Jonas WaldenstrÃ¶m

List of Publications by Year
in descending order

161 papers	11,021 citations	44069 48 h-index	32842 100 g-index
167 all docs	167 docs citations	167 times ranked	8847 citing authors

#	Article	IF	Citations
1	Global Patterns of Influenza A Virus in Wild Birds. Science, 2006, 312, 384-388.	12.6	1,619
2	A NEW PCR ASSAY FOR SIMULTANEOUS STUDIES OF LEUCOCYTOZOON, PLASMODIUM, AND HAEMOPROTEUS FROM AVIAN BLOOD. Journal of Parasitology, 2004, 90, 797-802.	0.7	812
3	Spatial, Temporal, and Species Variation in Prevalence of Influenza A Viruses in Wild Migratory Birds. PLoS Pathogens, 2007, 3, e61.	4.7	591
4	A New Nested Polymerase Chain Reaction Method Very Efficient in Detecting Plasmodium and Haemoproteus Infections From Avian Blood. Journal of Parasitology, 2004, 90, 191-194.	0.7	418
5	Rapid Advance of Spring Arrival Dates in Long-Distance Migratory Birds. Science, 2006, 312, 1959-1961.	12.6	399
6	Cross-species infection of blood parasites between resident and migratory songbirds in Africa. Molecular Ecology, 2002, 11, 1545-1554.	3.9	348
7	LINKAGE BETWEEN NUCLEAR AND MITOCHONDRIAL DNA SEQUENCES IN AVIAN MALARIA PARASITES: MULTIPLE CASES OF CRYPTIC SPECIATION?. Evolution; International Journal of Organic Evolution, 2004, 58, 1617-1621.	2.3	271
8	With Reference to Reference Genes: A Systematic Review of Endogenous Controls in Gene Expression Studies. PLoS ONE, 2015, 10, e0141853.	2.5	236
9	Prevalence of Campylobacter jejuni, Campylobacter lari, and Campylobacter coli in Different Ecological Guilds and Taxa of Migrating Birds. Applied and Environmental Microbiology, 2002, 68, 5911-5917.	3.1	233
10	Temporal dynamics and diversity of avian malaria parasites in a single host species. Journal of Animal Ecology, 2007, 76, 112-122.	2.8	218
11	Surveillance of Influenza Virus A in Migratory Waterfowl in Northern Europe. Emerging Infectious Diseases, 2007, 13, 404-411.	4.3	214
12	Dissemination of Escherichia coli with CTX-M Type ESBL between Humans and Yellow-Legged Gulls in the South of France. PLoS ONE, 2009, 4, e5958.	2.5	190
13	Detecting shifts of transmission areas in avian blood parasites - a phylogenetic approach. Molecular Ecology, 2007, 16, 1281-1290.	3.9	183
14	Effects of influenza A virus infection on migrating mallard ducks. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 1029-1036.	2.6	174
15	Associations between malaria and MHC genes in a migratory songbird. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 1511-1518.	2.6	172
16	Migrating Birds and Tickborne Encephalitis Virus. Emerging Infectious Diseases, 2007, 13, 1215-1218.	4.3	151
17	Influenza Virus in a Natural Host, the Mallard: Experimental Infection Data. PLoS ONE, 2010, 5, e8935.	2.5	130
18	Protozoan Acanthamoeba polyphaga as a Potential Reservoir for Campylobacter jejuni. Applied and Environmental Microbiology, 2005, 71, 987-992.	3.1	123

#	Article	IF	Citations
19	Dissemination of Spotted Fever Rickettsia Agents in Europe by Migrating Birds. PLoS ONE, 2010, 5, e8572.	2.5	120
20	Flying with the wind: scale dependency of speed and direction measurements in modelling wind support in avian flight. Movement Ecology, 2013, 1, 4.	2.8	111
21	Transatlantic spread of highly pathogenic avian influenza H5N1 by wild birds from Europe to North America in 2021. Scientific Reports, 2022, 12, .	3.3	106
22	Campylobacter jejuni in Black-Headed Gulls (Larus ridibundus): Prevalence, Genotypes, and Influence on C. jejuni Epidemiology. Journal of Clinical Microbiology, 2002, 40, 4594-4602.	3.9	104
