Red Colobus Monkeys, Western Uganda. Emerging Infectious Diseases, 2008, 14, 801-803.	2.0	23

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