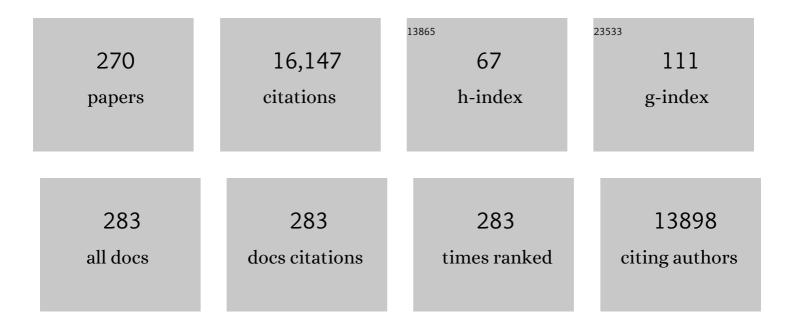
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Field evaluation of specific mycobacterial proteinâ€based skin test for the differentiation of <i>Mycobacterium bovisâ€</i> infected and Bacillus Calmette Guerinâ€vaccinated crossbred cattle in Ethiopia. Transboundary and Emerging Diseases, 2022, 69, .	3.0	6
2	HAM-ART: An optimised culture-free Hi-C metagenomics pipeline for tracking antimicrobial resistance genes in complex microbial communities. PLoS Genetics, 2022, 18, e1009776.	3.5	14
3	Productivity loss and cost of bovine tuberculosis for the dairy livestock sector in Ethiopia. Preventive Veterinary Medicine, 2022, 202, 105616.	1.9	4
4	Understanding the relative risks of zoonosis emergence under contrasting approaches to meeting livestock product demand. Royal Society Open Science, 2022, 9, .	2.4	9
5	A case of early neonate bovine tuberculosis in Ethiopia. Clinical Case Reports (discontinued), 2021, 9, 487-490.	0.5	2
6	Dynamics and risk of transmission of bovine tuberculosis in the emerging dairy regions of Ethiopia. Epidemiology and Infection, 2021, 149, e69.	2.1	1
7	A Meta-Analysis of the Effect of Bacillus Calmette-Guérin Vaccination Against Bovine Tuberculosis: Is Perfect the Enemy of Good?. Frontiers in Veterinary Science, 2021, 8, 637580.	2.2	19
8	Population structure and transmission of Mycobacterium bovis in Ethiopia. Microbial Genomics, 2021, 7, .	2.0	9
9	Global prevalence of <i>Mycobacterium bovis</i> infections among human tuberculosis cases: Systematic review and metaâ€analysis. Zoonoses and Public Health, 2021, 68, 704-718.	2.2	16
10	Epidemiology of Mycobacterium tuberculosis lineages and strain clustering within urban and peri-urban settings in Ethiopia. PLoS ONE, 2021, 16, e0253480.	2.5	5
11	Effect of Bovine Tuberculosis on Selected Productivity Parameters and Trading in Dairy Cattle Kept Under Intensive Husbandry in Central Ethiopia. Frontiers in Veterinary Science, 2021, 8, 698768.	2.2	4
12	The variable prevalence of bovine tuberculosis among dairy herds in Central Ethiopia provides opportunities for targeted intervention. PLoS ONE, 2021, 16, e0254091.	2.5	9
13	Evaluation of the Efficacy of BCG in Protecting Against Contact Challenge With Bovine Tuberculosis in Holstein-Friesian and Zebu Crossbred Calves in Ethiopia. Frontiers in Veterinary Science, 2021, 8, 702402.	2.2	11
14	Persistence of Multiple Paramyxoviruses in a Closed Captive Colony of Fruit Bats (Eidolon helvum). Viruses, 2021, 13, 1659.	3.3	6
15	Longitudinal Secretion of Paramyxovirus RNA in the Urine of Straw-Coloured Fruit Bats (Eidolon) Tj ETQq1 1 0.	784314 rgE	3T /Qverlock 1
16	A Longitudinal Study of Cattle Productivity in Intensive Dairy Farms in Central Ethiopia. Frontiers in Veterinary Science, 2021, 8, 698760.	2.2	10
17	80 questions for UK biological security. PLoS ONE, 2021, 16, e0241190.	2.5	8
18	Inferring Mycobacterium bovis transmission between cattle and badgers using isolates from the Randomised Badger Culling Trial. PLoS Pathogens, 2021, 17, e1010075.	4.7	20

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19	Disaster displacement and zoonotic disease dynamics: The impact of structural and chronic drivers in Sindh, Pakistan. PLOS Global Public Health, 2021, 1, e0000068.	1.6	5
20	Viral CpG Deficiency Provides No Evidence That Dogs Were Intermediate Hosts for SARS-CoV-2. Molecular Biology and Evolution, 2020, 37, 2706-2710.	8.9	18
21	Genotype Diversity of Mycobacterium bovis and Pathology of Bovine Tuberculosis in Selected Emerging Dairy Regions of Ethiopia. Frontiers in Veterinary Science, 2020, 7, 553940.	2.2	4
22	Achimota Pararubulavirus 3: A New Bat-Derived Paramyxovirus of the Genus Pararubulavirus. Viruses, 2020, 12, 1236.	3.3	6
23	Long-term follow-up of owned, free-roaming dogs in South Africa naturally exposed to Babesia rossi. International Journal for Parasitology, 2020, 50, 103-110.	3.1	4