23	The Ecology of Emerging Infectious Diseases in Migratory Birds: An Assessment of the Role of Climate Change and Priorities for Future Research. EcoHealth, 2012, 9, 80-88.	2.0	104
24	Within-Host Speciation of Malaria Parasites. PLoS ONE, 2007, 2, e235.	2.5	103
25	Long-term variation in influenza A virus prevalence and subtype diversity in migratory mallards in northern Europe. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20140098.	2.6	103
26	Marked host specificity and lack of phylogeographic population structure of Campylobacter jejuni in wild birds. Molecular Ecology, 2013, 22, 1463-1472.	3.9	96
27	Carriage of CTX-M type extended spectrum Î²-lactamases (ESBLs) in gulls across Europe. Acta Veterinaria Scandinavica, 2015, 57, 74.	1.6	87
28	Migratory Birds, Ticks, and Crimean-Congo Hemorrhagic Fever Virus. Emerging Infectious Diseases, 2012, 18, 2095-2097.	4.3	83
29	Antiviral Oseltamivir Is not Removed or Degraded in Normal Sewage Water Treatment: Implications for Development of Resistance by Influenza A Virus. PLoS ONE, 2007, 2, e986.	2.5	83
30	Antimicrobial Drugâ€“ResistantEscherichia coliin Wild Birds and Free-range Poultry, Bangladesh. Emerging Infectious Diseases, 2012, 18, 2055-2058.	4.3	75
31	What are malaria parasites?. Trends in Parasitology, 2005, 21, 209-211.	3.3	74
32	Haemosporidian Blood Parasites in European Birds of Prey and Owls. Journal of Parasitology, 2008, 94, 709-715.	0.7	74
33	Diversities and similarities in PFGE profiles of Campylobacter jejuni isolated from migrating birds and humans. Journal of Applied Microbiology, 2004, 96, 834-843.	3.1	72
34	Heterosubtypic Immunity to Influenza A Virus Infections in Mallards May Explain Existence of Multiple Virus Subtypes. PLoS Pathogens, 2013, 9, e1003443.	4.7	70
35	virF -Positive Yersinia pseudotuberculosis and Yersinia enterocolitica Found in Migratory Birds in Sweden. Applied and Environmental Microbiology, 2003, 69, 4670-4675.	3.1	69
36	Species diversity of campylobacteria in a wild bird community in Sweden. Journal of Applied Microbiology, 2007, 102, 424-32.	3.1	64

#	Article	IF	Citations
37	Sampling for low-pathogenic avian influenza A virus in wild Mallard ducks: Oropharyngeal versus cloacal swabbing. Vaccine, 2008, 26, 4414-4416.	3.8	62
38	Individual Variation in Influenza A Virus Infection Histories and Long-Term Immune Responses in Mallards. PLoS ONE, 2013, 8, e61201.	2.5	62
39	Prevalence and Phylogeny of Coronaviruses in Wild Birds from the Bering Strait Area (Beringia). PLoS ONE, 2010, 5, e13640.	2.5	61
40	Wild birdâ€associated <scp>C</scp>ampylobacter jejuni isolates are a consistent source of human disease, in <scp>O</scp>xfordshire, <scp>U</scp>nited <scp>K</scp>ingdom. Environmental Microbiology Reports, 2015, 7, 782-788.	2.4	61
41	Host and virus ecology as determinants of influenza A virus transmission in wild birds. Current Opinion in Virology, 2018, 28, 26-36.	5.4	58
42	Human-Associated Extended-Spectrum Î²-Lactamase in the Antarctic. Applied and Environmental Microbiology, 2012, 78, 2056-2058.	3.1	57
43	Extended-Spectrum Î²-Lactamases inEscherichia coli and Klebsiellapneumoniaein Gulls, Alaska, USA. Emerging Infectious Diseases, 2014, 20, 897-9.	4.3	57
44	Gene Segment Reassortment Between American and Asian Lineages of Avian Influenza Virus from Waterfowl in the Beringia Area. Vector-Borne and Zoonotic Diseases, 2008, 8, 783-790.	1.5	54
45	High Prevalence of Antibiotic Resistance in Pathogenic Escherichia coli from Large- and Small-Scale Poultry Farms in Bangladesh. Avian Diseases, 2011, 55, 689-692.	1.0	54
