Tony L Goldberg

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

 191
 6,077
 45
 70

 papers
 h-index
 g-index

 201
 7,160
 4.8
 5.62

 ext. papers
 ext. citations
 avg, IF
 L-index

#	Paper	IF	Citations
191	Viruses associated with ill health in wild chimpanzees <i>American Journal of Primatology</i> , 2022 , e23358	2.5	1
190	A Novel Gonadotropic Microsporidian Parasite (Microsporidium clinchi n. sp.) Infecting a Declining Population of Pheasantshell Mussels (Actinonaias pectorosa) (Unioinidae) from the Clinch River, USA. <i>Parasitologia</i> , 2022 , 2, 1-12		2
189	Typical intracranial myiasis in Nigerian red river hogs () caused by an unknown bot fly (Diptera: Oestridae) <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2022 , 17, 14-19	2.6	1
188	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 <i>One Health Outlook</i> , 2022 , 4, 5	5	1
187	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. <i>Viruses</i> , 2021 , 13,	6.2	1
186	Bat Flies of the Family (Diptera: Hippoboscoidea) Host Relatives of Medically and Agriculturally Important "Bat-Associated" Viruses. <i>Viruses</i> , 2021 , 13,	6.2	2
185	Skin fungal assemblages of bats vary based on susceptibility to white-nose syndrome. <i>ISME Journal</i> , 2021 , 15, 909-920	11.9	5
184	Age Patterning in Wild Chimpanzee Gut Microbiota Diversity Reveals Differences from Humans in Early Life. <i>Current Biology</i> , 2021 , 31, 613-620.e3	6.3	11
183	Opportunities for respiratory disease transmission from people to chimpanzees at an East African tourism site. <i>American Journal of Primatology</i> , 2021 , 83, e23228	2.5	6
182	A multi-omic investigation of male lower urinary tract symptoms: Potential role for JC virus. <i>PLoS ONE</i> , 2021 , 16, e0246266	3.7	1
181	A Sarcina bacterium linked to lethal disease in sanctuary chimpanzees in Sierra Leone. <i>Nature Communications</i> , 2021 , 12, 763	17.4	5
180	Widespread Seropositivity to Viral Hemorrhagic Septicemia Virus (VHSV) in Four Species of Inland Sport Fishes in Wisconsin. <i>Journal of Aquatic Animal Health</i> , 2021 , 33, 53-65	2.6	
179	Nephroblastoma in a Common Mudpuppy Necturus maculosus simultaneously Present with a Mollicute Bacterium of the Genus Acholeplasma. <i>Journal of Aquatic Animal Health</i> , 2021 , 33, 44-52	2.6	
178	ICTV Virus Taxonomy Profile: 2021. <i>Journal of General Virology</i> , 2021 , 102,	4.9	12
177	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021 , 166, 3513-3566	2.6	10
176	Mycobiome Traits Associated with Disease Tolerance Predict Many Western North American Bat Species Will Be Susceptible to White-Nose Syndrome. <i>Microbiology Spectrum</i> , 2021 , 9, e0025421	8.9	1
175	Mussel Mass Mortality and the Microbiome: Evidence for Shifts in the Bacterial Microbiome of a Declining Freshwater Bivalve. <i>Microorganisms</i> , 2021 , 9,	4.9	5

(2020-2021)