24	Experimental Lagos bat virus infection in straw-colored fruit bats: A suitable model for bat rabies in a natural reservoir species. PLoS Neglected Tropical Diseases, 2020, 14, e0008898.	3.0	8
25	Network analysis of dairy cattle movement and associations with bovine tuberculosis spread and control in emerging dairy belts of Ethiopia. BMC Veterinary Research, 2019, 15, 262.	1.9	23
26	Demodicosis in a captive African straw-coloured fruit bat (Eidolon helvum). Experimental and Applied Acarology, 2019, 78, 547-554.	1.6	3
27	Impacts of environmental and socio-economic factors on emergence and epidemic potential of Ebola in Africa. Nature Communications, 2019, 10, 4531.	12.8	63
28	What is stirring in the reservoir? Modelling mechanisms of henipavirus circulation in fruit bat hosts. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20190021.	4.0	29
29	Estimating undetected Ebola spillovers. PLoS Neglected Tropical Diseases, 2019, 13, e0007428.	3.0	22
30	New methodologies for the estimation of population vulnerability to diseases: a case study of Lassa fever and Ebola in Nigeria and Sierra Leone. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20180265.	4.0	5
31	Seasonal variation in food availability and relative importance of dietary items in the Gambian epauletted fruit bat (<i>Epomophorus gambianus</i>). Ecology and Evolution, 2019, 9, 5683-5693.	1.9	5
32	Prevalence of bovine tuberculosis and its associated risk factors in the emerging dairy belts of regional cities in Ethiopia. Preventive Veterinary Medicine, 2019, 168, 81-89.	1.9	42
33	Disentangling serology to elucidate henipa―and filovirus transmission in Madagascar fruit bats. Journal of Animal Ecology, 2019, 88, 1001-1016.	2.8	36
34	Policy and Science for Global Health Security: Shaping the Course of International Health. Tropical Medicine and Infectious Disease, 2019, 4, 60.	2.3	12
35	Support for viral persistence in bats from age-specific serology and models of maternal immunity. Scientific Reports, 2018, 8, 3859.	3.3	37
36	Domesticated animals as hosts of henipaviruses and filoviruses: A systematic review. Veterinary Journal, 2018, 233, 25-34.	1.7	32

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37	Whole Genome Sequencing for Determining the Source of Mycobacterium bovis Infections in Livestock Herds and Wildlife in New Zealand. Frontiers in Veterinary Science, 2018, 5, 272.	2.2	44
38	Risk factors and variations in detection of new bovine tuberculosis breakdowns via slaughterhouse surveillance in Great Britain. PLoS ONE, 2018, 13, e0198760.	2.5	10
39	The Gambian epauletted fruit bat shows increased genetic divergence in the Ethiopian highlands and in an area of rapid urbanization. Ecology and Evolution, 2018, 8, 12803-12820.	1.9	6
40	Brucellosis in the Addis Ababa dairy cattle: the myths and the realities. BMC Veterinary Research, 2018, 14, 396.	1.9	18
41	Environmental limits of Rift Valley fever revealed using ecoepidemiological mechanistic models. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E7448-E7456.	7.1	19
42	Animal infection studies of two recently discovered African bat paramyxoviruses, Achimota 1 and Achimota 2. Scientific Reports, 2018, 8, 12744.	3.3	9
43	Maternal antibody and the maintenance of a lyssavirus in populations of seasonally breeding African bats. PLoS ONE, 2018, 13, e0198563.	2.5	16
44	Pathogenesis of bat rabies in a natural reservoir: Comparative susceptibility of the straw-colored fruit bat (Eidolon helvum) to three strains of Lagos bat virus. PLoS Neglected Tropical Diseases, 2018, 12, e0006311.	3.0	21
45	The intractable challenge of evaluating cattle vaccination as a control for bovine Tuberculosis. ELife, 2018, 7, .	6.0	8
46	How Does Africa's Most Hunted Bat Vary Across the Continent? Population Traits of the Straw-Coloured Fruit Bat (Eidolon helvum) and Its Interactions with Humans. Acta Chiropterologica, 2017, 19, 77.	0.6	23