46	Environmental Levels of the Antiviral Oseltamivir Induce Development of Resistance Mutation H274Y in Influenza A/H1N1 Virus in Mallards. PLoS ONE, 2011, 6, e24742.	2.5	54
47	Frequency and patterns of reassortment in natural influenza A virus infection in a reservoir host. Virology, 2013, 443, 150-160.	2.4	54
48	Potential disease transmission from wild geese and swans to livestock, poultry and humans: a review of the scientific literature from a One Health perspective. Infection Ecology and Epidemiology, 2017, 7, 1300450.	0.8	54
49	Disease Dynamics and Bird Migrationâ€"Linking Mallards Anas platyrhynchos and Subtype Diversity of the Influenza A Virus in Time and Space. PLoS ONE, 2012, 7, e35679.	2.5	53
50	Movements, Home-Range Size and Habitat Selection of Mallards during Autumn Migration. PLoS ONE, 2014, 9, e100764.	2.5	52
51	Campylobacter jejuni Colonization in Wild Birds: Results from an Infection Experiment. PLoS ONE, 2010, 5, e9082.	2.5	52
52	Prevalence of Campylobacter in Wild Birds of the Mid-Atlantic Region, USA. Journal of Wildlife Diseases, 2011, 47, 750-754.	0.8	51
53	Biological Earth observation with animal sensors. Trends in Ecology and Evolution, 2022, 37, 293-298.	8.7	49
54	Garden Warbler Sylvia borin migration in sub-Saharan West Africa: phenology and body mass changes. Ibis, 2005, 147, 750-757.	1.9	48

#	Article	IF	Citations
55	Temporal patterns of occurrence and transmission of the blood parasite Haemoproteus payevskyi in the great reed warbler Acrocephalus arundinaceus. Journal of Ornithology, 2007, 148, 401-409.	1.1	48
56	Comparison of Extended-Spectrum Î²-Lactamase (ESBL) CTX-M Genotypes in Franklin Gulls from Canada and Chile. PLoS ONE, 2015, 10, e0141315.	2.5	45
57	Avian Reservoirs and Zoonotic Potential of theEmerging Human Pathogen Helicobactercanadensis. Applied and Environmental Microbiology, 2003, 69, 7523-7526.	3.1	43
58	Salmonellain Birds Migrating through Sweden. Emerging Infectious Diseases, 2003, 9, 753-755.	4.3	42
59	Antibiotic resistance patterns in Escherichia coli from gulls in nine European countries. Infection Ecology and Epidemiology, 2014, 4, 21565.	0.8	42
60	Transient Expression of Hemagglutinin Antigen from Low Pathogenic Avian Influenza A (H7N7) in Nicotiana benthamiana. PLoS ONE, 2012, 7, e33010.	2.5	41
61	Multilocus sequence typing of Campylobacter jejuni from broilers. Veterinary Microbiology, 2010, 140, 180-185.	1.9	38
62	The Evolution of Innate Immune Genes: Purifying and Balancing Selection on Î²-Defensins in Waterfowl. Molecular Biology and Evolution, 2016, 33, 3075-3087.	8.9	38
63	Migratory birds as disseminators of ticks and the tick-borne pathogens Borrelia bacteria and tick-borne encephalitis (TBE) virus: a seasonal study at Ottenby Bird Observatory in South-eastern Sweden. Parasites and Vectors, 2020, 13, 607.	2.5	38
64	Mounting evidence for the presence of influenza A virus in the avifauna of the Antarctic region. Antarctic Science, 2006, 18, 353-356.	0.9	36
65	Isotope signatures in winter moulted feathers predict malaria prevalence in a breeding avian host. Oecologia, 2008, 158, 299-306.	2.0	36
66	Circannual variation in blood parasitism in a subâ€Saharan migrant passerine bird, the garden warbler. Journal of Evolutionary Biology, 2013, 26, 1047-1059.	1.7	36
67	Direct and indirect effects of winter harshness on the survival of Mallards Anas platyrhynchos in northwest Europe. Ibis, 2012, 154, 307-317.	1.9	35
68	In Search of Human-associated Bacterial Pathogens in Antarctic Wildlife: Report from Six Penguin Colonies Regularly Visited by Tourists. Ambio, 2005, 34, 430-432.	5.5	34