174	Wisconsin dairy farm worker perceptions and practices related to antibiotic use, resistance, and infection prevention using a systems engineering framework <i>PLoS ONE</i> , 2021 , 16, e0258290	3.7	0
173	Discovery of Lanama Virus, a Distinct Member of Species (:), in Wild Vervet Monkeys (). <i>Viruses</i> , 2020 , 12,	6.2	1
172	Diversity, Transmission, and Cophylogeny of Ledanteviruses (:) and Nycteribiid Bat Flies Parasitizing Angolan Soft-Furred Fruit Bats in Bundibugyo District, Uganda. <i>Microorganisms</i> , 2020 , 8,	4.9	6
171	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. <i>Marine Biology</i> , 2020 , 167, 1	2.5	1
170	Assessment of a Serologic Diagnostic Test and Kinetics of Antibody Development in Northern Pike Experimentally Infected with Viral Hemorrhagic Septicemia Virus. <i>Journal of Aquatic Animal Health</i> , 2020 , 32, 3-10	2.6	1
169	Eating Bushmeat Improves Food Security in a Biodiversity and Infectious Disease "Hotspot". <i>EcoHealth</i> , 2020 , 17, 125-138	3.1	27
168	One Health: Farmworker Perceptions of Antibiotic Resistance and Personal Protective Practices on Wisconsin Dairy Farms. <i>Infection Control and Hospital Epidemiology</i> , 2020 , 41, s448-s449	2	
167	Dynamically evolving novel overlapping gene as a factor in the SARS-CoV-2 pandemic. <i>ELife</i> , 2020 , 9,	8.9	35
166	Use of a systems engineering framework to assess perceptions and practices about antimicrobial resistance of workers on large dairy farms in Wisconsin 2020 ,		2
165	Biodiversity of protists and nematodes in the wild nonhuman primate gut. ISME Journal, 2020, 14, 609-6	5 22 .9	16
165 164	Biodiversity of protists and nematodes in the wild nonhuman primate gut. <i>ISME Journal</i> , 2020 , 14, 609-60. Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428	50.4	16 27
164	Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428 Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat () in Uganda Dramatically Expands the	50.4	27
164	Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428 Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat () in Uganda Dramatically Expands the Global Distribution of an Emerging Bat-Borne Respiratory Virus. <i>Viruses</i> , 2020 , 12, Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in	50.4	27
164 163 162	Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428 Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat () in Uganda Dramatically Expands the Global Distribution of an Emerging Bat-Borne Respiratory Virus. <i>Viruses</i> , 2020 , 12, Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in Hepatocystis parasites from naturally infected red colobus monkeys. <i>PLoS Pathogens</i> , 2020 , 16, e10087 Mass mortality in freshwater mussels (Actinonaias pectorosa) in the Clinch River, USA, linked to a	50.4 6.2	27 3 11
164 163 162	Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428 Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat () in Uganda Dramatically Expands the Global Distribution of an Emerging Bat-Borne Respiratory Virus. <i>Viruses</i> , 2020 , 12, Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in Hepatocystis parasites from naturally infected red colobus monkeys. <i>PLoS Pathogens</i> , 2020 , 16, e10087 Mass mortality in freshwater mussels (Actinonaias pectorosa) in the Clinch River, USA, linked to a novel densovirus. <i>Scientific Reports</i> , 2020 , 10, 14498 Demography, life-history trade-offs, and the gastrointestinal virome of wild chimpanzees.	50.4 6.2 17 ⁶ 4.9	27 3 11
164 163 162 161	Relatives of rubella virus in diverse mammals. <i>Nature</i> , 2020 , 586, 424-428 Pteropine Orthoreovirus in an Angolan Soft-Furred Fruit Bat () in Uganda Dramatically Expands the Global Distribution of an Emerging Bat-Borne Respiratory Virus. <i>Viruses</i> , 2020 , 12, Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in Hepatocystis parasites from naturally infected red colobus monkeys. <i>PLoS Pathogens</i> , 2020 , 16, e10087 Mass mortality in freshwater mussels (Actinonaias pectorosa) in the Clinch River, USA, linked to a novel densovirus. <i>Scientific Reports</i> , 2020 , 10, 14498 Demography, life-history trade-offs, and the gastrointestinal virome of wild chimpanzees. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020 , 375, 20190613 Faecal parasites increase with age but not reproductive effort in wild female chimpanzees.	50.4 6.2 17 ⁶ 4.9 5.8	27 3 11 11 8

Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in 156 Hepatocystis parasites from naturally infected red colobus monkeys 2020, 16, e1008717 Genomic and transcriptomic evidence for descent from Plasmodium and loss of blood schizogony in 155 Hepatocystis parasites from naturally infected red colobus monkeys **2020**, 16, e1008717 Reference intervals for blood-based biochemical analytes of southern Beaufort Sea polar bears 154 1 2019, 7, coz040 Simultaneous outbreaks of respiratory disease in wild chimpanzees caused by distinct viruses of 18.9 153 44 human origin. Emerging Microbes and Infections, 2019, 8, 139-149 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 152 3.9 5 Evolution, 2019, 11, 1630-1643 The evolution of a super-swarm of foot-and-mouth disease virus in cattle. PLoS ONE, 2019, 14, e02108473.7 151 13 Evolutionary trends in host physiology outweigh dietary niche in structuring primate gut 150 11.9 132 microbiomes. ISME Journal, 2019, 13, 576-587 Molecular Analysis of the Complete Genome of a Simian Foamy Virus Infecting (pileated gibbon) 6.2 149 Reveals Ancient Co-Evolution with Lesser Apes. Viruses, 2019, 11, Life on the Rainforest Edge: Food Security in the Agricultural-Forest Frontier of Cross River State, 148 4.8 10 Nigeria. Frontiers in Sustainable Food Systems, 2019, 3, Multidecade Mortality and a Homolog of Hepatitis C Virus in Bald Eagles (Haliaeetus 147 4.9 9 leucocephalus), the National Bird of the USA. Scientific Reports, 2019, 9, 14953 Diverse RNA viruses of arthropod origin in the blood of fruit bats suggest a link between bat and 146 3.6 21 arthropod viromes. Virology, 2019, 528, 64-72 Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294 2.6 145 111 Co-circulation of Flanders Virus and West Nile Virus in Culex Mosquitoes (Diptera: Culicidae) from 144 2.2 2 Chicago, Illinois. Journal of Medical Entomology, 2018, 55, 1062-1066 Lethal Respiratory Disease Associated with Human Rhinovirus C in Wild Chimpanzees, Uganda, 143 57 2013. Emerging Infectious Diseases, **2018**, 24, 267-274 Spatial configuration becomes more important with increasing habitat loss: a simulation study of 142 4.3 2 environmentally-transmitted parasites. Landscape Ecology, 2018, 33, 1259-1272 Risk factors for respiratory illness in a community of wild chimpanzees (). Royal Society Open Science 141 3.3 22 , **2018**, 5, 180840 Selective constraint and adaptive potential of West Nile virus within and among naturally infected 140 3.7 12 avian hosts and mosquito vectors. Virus Evolution, 2018, 4, vey013 Antibiotic-Resistant Escherichia coli and Class 1 Integrons in Humans, Domestic Animals, and Wild 4.8 18 139 Primates in Rural Uganda. Applied and Environmental Microbiology, 2018, 84,