47	One Health for a changing world: new perspectives from Africa. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160162.	4.0	45
48	Integrative modelling for One Health: pattern, process and participation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160164.	4.0	43
49	One Health, emerging infectious diseases and wildlife: two decades of progress?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160167.	4.0	334
50	Seasonality in oestrus and litter size in an assistance dog breeding colony in the United Kingdom. Veterinary Record, 2017, 181, 371-371.	0.3	8
51	Mathematical Models for the Epidemiology and Evolution of Mycobacterium tuberculosis. Advances in Experimental Medicine and Biology, 2017, 1019, 281-307.	1.6	1
52	Lagos Bat Virus Infection Dynamics in Free-Ranging Straw-Colored Fruit Bats (Eidolon helvum). Tropical Medicine and Infectious Disease, 2017, 2, 25.	2.3	16
53	A11 Phylogenetic and phylogeographic analysis of viral surveillance data to inform rabies control programmes in Cambodia. Virus Evolution, 2017, 3, .	4.9	0
54	Facility-based surveillance for emerging infectious diseases; diagnostic practices in rural West African hospital settings: observations from Ghana. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160544.	4.0	14

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55	The equine Hendra virus vaccine remains a highly effective preventative measure against infection in horses and humans: †The imperative to develop a human vaccine for the Hendra virus in Australia'. Infection Ecology and Epidemiology, 2016, 6, 31658.	0.8	6
56	Exposure to Bat-Associated <i>Bartonella</i> spp. among Humans and Other Animals, Chana. Emerging Infectious Diseases, 2016, 22, 922-924.	4.3	15
57	Transmission or Within-Host Dynamics Driving Pulses of Zoonotic Viruses in Reservoir–Host Populations. PLoS Neglected Tropical Diseases, 2016, 10, e0004796.	3.0	152
58	High turnover drives prolonged persistence of influenza in managed pig herds. Journal of the Royal Society Interface, 2016, 13, 20160138.	3.4	33
59	Environmentalâ€mechanistic modelling of the impact of global change on human zoonotic disease emergence: a case study of Lassa fever. Methods in Ecology and Evolution, 2016, 7, 646-655.	5.2	60
60	Bat trait, genetic and pathogen data from large-scale investigations of African fruit bats, Eidolon helvum. Scientific Data, 2016, 3, 160049.	5.3	9
61	Testing the dairy difference. Veterinary Journal, 2016, 217, 134-135.	1.7	2
62	The complete mitochondrial genome of <i>Epomophorus gambianus</i> (Chiroptera: Pteropodidae) and its phylogenetic analysis. Mitochondrial DNA Part B: Resources, 2016, 1, 447-449.	0.4	1
63	Increased risk of A(H1N1)pdm09 influenza infection in UK pig industry workers compared to a general population cohort. Influenza and Other Respiratory Viruses, 2016, 10, 291-300.	3.4	18
64	Ebola, Bats and Evidence-Based Policy. EcoHealth, 2016, 13, 9-11.	2.0	7
65	A Unified Framework for the Infection Dynamics of Zoonotic Spillover and Spread. PLoS Neglected Tropical Diseases, 2016, 10, e0004957.	3.0	52
66	Network analysis of host–virus communities in bats and rodents reveals determinants of crossâ€species transmission. Ecology Letters, 2015, 18, 1153-1162.	6.4	120
67	Bayesian Model Choice in Cumulative Link Ordinal Regression Models. Bayesian Analysis, 2015, 10, .	3.0	15
68	Eliminating bovine tuberculosis in cattle and badgers: insight from a dynamic model. Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20150374.	2.6	28
69	Can insecticide-treated netting provide protection for Equids from Culicoides biting midges in the United Kingdom?. Parasites and Vectors, 2015, 8, 604.	2.5	19
70	Clinical research: developing an appropriate career structure. Veterinary Record, 2015, 177, 544-547.	0.3	0
71	Characteristics and Risk Perceptions of Ghanaians Potentially Exposed to Bat-Borne Zoonoses through Bushmeat. EcoHealth, 2015, 12, 104-120.	2.0	76