69	Temporal dynamics, diversity, and interplay in three components of the virodiversity of a Mallard population: Influenza A virus, avian paramyxovirus and avian coronavirus. Infection, Genetics and Evolution, 2015, 29, 129-137.	2.3	34
70	Assessing the Role of Seabirds in the Ecology of Influenza A Viruses. Avian Diseases, 2016, 60, 378.	1.0	34
71	Co-infection with Babesia divergens and Anaplasma phagocytophilum in cattle (Bos taurus), Sweden. Ticks and Tick-borne Diseases, 2017, 8, 933-935.	2.7	34
72	Does influenza A virus infection affect movement behaviour during stopover in its wild reservoir host?. Royal Society Open Science, 2016, 3, 150633.	2.4	33

#	Article	IF	Citations
73	Antimicrobial Resistance Profiles of Campylobacter jejuni Isolates from Wild Birds in Sweden. Applied and Environmental Microbiology, 2005, 71, 2438-2441.	3.1	30
74	Evidence for continental-scale dispersal of antimicrobial resistant bacteria by landfill-foraging gulls. Science of the Total Environment, 2021, 764, 144551.	8.0	30
75	Phylogenetic analysis of the non-structural (NS) gene of influenza A viruses isolated from mallards in Northern Europe in 2005. Virology Journal, 2008, 5, 147.	3.4	29
76	The Pattern of Influenza Virus Attachment Varies among Wild Bird Species. PLoS ONE, 2011, 6, e24155.	2.5	29
77	Avian Influenza Surveillance with FTA Cards: Field Methods, Biosafety, and Transportation Issues Solved. Journal of Visualized Experiments, 2011, , .	0.3	28
78	A Comprehensive Model for the Quantitative Estimation of Seed Dispersal by Migratory Mallards. Frontiers in Ecology and Evolution, 2019, 7, .	2.2	28
79	Antibiotic susceptibility of faecal bacteria in Antarctic penguins. Polar Biology, 2008, 31, 759-763.	1.2	27
80	Amoebae and algae can prolong the survival of Campylobacter species in co-culture. Experimental Parasitology, 2010, 126, 59-64.	1.2	27
81	Candidatus Neoehrlichia mikurensis in Ticks from Migrating Birds in Sweden. PLoS ONE, 2015, 10, e0133250.	2.5	27
82	Babesia, Theileria, and Hepatozoon species in ticks infesting animal hosts in Romania. Parasitology Research, 2017, 116, 2291-2297.	1.6	27
83	Canine tick-borne diseases in pet dogs from Romania. Parasites and Vectors, 2017, 10, 155.	2.5	27
84	Acanthamoeba - Campylobacter Coculture as a Novel Method for Enrichment of Campylobacter Species. Applied and Environmental Microbiology, 2007, 73, 6864-6869.	3.1	26
85	Increase in Acid Tolerance of Campylobacter jejuni through Coincubation with Amoebae. Applied and Environmental Microbiology, 2010, 76, 4194-4200.	3.1	26
86	A Panel of Stably Expressed Reference Genes for Real-Time qPCR Gene Expression Studies of Mallards (Anas platyrhynchos). PLoS ONE, 2016, 11, e0149454.	2.5	26
87	Response to Comment on "Rapid Advance of Spring Arrival Dates in Long-Distance Migratory Birds". Science, 2007, 315, 598c-598c.	12.6	24
88	Flexibility of Continental Navigation and Migration in European Mallards. PLoS ONE, 2013, 8, e72629.	2.5	24
89	Surveillance for West Nile Virus in Wild Birds from Northern Europe. Vector-Borne and Zoonotic Diseases, 2011, 11, 77-79.	1.5	23
90	Oseltamivir-Resistant Influenza A (H1N1) Virus Strain with an H274Y Mutation in Neuraminidase Persists without Drug Pressure in Infected Mallards. Applied and Environmental Microbiology, 2015, 81, 2378-2383.	3.1	23

#	Article	IF	Citations
91	Influenza A virus evolution and spatio-temporal dynamics in Eurasian wild birds: a phylogenetic and phylogeographical study of whole-genome sequence data. Journal of General Virology, 2015, 96, 2050-2060.	2.9	23