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138	Severe neurologic disease and chick mortality in crested screamers (Chauna torquata) infected with a novel Gyrovirus. <i>Virology</i> , 2018 , 520, 111-115	3.6	13	
137	Demography and health of "village dogs" in rural Western Uganda. <i>Preventive Veterinary Medicine</i> , 2017 , 137, 24-27	3.1	6	
136	Combining Footwear with Public Health Iconography to Prevent Soil-Transmitted Helminth Infections. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017 , 96, 205-213	3.2	10	
135	Identification by next-generation sequencing of Aichivirus B in a calf with enterocolitis and neurologic signs. <i>Journal of Veterinary Diagnostic Investigation</i> , 2017 , 29, 208-211	1.5	6	
134	Culex Flavivirus During West Nile Virus Epidemic and Interepidemic Years in Chicago, United States. <i>Vector-Borne and Zoonotic Diseases</i> , 2017 , 17, 567-575	2.4	12	
133	Within-Host Evolution of Simian Arteriviruses in Crab-Eating Macaques. <i>Journal of Virology</i> , 2017 , 91,	6.6	2	
132	Metagenomic assessment of adventitious viruses in commercial bovine sera. <i>Biologicals</i> , 2017 , 47, 64-68	1.8	23	
131	Spatial Overlap Between People and Non-human Primates in a Fragmented Landscape. <i>EcoHealth</i> , 2017 , 14, 88-99	3.1	6	
130	Comparison of gastrointestinal parasite communities in vervet monkeys. <i>Integrative Zoology</i> , 2017 , 12, 512-520	1.9	8	
129	Kanyawara Virus: A Novel Rhabdovirus Infecting Newly Discovered Nycteribiid Bat Flies Infesting Previously Unknown Pteropodid Bats in Uganda. <i>Scientific Reports</i> , 2017 , 7, 5287	4.9	22	
128	Cis-regulatory evolution in a wild primate: Infection-associated genetic variation drives differential expression of MHC-DQA1 in vitro. <i>Molecular Ecology</i> , 2017 , 26, 4523-4535	5.7	1	
127	Prevalence and Risk Factors Associated with Hemoparasites in Cattle and Goats at the Edge of Kibale National Park, Western Uganda. <i>Journal of Parasitology</i> , 2017 , 103, 69-74	0.9	15	
126	A Multicomponent Animal Virus Isolated from Mosquitoes. <i>Cell Host and Microbe</i> , 2016 , 20, 357-367	23.4	76	
125	Overlap in the Seasonal Infection Patterns of Avian Malaria Parasites and West Nile Virus in Vectors and Hosts. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016 , 95, 1121-1129	3.2	8	
124	Changes in physiological stress and behaviour in semi-free-ranging red-capped mangabeys (Cercocebus torquatus) following antiparasitic treatment. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016 , 283,	4.4	10	
123	Drivers for emerging issues in animal and plant health. <i>EFSA Journal</i> , 2016 , 14, e00512	2.3	13	
122	Predicting West Nile Virus Infection Risk From the Synergistic Effects of Rainfall and Temperature. Journal of Medical Entomology, 2016 , 53, 935-944	2.2	27	
121	Primate reinfection with gastrointestinal parasites: behavioural and physiological predictors of parasite acquisition. <i>Animal Behaviour</i> , 2016 , 117, 105-113	2.8	14	

120	Zoonotic Potential of Simian Arteriviruses. <i>Journal of Virology</i> , 2016 , 90, 630-5	6.6	35
119	Rapid Identification of Major Histocompatibility Complex Class I Haplotypes Using Deep Sequencing in an Endangered Old World Monkey. <i>Conservation Genetics Resources</i> , 2016 , 8, 23-26	0.8	3
118	Reorganization and expansion of the nidoviral family Arteriviridae. Archives of Virology, 2016, 161, 755	- 68 .6	167
117	Decreased Flight Activity in Culex pipiens (Diptera: Culicidae) Naturally Infected With Culex flavivirus. <i>Journal of Medical Entomology</i> , 2016 , 53, 233-6	2.2	14
116	Specific Detection of Two Divergent Simian Arteriviruses Using RNAscope In Situ Hybridization. <i>PLoS ONE</i> , 2016 , 11, e0151313	3.7	4
115	Social Behaviours and Networks of Vervet Monkeys Are Influenced by Gastrointestinal Parasites. <i>PLoS ONE</i> , 2016 , 11, e0161113	3.7	43
114	Novel reovirus associated with epidemic mortality in wild largemouth bass (Micropterus salmoides). Journal of General Virology, 2016 , 97, 2482-2487	4.9	23
113	Divergent Simian Arteriviruses Cause Simian Hemorrhagic Fever of Differing Severities in Macaques. <i>MBio</i> , 2016 , 7, e02009-15	7.8	9
112	Definitive Hosts of Versteria Tapeworms (Cestoda: Taeniidae) Causing Fatal Infection in North America. <i>Emerging Infectious Diseases</i> , 2016 , 22, 707-10	10.2	13
111	Naturally Circulating Hepatitis A Virus in Olive Baboons, Uganda. <i>Emerging Infectious Diseases</i> , 2016 , 22, 1308-10	10.2	9
110	Assessing Commitment and Reporting Fidelity to a Text Message-Based Participatory Surveillance in Rural Western Uganda. <i>PLoS ONE</i> , 2016 , 11, e0155971	3.7	4
109	Arteriviruses, Pegiviruses, and Lentiviruses Are Common among Wild African Monkeys. <i>Journal of Virology</i> , 2016 , 90, 6724-6737	6.6	19
108	Safeguarding biodiversity: what is perceived as working, according to the conservation community?. <i>Oryx</i> , 2016 , 50, 302-307	1.5	11
107	Spatial patterns of persistence for environmentally transmitted parasites: Effects of regional climate and local landscape. <i>Ecological Modelling</i> , 2016 , 338, 78-89	3	3
106	The need for a global health ethic. <i>Lancet, The</i> , 2015 , 386, e37-e39	40	52
105	Drivers of bushmeat hunting and perceptions of zoonoses in Nigerian hunting communities. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0003792	4.8	55
104	Genomic resources notes accepted 1 December 2014 - 31 January 2015. <i>Molecular Ecology Resources</i> , 2015 , 15, 684	8.4	2
103	Identification of Avian and Hemoparasite DNA in Blood-Engorged Abdomens of Culex pipiens (Diptera; Culicidae) from a West Nile Virus Epidemic region in Suburban Chicago, Illinois. <i>Journal of Medical Entomology</i> , 2015 , 52, 461-8	2.2	10