72	Effective vaccination against rabies in puppies in rabies endemic regions. Veterinary Record, 2015, 177, 150-150.	0.3	32

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73	Potential Benefits of Cattle Vaccination as a Supplementary Control for Bovine Tuberculosis. PLoS Computational Biology, 2015, 11, e1004038.	3.2	28
74	Using Modelling to Disentangle the Relative Contributions of Zoonotic and Anthroponotic Transmission: The Case of Lassa Fever. PLoS Neglected Tropical Diseases, 2015, 9, e3398.	3.0	96
75	Seven challenges in modeling pathogen dynamics within-host and across scales. Epidemics, 2015, 10, 45-48.	3.0	79
76	Eight challenges in modelling infectious livestock diseases. Epidemics, 2015, 10, 1-5.	3.0	72
77	Nine challenges in modelling the emergence of novel pathogens. Epidemics, 2015, 10, 35-39.	3.0	60
78	Specificity of the comparative skin test for bovine tuberculosis in Great Britain. Veterinary Record, 2015, 177, 258-258.	0.3	69
79	Achieving Population-Level Immunity to Rabies in Free-Roaming Dogs in Africa and Asia. PLoS Neglected Tropical Diseases, 2014, 8, e3160.	3.0	45
80	Viral antibody dynamics in a chiropteran host. Journal of Animal Ecology, 2014, 83, 415-428.	2.8	43
81	The demography of freeâ€roaming dog populations and applications to disease and population control. Journal of Applied Ecology, 2014, 51, 1096-1106.	4.0	101
82	The effect of seasonal birth pulses on pathogen persistence in wild mammal populations. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20132962.	2.6	85
83	Antigenic variation of foot-and-mouth disease virus serotype A. Journal of General Virology, 2014, 95, 384-392.	2.9	42
84	Participatory methods for the assessment of the ownership status of free-roaming dogs in Bali, Indonesia, for disease control and animal welfare. Preventive Veterinary Medicine, 2014, 116, 203-208.	1.9	28
85	Prevalence and Risk Factors of Feather Plucking in African Grey Parrots (Psittacus erithacus) Tj ETQq1 1 0.78431 Medicine, 2014, 23, 250-257.	.4 rgBT /O 0.4	verlock 10 Tf 30
86	Laboratory animal models to study foot-and-mouth disease: a review with emphasis on natural and vaccine-induced immunity. Journal of General Virology, 2014, 95, 2329-2345.	2.9	41
87	Infectious risk factors and clinical indicators for tracheal mucus in <scp>B</scp> ritish <scp>N</scp> ational <scp>H</scp> unt racehorses. Equine Veterinary Journal, 2014, 46, 150-155.	1.7	22
88	5. Emerging risks from bat bushmeat in West Africa. , 2014, , 91-106.		0
89	Bat Flight and Zoonotic Viruses. Emerging Infectious Diseases, 2014, 20, 741-745.	4.3	269
90	Deciphering Serology to Understand the Ecology of Infectious Diseases in Wildlife. EcoHealth, 2013, 10, 298-313.	2.0	156

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91	Use of cross-reactive serological assays for detecting novel pathogens in wildlife: Assessing an appropriate cutoff for henipavirus assays in African bats. Journal of Virological Methods, 2013, 193, 295-303.	2.1	50
92	Continent-wide panmixia of an African fruit bat facilitates transmission of potentially zoonotic viruses. Nature Communications, 2013, 4, 2770.	12.8	105
93	Age-dependent patterns of bovine tuberculosis in cattle. Veterinary Research, 2013, 44, 97.	3.0	57
94	Evidenceâ€based control of canine rabies: a critical review of population density reduction. Journal of Animal Ecology, 2013, 82, 6-14.	2.8	163
95	Ecology of Zoonotic Infectious Diseases in Bats: Current Knowledge and Future Directions. Zoonoses and Public Health, 2013, 60, 2-21.	2.2	150
96	Exercise affects joint injury risk in young Thoroughbreds in training. Veterinary Journal, 2013, 196, 339-344.	1.7	16
97	A comparison of bats and rodents as reservoirs of zoonotic viruses: are bats special?. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20122753.	2.6	508