92	Genetic diversity and host associations in Campylobacter jejuni from human cases and broilers in 2000 and 2008. Veterinary Microbiology, 2015, 178, 94-98.	1.9	23
93	Where do all the subtypes go? Temporal dynamics of H8â€“H12 influenza A viruses in waterfowl. Virus Evolution, 2018, 4, vey025.	4.9	23
94	Host Range of Influenza A Virus H1 to H16 in Eurasian Ducks Based on Tissue and Receptor Binding Studies. Journal of Virology, 2021, 95, .	3.4	23
95	Resistance Mutation R292K Is Induced in Influenza A(H6N2) Virus by Exposure of Infected Mallards to Low Levels of Oseltamivir. PLoS ONE, 2013, 8, e71230.	2.5	22
96	As the Duck Fliesâ€"Estimating the Dispersal of Low-Pathogenic Avian Influenza Viruses by Migrating Mallards. Frontiers in Ecology and Evolution, 2018, 6, .	2.2	22
97	Red fox and tick-borne encephalitis (TBE) in humans: Can predators influence public health?. Scandinavian Journal of Infectious Diseases, 2008, 40, 527-532.	1.5	20
98	Campylobacter jejuniin Penguins, Antarctica. Emerging Infectious Diseases, 2009, 15, 847-849.	4.3	20
99	Influenza A(H7N9) Virus Acquires Resistance-Related Neuraminidase I222T Substitution When Infected Mallards Are Exposed to Low Levels of Oseltamivir in Water. Antimicrobial Agents and Chemotherapy, 2015, 59, 5196-5202.	3.2	20
100	Characterization of avian influenza virus attachment patterns to human and pig tissues. Scientific Reports, 2018, 8, 12215.	3.3	20
101	Hotspots in the grid: Avian sensitivity and vulnerability to collision risk from energy infrastructure interactions in Europe and North Africa. Journal of Applied Ecology, 2022, 59, 1496-1512.	4.0	20
102	Does influenza A affect body condition of wild mallard ducks, or vice versa ? A reply to Flint and Franson. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 2347-2349.	2.6	19
103	Forecasting risk of tick-borne encephalitis (TBE): Using data from wildlife and climate to predict next year's number of human victims. Scandinavian Journal of Infectious Diseases, 2011, 43, 366-372.	1.5	19
104	Prevalence of avian paramyxovirus type 1 in Mallards during autumn migration in the western Baltic Sea region. Virology Journal, 2013, 10, 285.	3.4	19
105	LINKAGE BETWEEN NUCLEAR AND MITOCHONDRIAL DNA SEQUENCES IN AVIAN MALARIA PARASITES: MULTIPLE CASES OF CRYPTIC SPECIATION?. Evolution; International Journal of Organic Evolution, 2004, 58, 1617.	2.3	18
106	The â€œhuman influenza receptorâ€•Neu5AcÎ±2,6Gal is expressed among different taxa of wild birds. Archives of Virology, 2009, 154, 1533-1537.	2.1	18
107	The Potential of Isolation Source to Predict Colonization in Avian Hosts: A Case Study in Campylobacter jejuni Strains From Three Bird Species. Frontiers in Microbiology, 2018, 9, 591.	3.5	18
108	A novel Chlamydiaceaeâ€like bacterium found in faecal specimens from sea birds from the Bering Sea. Environmental Microbiology Reports, 2010, 2, 605-610.	2.4	17

#	Article	IF	Citations
109	Host ecology regulates interspecies recombination in bacteria of the genus Campylobacter. ELife, 2022, 11, .	6.0	17
110	Salmonella Amager, Campylobacter jejuni, and Urease-positive Thermophilic Campylobacter Found in Free-flying Peregrine Falcons (Falco peregrinus) in Sweden. Journal of Wildlife Diseases, 2004, 40, 583-587.	0.8	16
111	Chlamydia psittaci in Swedish Wetland Birds: A Risk to Zoonotic Infection?. Avian Diseases, 2012, 56, 737-740.	1.0	16
112	Capturing individualâ€level parameters of influenza A virus dynamics in wild ducks using multistate models. Journal of Applied Ecology, 2016, 53, 1289-1297.	4.0	16