(2014-2015)

102	Sickness behaviour associated with non-lethal infections in wild primates. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015 , 282,	4.4	38	
101	Historical Outbreaks of Simian Hemorrhagic Fever in Captive Macaques Were Caused by Distinct Arteriviruses. <i>Journal of Virology</i> , 2015 , 89, 8082-7	6.6	15	
100	Lung fluke (Paragonimus africanus) infects Nigerian red-capped mangabeys and causes respiratory disease. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2015 , 4, 329-32	2.6	7	
99	WEST NILE VIRUS ANTIBODY DECAY RATE IN FREE-RANGING BIRDS. <i>Journal of Wildlife Diseases</i> , 2015 , 51, 601-8	1.3	9	
98	Uncovering zoonoses awareness in an emerging disease 'hotspot'. <i>Social Science and Medicine</i> , 2015 , 129, 78-86	5.1	19	
97	Providing health care to improve community perceptions of protected areas. <i>Oryx</i> , 2015 , 49, 636-642	1.5	18	
96	GB virus C coinfections in west African Ebola patients. <i>Journal of Virology</i> , 2015 , 89, 2425-9	6.6	47	
95	Durable sequence stability and bone marrow tropism in a macaque model of human pegivirus infection. <i>Science Translational Medicine</i> , 2015 , 7, 305ra144	17.5	18	
94	Temporal Variation in Viral Hemorrhagic Septicemia Virus Antibodies in Freshwater Drum (Aplodinotus grunniens) Indicates Cyclic Transmission in Lake Winnebago, Wisconsin. <i>Journal of Clinical Microbiology</i> , 2015 , 53, 2889-94	9.7	4	
93	Simian hemorrhagic fever virus cell entry is dependent on CD163 and uses a clathrin-mediated endocytosis-like pathway. <i>Journal of Virology</i> , 2015 , 89, 844-56	6.6	30	
92	Fecal microbiomes of non-human primates in Western Uganda reveal species-specific communities largely resistant to habitat perturbation. <i>American Journal of Primatology</i> , 2014 , 76, 347-54	2.5	56	
91	Broad protection against avian influenza virus by using a modified vaccinia Ankara virus expressing a mosaic hemagglutinin gene. <i>Journal of Virology</i> , 2014 , 88, 13300-9	6.6	34	
90	Two novel simian arteriviruses in captive and wild baboons (Papio spp.). <i>Journal of Virology</i> , 2014 , 88, 13231-9	6.6	23	
89	Host group formation decreases exposure to vector-borne disease: a field experiment in a 'hotspot' of West Nile virus transmission. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014 , 281, 2014	15 8 6	16	
88	Beyond bushmeat: animal contact, injury, and zoonotic disease risk in Western Uganda. <i>EcoHealth</i> , 2014 , 11, 534-43	3.1	43	
87	High genetic diversity and adaptive potential of two simian hemorrhagic fever viruses in a wild primate population. <i>PLoS ONE</i> , 2014 , 9, e90714	3.7	29	
86	Dispersal of adult culex mosquitoes in an urban west nile virus hotspot: a mark-capture study incorporating stable isotope enrichment of natural larval habitats. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e2768	4.8	37	
85	Nodule worm infection in humans and wild primates in Uganda: cryptic species in a newly identified region of human transmission. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e2641	4.8	59	