98	The evolutionary dynamics of influenza A virus adaptation to mammalian hosts. Philosophical Transactions of the Royal Society B: Biological Sciences, 2013, 368, 20120382.	4.0	40
99	Metagenomic study of the viruses of African straw-coloured fruit bats: Detection of a chiropteran poxvirus and isolation of a novel adenovirus. Virology, 2013, 441, 95-106.	2.4	121
100	Novel, Potentially Zoonotic Paramyxoviruses from the African Straw-Colored Fruit Bat Eidolon helvum. Journal of Virology, 2013, 87, 1348-1358.	3.4	75
101	Evolution of Equine Influenza Virus in Vaccinated Horses. Journal of Virology, 2013, 87, 4768-4771.	3.4	34
102	A longitudinal study of respiratory infections in British National Hunt racehorses. Veterinary Record, 2013, 172, 637-637.	0.3	15
103	A restatement of the natural science evidence base relevant to the control of bovine tuberculosis in Great Britain ^{â€} . Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20131634.	2.6	118
104	Where are the horses? With the sheep or cows? Uncertain host location, vector-feeding preferences and the risk of African horse sickness transmission in Great Britain. Journal of the Royal Society Interface, 2013, 10, 20130194.	3.4	42
105	What can mathematical models bring to the control of equine influenza?. Equine Veterinary Journal, 2013, 45, 784-788.	1.7	18
106	The accuracy of the National Equine Database in relation to vector-borne disease risk modelling of horses in Great Britain. Equine Veterinary Journal, 2013, 45, 302-308.	1.7	11
107	Inferring the inter-host transmission of influenza A virus using patterns of intra-host genetic variation. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20122173.	2.6	45
108	Transmission of Equine Influenza Virus during an Outbreak Is Characterized by Frequent Mixed Infections and Loose Transmission Bottlenecks. PLoS Pathogens, 2012, 8, e1003081.	4.7	57

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109	Estimating the Hidden Burden of Bovine Tuberculosis in Great Britain. PLoS Computational Biology, 2012, 8, e1002730.	3.2	117
110	Evolution of an Eurasian Avian-like Influenza Virus in NaÃ ⁻ ve and Vaccinated Pigs. PLoS Pathogens, 2012, 8, e1002730.	4.7	79
111	Dynamics of <i>Salmonella</i> infection of macrophages at the single cell level. Journal of the Royal Society Interface, 2012, 9, 2696-2707.	3.4	70
112	Descriptive clinical and epidemiological characteristics of influenza A H1N1 2009 virus infections in pigs in England. Veterinary Record, 2012, 171, 271-271.	0.3	22
113	<i>Bartonella</i> species in bat flies (Diptera: Nycteribiidae) from western Africa. Parasitology, 2012, 139, 324-329.	1.5	82
114	Descriptive epidemiology of joint injuries in Thoroughbred racehorses in training. Equine Veterinary Journal, 2012, 44, 13-19.	1.7	49
115	Endemic Lagos bat virus infection inEidolon helvum. Epidemiology and Infection, 2012, 140, 2163-2171.	2.1	41
116	Demography of straw-colored fruit bats in Ghana. Journal of Mammalogy, 2012, 93, 1393-1404.	1.3	66
117	Estimating reassortment rates in co-circulating Eurasian swine influenza viruses. Journal of General Virology, 2012, 93, 2326-2336.	2.9	42
118	Estimation of the Relative Sensitivity of the Comparative Tuberculin Skin Test in Tuberculous Cattle Herds Subjected to Depopulation. PLoS ONE, 2012, 7, e43217.	2.5	39
119	A framework for the study of zoonotic disease emergence and its drivers: spillover of bat pathogens as a case study. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 2881-2892.	4.0	156
120	Factors Associated with Pleurisy in Pigs: A Case-Control Analysis of Slaughter Pig Data for England and Wales. PLoS ONE, 2012, 7, e29655.	2.5	31
121	Henipavirus Neutralising Antibodies in an Isolated Island Population of African Fruit Bats. PLoS ONE, 2012, 7, e30346.	2.5	71
122	Quantifying Transmission of Highly Pathogenic and Low Pathogenicity H7N1 Avian Influenza in Turkeys. PLoS ONE, 2012, 7, e45059.	2.5	34