113	Molecular survey of neglected bacterial pathogens reveals an abundant diversity of species and genotypes in ticks collected from animal hosts across Romania. Parasites and Vectors, 2018, 11, 144.	2.5	16
114	A simple method for long-term storage of Acanthamoeba species. Parasitology Research, 2009, 104, 935-937.	1.6	15
115	A rapid and transient innate immune response to avian influenza infection in mallards. Molecular Immunology, 2018, 95, 64-72.	2.2	15
116	Infected or not: are PCR-positive oropharyngeal swabs indicative of low pathogenic influenza A virus infection in the respiratory tract of Mallard Anas platyrhynchos?. Veterinary Research, 2014, 45, 53.	3.0	14
117	Enteropathogenic Escherichia coli (EPEC) in Antarctic fur seals Arctocephalus gazella. Polar Biology, 2007, 30, 1227-1229.	1.2	13
118	Antibiotic Resistance Patterns in Fecal Bacteria Isolated from Christmas Shearwater (Puffinus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 387 486-489.	1.0	13
119	Chlamydia psittaciin birds of prey, Sweden. Infection Ecology and Epidemiology, 2012, 2, 8435.	0.8	13
120	How Does Sampling Methodology Influence Molecular Detection and Isolation Success in Influenza A Virus Field Studies?. Applied and Environmental Microbiology, 2016, 82, 1147-1153.	3.1	13
121	In search of human-associated bacterial pathogens in Antarctic wildlife: report from six penguin colonies regularly visited by tourists. Ambio, 2005, 34, 430-2.	5.5	13
122	Attachment Patterns of Human and Avian Influenza Viruses to Trachea and Colon of 26 Bird Species â€“ Support for the Community Concept. Frontiers in Microbiology, 2019, 10, 815.	3.5	12
123	Birds and Viruses at a Crossroad - Surveillance of Influenza A Virus in Portuguese Waterfowl. PLoS ONE, 2012, 7, e49002.	2.5	12
124	Differentiation and phylogeny of the olivaceous warbler Hippolais pallida species complex. Journal Fur Ornithologie, 2005, 146, 127-136.	1.2	11
125	Trends in Body Mass of Ducks over Time: The Hypotheses in Guillemain et al. Revisited. Ambio, 2011, 40, 338-340.	5.5	10
126	How to track a flu virus. Nature, 2012, 483, 535-536.	27.8	10

#	Article	IF	Citations
127	No evidence for homosubtypic immunity of influenza H3 in Mallards following vaccination in a natural experimental system. Molecular Ecology, 2017, 26, 1420-1431.	3.9	10
128	Haemosporidian Blood Parasites in European Birds of Prey and Owls. Journal of Parasitology, 2008, 94, 709.	0.7	10
129	Roadside ecology and epidemiology of tick-borne diseases. Scandinavian Journal of Infectious Diseases, 2008, 40, 853-858.	1.5	9
130	On the potential roles of ticks and migrating birds in the ecology of West Nile virus. Infection Ecology and Epidemiology, 2014, 4, 20943.	0.8	9
131	Zero Prevalence of Influenza A Virus in Two Raptor Species by Standard Screening. Vector-Borne and Zoonotic Diseases, 2010, 10, 387-390.	1.5	8
132	Three Babesia species in Ixodes ricinus ticks from migratory birds in Sweden. Parasites and Vectors, 2021, 14, 183.	2.5	8
133	Non-breeding ecology of the Whinchat Saxicola rubetra in Nigeria. Ornis Svecica, 2012, 22, 25-32.	0.1	8
134	Campylobacter jejuni sequence types show remarkable spatial and temporal stability in Blackbirds. Infection Ecology and Epidemiology, 2015, 5, 28383.	0.8	7
135	Molecular identification of papillomavirus in ducks. Scientific Reports, 2018, 8, 9096.	3.3	7
136	A Comparative Study of the Innate Humoral Immune Response to Avian Influenza Virus in Wild and Domestic Mallards. Frontiers in Microbiology, 2020, 11, 608274.	3.5	7
137	Migration patterns, population trends and morphometrics of Ruddy Turnstones Arenaria interpres passing through Ottenby in south-eastern Sweden. Ornis Svecica, 2005, 15, 63-72.	0.1	7