84	Hidden population structure and cross-species transmission of whipworms (Trichuris sp.) in humans and non-human primates in Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e3256	4.8	56
83	Genome Sequences of Simian Hemorrhagic Fever Virus Variant NIH LVR42-0/M6941 Isolates (Arteriviridae: Arterivirus). <i>Genome Announcements</i> , 2014 , 2,		8
82	Fatal metacestode infection in Bornean orangutan caused by unknown Versteria species. <i>Emerging Infectious Diseases</i> , 2014 , 20, 109-13	10.2	14
81	Development and evaluation of a blocking enzyme-linked immunosorbent assay and virus neutralization assay to detect antibodies to viral hemorrhagic septicemia virus. <i>Vaccine Journal</i> , 2014 , 21, 435-42		12
80	Discovery and full genome characterization of a new SIV lineage infecting red-tailed guenons (Cercopithecus ascanius schmidti) in Kibale National Park, Uganda. <i>Retrovirology</i> , 2014 , 11, 55	3.6	10
79	Discovery and characterization of distinct simian pegiviruses in three wild African Old World monkey species. <i>PLoS ONE</i> , 2014 , 9, e98569	3.7	38
78	Deep sequencing identifies two genotypes and high viral genetic diversity of human pegivirus (GB virus C) in rural Ugandan patients. <i>Journal of General Virology</i> , 2013 , 94, 2670-2678	4.9	13
77	Is 's abundance determined by animal foraging?. Forest Ecology and Management, 2013 , 308, 62-62	3.9	13
76	Physiological and Behavioral Effects of Capture Darting on Red Colobus Monkeys (Procolobus rufomitratus) with a Comparison to Chimpanzee (Pan troglodytes) Predation. <i>International Journal of Primatology</i> , 2013 , 34, 1020-1031	2	15
75	Movement patterns of bonefish (Albula vulpes) in tidal creeks and coastal waters of Eleuthera, The Bahamas. <i>Fisheries Research</i> , 2013 , 147, 404-412	2.3	44
74	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. <i>Retrovirology</i> , 2013 , 10, 107	3.6	28
73	Prevalence of filarioid nematodes and trypanosomes in American robins and house sparrows, Chicago USA. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2013 , 2, 42-9	2.6	21
72	Co-infection and cross-species transmission of divergent Hepatocystis lineages in a wild African primate community. <i>International Journal for Parasitology</i> , 2013 , 43, 613-9	4.3	25
71	Exceptional simian hemorrhagic fever virus diversity in a wild African primate community. <i>Journal of Virology</i> , 2013 , 87, 688-91	6.6	51
70	A novel hepacivirus with an unusually long and intrinsically disordered NS5A protein in a wild Old World primate. <i>Journal of Virology</i> , 2013 , 87, 8971-81	6.6	73
69	Coincident tick infestations in the nostrils of wild chimpanzees and a human in Uganda. <i>American Journal of Tropical Medicine and Hygiene</i> , 2013 , 89, 924-7	3.2	10
68	Going, Going, Gone: A 15-Year History of the Decline of Primates in Forest Fragments near Kibale National Park, Uganda 2013 , 89-100		68
67	An outbreak of the 2009 influenza a (H1N1) virus in a children's hospital. <i>Influenza and Other Respiratory Viruses</i> , 2012 , 6, 374-9	5.6	19