123	The Effect of Badger Culling on Breakdown Prolongation and Recurrence of Bovine Tuberculosis in Cattle Herds in Great Britain. PLoS ONE, 2012, 7, e51342.	2.5	4
124	Co-circulation of diverse paramyxoviruses in an urban African fruit bat population. Journal of General Virology, 2012, 93, 850-856.	2.9	60
125	Participatory epidemiology: Approaches, methods, experiences. Veterinary Journal, 2012, 191, 151-160.	1.7	187
126	Effects of essential fatty acid supplementation in dogs with idiopathic epilepsy: A clinical trial. Veterinary Journal, 2012, 191, 396-398.	1.7	18

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127	Modelâ€guided fieldwork: practical guidelines for multidisciplinary research on wildlife ecological and epidemiological dynamics. Ecology Letters, 2012, 15, 1083-1094.	6.4	131
128	Ebola Virus Antibodies in Fruit Bats, Ghana, West Africa. Emerging Infectious Diseases, 2012, 18, 1207-1209.	4.3	126
129	Uncovering the fruit bat bushmeat commodity chain and the true extent of fruit bat hunting in Ghana, West Africa. Biological Conservation, 2011, 144, 3000-3008.	4.1	139
130	Making use of equine population demography for disease control purposes: Preliminary observations on the difficulties of counting and locating horses in Great Britain. Equine Veterinary Journal, 2011, 43, 372-375.	1.7	22
131	Descriptive results from a longitudinal study of airway inflammation in British National Hunt racehorses. Equine Veterinary Journal, 2011, 43, 750-755.	1.7	28
132	What's in a name? Inflammatory airway disease in racehorses in training. Equine Veterinary Journal, 2011, 43, 756-758.	1.7	12
133	Antibodies to Henipavirus or Henipa-Like Viruses in Domestic Pigs in Ghana, West Africa. PLoS ONE, 2011, 6, e25256.	2.5	72
134	Experimental transmission of avianâ€like swine H1N1 influenza virus between immunologically naÃ⁻ve and vaccinated pigs. Influenza and Other Respiratory Viruses, 2011, 5, 357-364.	3.4	6
135	Recurrence of bovine tuberculosis breakdowns in Great Britain: Risk factors and prediction. Preventive Veterinary Medicine, 2011, 102, 22-29.	1.9	94
136	A universal real-time assay for the detection of Lyssaviruses. Journal of Virological Methods, 2011, 177, 87-93.	2.1	76
137	Evidence for several waves of global transmission in the seventh cholera pandemic. Nature, 2011, 477, 462-465.	27.8	649
138	Diagnostic assays for glanders. Veterinary Record, 2011, 169, 663-663.	0.3	0
139	Genome-Scale Evolution and Phylodynamics of Equine H3N8 Influenza A Virus. Journal of Virology, 2011, 85, 5312-5322.	3.4	90
140	National Hunt Racehorse, Point to Point Horse, and Timber Racing Horse. , 2011, , 1062-1075.		0
141	Antigenic and Genetic Evolution of Equine Influenza A (H3N8) Virus from 1968 to 2007. Journal of Virology, 2011, 85, 12742-12749.	3.4	89
142	Bluetongue serotype 8 vaccine coverage in northern and southâ€eastern England in 2008. Veterinary Record, 2011, 168, 428-428.	0.3	6
143	A Bayesian Approach to Analyse Genetic Variation within RNA Viral Populations. PLoS Computational Biology, 2011, 7, e1002027.	3.2	8
144	Prevalence and risk factors for swine influenza virus infection in the English pig population. PLOS Currents, 2011, 3, RRN1209.	1.4	36

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145	Effect of age and training on murmurs of atrioventricular valvular regurgitation in young Thoroughbreds. Equine Veterinary Journal, 2010, 32, 195-199.	1.7	56
146	Control of strangles outbreaks by isolation of guttural pouch carriers identified using PCR and culture of Streptococcus equi. Equine Veterinary Journal, 2010, 32, 515-526.	1.7	121
147	Coughing: a source of irritation?. Equine Veterinary Journal, 2010, 33, 221-223.	1.7	1
148	A case-control study of respiratory disease in Thoroughbred racehorses in Sydney, Australia. Equine Veterinary Journal, 2010, 33, 256-264.	1.7	89