138	Barn Swallows (Hirundo rustica) Test Negative for Salmonella. Vector-Borne and Zoonotic Diseases, 2008, 8, 451-454.	1.5	6
139	Characterization of Campylobacter spp. isolated from wild birds in the Antarctic and Sub-Antarctic. PLoS ONE, 2018, 13, e0206502.	2.5	6
140	The timing of spring migration in trans-Saharan migrants: a comparison between Ottenby, Sweden and Capri, Italy. Ornis Svecica, 2006, 16, 27-33.	0.1	6
141	Moult strategies in the Common Whitethroat Sylvia c. communis in northern Nigeria. Ibis, 2002, 144, E11-E18.	1.9	5
142	Intestinal spirochaetes (genusBrachyspira) colonise wild birds in the southern Atlantic region and Antarctica. Infection Ecology and Epidemiology, 2015, 5, 29296.	0.8	5
143	Migration distance affects how closely Eurasian wigeons follow spring phenology during migration. Movement Ecology, 2021, 9, 61.	2.8	5
144	A novelSalmonellaserovar isolated from Peregrine Falcon (Falco peregrinus) nestlings in Sweden:Salmonella enterica entericaserovar Pajala (SalmonellaPajala). Infection Ecology and Epidemiology, 2012, 2, 7373.	0.8	4

#	Article	IF	Citations
145	Multilocus Sequence Typing and FlaA Sequencing Reveal the Genetic Stability of Campylobacter jejuni Enrichment during Coculture with Acanthamoeba polyphaga. Applied and Environmental Microbiology, 2013, 79, 2477-2479.	3.1	4
146	Of Ducks and Men: Ecology and Evolution of a Zoonotic Pathogen in a Wild Reservoir Host. Advances in Environmental Microbiology, 2017, , 247-286.	0.3	4
147	Narrative overview on wild bird migration in the context of highly pathogenic avian influenza incursion into the European Union. EFSA Supporting Publications, 2017, 14, 1283E.	0.7	4
148	Expression of immune genes RIG-I and Mx in mallard ducks infected with low pathogenic avian influenza (LPAI): A dataset. Data in Brief, 2018, 18, 1562-1566.	1.0	3
149	The accuracy of field sex determination in the Common Whitethroat Sylvia c. communis. Ornis Svecica, 2000, 10, 67-70.	0.1	3
150	The stopover behaviour of the Garden Warbler Sylvia borin in Obudu, southeast Nigeria. Ornis Svecica, 2011, 21, 29-36.	0.1	3
151	Freeze-drying can replace cold-chains for transport and storage of fecal microbiome samples. PeerJ, 2022, 10, e13095.	2.0	3
152	Population fluctuations and timing of spring migration of the Scandinavian Bluethroat Luscinia svecica svecica at Ottenby Bird Observatory, Sweden, 1955â€“2008. Ornis Svecica, 2011, 21, 92-100.	0.1	2
153	Age and sex determination of Mallards Anas platyrhynchos in autumn. Ornis Svecica, 2016, 26, 61-81.	0.1	2
154	Detection of Neoehrlichia mikurensis DNA in blood donors in southeastern Sweden. Infectious Diseases, 2022, 54, 748-759.	2.8	2
155	A Bayesian semiparametric Jollyâ€“Seber model with individual heterogeneity: An application to migratory mallards at stopover. Annals of Applied Statistics, 2021, 15, .	1.1	1
156	Recension av â€œEuropean Breeding Bird Atlas 2: Distribution, Abundance and Changeâ€•(Keller V,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.1	0
157	Recension av â€œAgeing and Sexing of Migratory East Asian Passerinesâ€•(Norevik G, HellstrÃ¶m M, Liu D &) Tj ETQq1 1 0.784314 rgBT	0.1	0
158	Recension av â€œMoult and Ageing of European Passerines, 2nd editionâ€•(Jenni L & Winkler R, 2020). Ornis Svecica, 0, 31, .	0.1	0
159	Recension av â€œFugleatlas: de danske ynglefugles udbredelse 2014â€“2017â€•(VikstrÃ¸m T & MoshÃ¸j CM,) Tj ETQq1 1 0.784314	0.1	0
160	Ornis Svecica moulting into its new plumage. Ornis Svecica, 0, 30, .	0.1	0
161	On the wing. Ornis Svecica, 0, 32, 1-4.	0.1	0