(2010-2012)

66	Wild birds and urban ecology of ticks and tick-borne pathogens, Chicago, Illinois, USA, 2005-2010. <i>Emerging Infectious Diseases</i> , 2012 , 18, 1589-95	10.2	71
65	Protozoan parasites in group-living primates: testing the biological island hypothesis. <i>American Journal of Primatology</i> , 2012 , 74, 510-7	2.5	13
64	Epidemiology and molecular relationships of Cryptosporidium spp. in people, primates, and livestock from Western Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1597	4.8	56
63	Evaluation of a stable isotope method to mark naturally-breeding larval mosquitoes for adult dispersal studies. <i>Journal of Medical Entomology</i> , 2012 , 49, 61-70	2.2	32
62	Diversity of Wolbachia pipientis strain wPip in a genetically admixtured, above-ground Culex pipiens (Diptera: Culicidae) population: association with form molestus ancestry and host selection patterns. <i>Journal of Medical Entomology</i> , 2012 , 49, 474-81	2.2	7
61	Culex flavivirus and West Nile virus mosquito coinfection and positive ecological association in Chicago, United States. <i>Vector-Borne and Zoonotic Diseases</i> , 2011 , 11, 1099-105	2.4	90
60	Novel, divergent simian hemorrhagic fever viruses in a wild Ugandan red colobus monkey discovered using direct pyrosequencing. <i>PLoS ONE</i> , 2011 , 6, e19056	3.7	55
59	Fine-scale variation in vector host use and force of infection drive localized patterns of West Nile virus transmission. <i>PLoS ONE</i> , 2011 , 6, e23767	3.7	94
58	Impacts of dissolved oxygen on the behavior and physiology of bonefish: Implications for live-release angling tournaments. <i>Journal of Experimental Marine Biology and Ecology</i> , 2011 , 402, 19-26	2.1	28
57	Genetic and antigenic relationships of vesicular stomatitis viruses from South America. <i>Archives of Virology</i> , 2011 , 156, 1961-8	2.6	16
56	Spatial ecology and residency patterns of adult great barracuda (Sphyraena barracuda) in coastal waters of The Bahamas. <i>Marine Biology</i> , 2011 , 158, 2227-2237	2.5	41
55	Evaluation of a novel emergence trap to study Culex mosquitoes in urban catch basins. <i>Journal of the American Mosquito Control Association</i> , 2011 , 27, 142-7	0.9	17
54	Field investigation of innate immunity in passerine birds in suburban Chicago, Illinois, USA. <i>Journal of Wildlife Diseases</i> , 2011 , 47, 603-11	1.3	11
53	Detection of known and novel adenoviruses in cattle wastes via broad-spectrum primers. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 5001-8	4.8	20
52	West Nile virus may have hitched a ride across the Western United States on Culex tarsalis mosquitoes. <i>Molecular Ecology</i> , 2010 , 19, 1518-9	5.7	9
51	Characterization of the fecal microbiome from non-human wild primates reveals species specific microbial communities. <i>PLoS ONE</i> , 2010 , 5, e13963	3.7	169
50	Multi-year evolutionary dynamics of West Nile virus in suburban Chicago, USA, 2005-2007. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2010 , 365, 1871-8	5.8	30
49	Molecular epidemiology of cross-species Giardia duodenalis transmission in western Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e683	4.8	101

48	Local impact of temperature and precipitation on West Nile virus infection in Culex species mosquitoes in northeast Illinois, USA. <i>Parasites and Vectors</i> , 2010 , 3, 19	4	174
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