149	Racehorse injuries, clinical problems and fatalities recorded on British racecourses from flat racing and National Hunt racing during 1996, 1997 and 1998. Equine Veterinary Journal, 2010, 33, 478-486.	1.7	308
150	It's all in the mix: infection transmission in populations. Equine Veterinary Journal, 2010, 35, 526-528.	1.7	1
151	Gender differences in bone turnover in 2-year-old Thoroughbreds. Equine Veterinary Journal, 2010, 35, 702-706.	1.7	19
152	Is isoflurane safer than halothane in equine anaesthesia? Results from a prospective multicentre randomised controlled trial. Equine Veterinary Journal, 2010, 36, 64-71.	1.7	101
153	An epidemiological study of risk factors associated with the recurrence of equine grass sickness (dysautonomia) on previously affected premises. Equine Veterinary Journal, 2010, 36, 105-112.	1.7	43
154	Descriptive epidemiology of fractures occurring in British Thoroughbred racehorses in training. Equine Veterinary Journal, 2010, 36, 167-173.	1.7	112
155	Fractures and tendon injuries in National Hunt horses in training in the UK: a pilot study. Equine Veterinary Journal, 2010, 36, 365-367.	1.7	48
156	End-to-end jejuno-ileal anastomosis following resection of strangulated small intestine in horses: a comparative study. Equine Veterinary Journal, 2010, 37, 356-359.	1.7	26
157	Inflammatory airway disease, nasal discharge and respiratory infections in young British racehorses. Equine Veterinary Journal, 2010, 37, 236-242.	1.7	94
158	Risk factors for epistaxis on British racecourses: evidence for locomotory impact-induced trauma contributing to the aetiology of exercise-induced pulmonary haemorrhage. Equine Veterinary Journal, 2010, 37, 402-411.	1.7	30
159	Training-related factors associated with dorsometacarpal disease in young Thoroughbred racehorses in the UK. Equine Veterinary Journal, 2010, 37, 442-448.	1.7	45
160	Factors associated with failure of Thoroughbred horses to train and race. Equine Veterinary Journal, 2010, 38, 113-118.	1.7	133
161	Characterization of microsatellite loci in the straw-colored fruit bat, Eidolon helvum (Pteropodidae). Conservation Genetics Resources, 2010, 2, 279-282.	0.8	4
162	Predicting prolonged bovine tuberculosis breakdowns in Great Britain as an aid to control. Preventive Veterinary Medicine, 2010, 97, 183-190.	1.9	47

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163	Molecular epidemiology of Streptococcus zooepidemicus isolated from the respiratory tracts of Thoroughbred racehorses in training. Veterinary Journal, 2010, 183, 348-351.	1.7	12
164	Virus neutralising activity of African fruit bat (Eidolon helvum) sera against emerging lyssaviruses. Virology, 2010, 408, 183-189.	2.4	53
165	Methods and mortality results of a health survey of purebred dogs in the UK. Journal of Small Animal Practice, 2010, 51, 512-524.	1.2	195
166	An ex vivo swine tracheal organ culture for the study of influenza infection. Influenza and Other Respiratory Viruses, 2010, 4, 7-15.	3.4	29
167	The Spread of Bluetongue Virus Serotype 8 in Great Britain and Its Control by Vaccination. PLoS ONE, 2010, 5, e9353.	2.5	60
168	Intra- and Interhost Evolutionary Dynamics of Equine Influenza Virus. Journal of Virology, 2010, 84, 6943-6954.	3.4	97
169	Control of equine influenza: scenario testing using a realistic metapopulation model of spread. Journal of the Royal Society Interface, 2010, 7, 67-79.	3.4	54
170	Quantifying Antigenic Relationships among the Lyssaviruses. Journal of Virology, 2010, 84, 11841-11848.	3.4	83
171	Intrahost Evolutionary Dynamics of Canine Influenza Virus in Nail̂^ve and Partially Immune Dogs. Journal of Virology, 2010, 84, 5329-5335.	3.4	61
172	Use of a novel serological test for exposure to <i>Streptococcus equi</i> subspecies <i>equi</i> in hospitalised horses. Veterinary Record, 2010, 166, 294-297.	0.3	12
173	Dynamics of Influenza Virus Infection and Pathology. Journal of Virology, 2010, 84, 3974-3983.	3.4	172
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