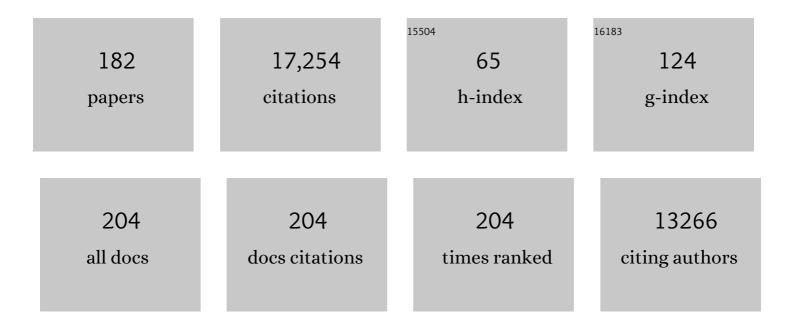
Josephine M Pemberton

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Age, Sex, Density, Winter Weather, and Population Crashes in Soay Sheep. Science, 2001, 292, 1528-1531.	12.6	820
2	Phenological sensitivity to climate across taxa and trophic levels. Nature, 2016, 535, 241-245.	27.8	705
3	Nonamplifying alleles at microsatellite loci: a caution for parentage and population studies. Molecular Ecology, 1995, 4, 249-252.	3.9	684
4	Noise and determinism in synchronized sheep dynamics. Nature, 1998, 394, 674-677.	27.8	498
5	PARASITE-MEDIATED SELECTION AGAINST INBRED SOAY SHEEP IN A FREE-LIVING ISLAND POPULATON. Evolution; International Journal of Organic Evolution, 1999, 53, 1259-1267.	2.3	466
6	ANTLER SIZE IN RED DEER: HERITABILITY AND SELECTION BUT NO EVOLUTION. Evolution; International Journal of Organic Evolution, 2002, 56, 1683-1695.	2.3	445
7	Heritability of fitness in a wild mammal population. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 698-703.	7.1	443
8	A microsatellite polymorphism in the gamma interferon gene is associated with resistance to gastrointestinal nematodes in a naturally-parasitized population of Soay sheep. Parasitology, 2001, 122, 571-582.	1.5	431
9	Major histocompatibility complex variation associated with juvenile survival and parasite resistance in a large unmanaged ungulate population (Ovis aries L.). Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 3714-3719.	7.1	408
10	Microsatellites reveal heterosis in red deer. Proceedings of the Royal Society B: Biological Sciences, 1998, 265, 489-495.	2.6	351
11	Population density affects sex ratio variation in red deer. Nature, 1999, 399, 459-461.	27.8	343
12	Dominant rams lose out by sperm depletion. Nature, 2001, 409, 681-682.	27.8	342
13	Sexually antagonistic genetic variation for fitness in red deer. Nature, 2007, 447, 1107-1110.	27.8	336
14	Wild pedigrees: the way forward. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 613-621.	2.6	308
15	Life history trade-offs at a single locus maintain sexually selected genetic variation. Nature, 2013, 502, 93-95.	27.8	296
16	A retrospective assessment of the accuracy of the paternity inference program cervus. Molecular Ecology, 2000, 9, 801-808.	3.9	282
17	Overt and covert competition in a promiscuous mammal: the importance of weaponry and testes size to male reproductive success. Proceedings of the Royal Society B: Biological Sciences, 2003, 270, 633-640.	2.6	278
18	The Dynamics of Phenotypic Change and the Shrinking Sheep of St. Kilda. Science, 2009, 325, 464-467.	12.6	271

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19	Performance of Marker-Based Relatedness Estimators in Natural Populations of Outbred Vertebrates. Genetics, 2006, 173, 2091-2101.	2.9	250
20	Genomeâ€wide association mapping identifies the genetic basis of discrete and quantitative variation in sexual weaponry in a wild sheep population. Molecular Ecology, 2011, 20, 2555-2566.	3.9	217
21	Environmental Coupling of Selection and Heritability Limits Evolution. PLoS Biology, 2006, 4, e216.	5.6	217
22	Introgression Through Rare Hybridization: A Genetic Study of a Hybrid Zone Between Red and Sika Deer (Genus Cervus) in Argyll, Scotland. Genetics, 1999, 152, 355-371.	2.9	210
23	Inbreeding depression across the lifespan in a wild mammal population. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 3585-3590.	7.1	208
24	Measuring inbreeding depression in the wild: the old ways are the best. Trends in Ecology and Evolution, 2004, 19, 613-615.	8.7	200
25	Estimating quantitative genetic parameters in wild populations: a comparison of pedigree and genomic approaches. Molecular Ecology, 2014, 23, 3434-3451.	3.9	199
26	The Evolutionary Demography of Ecological Change: Linking Trait Variation and Population Growth. Science, 2007, 315, 1571-1574.	12.6	196
27	Landscape features affect gene flow of Scottish Highland red deer (<i>Cervus elaphus</i>). Molecular Ecology, 2008, 17, 981-996.	3.9	182
28	Fitness Correlates of Heritable Variation in Antibody Responsiveness in a Wild Mammal. Science, 2010, 330, 662-665.	12.6	182
29	Inbreeding avoidance, tolerance, or preference in animals?. Trends in Ecology and Evolution, 2013, 28, 205-211.	8.7	176
30	Maternal genetic effects set the potential for evolution in a free-living vertebrate population. Journal of Evolutionary Biology, 2004, 18, 405-414.	1.7	169
31	Life history correlates of oxidative damage in a freeâ€living mammal population. Functional Ecology, 2009, 23, 809-817.	3.6	169
32	Comparing molecular measures for detecting inbreeding depression. Journal of Evolutionary Biology, 2002, 15, 20-31.	1.7	160
33	Inter―and Intrasexual Variation in Aging Patterns across Reproductive Traits in a Wild Red Deer Population. American Naturalist, 2009, 174, 342-357.	2.1	156
34	Conserved Genetic Architecture Underlying Individual Recombination Rate Variation in a Wild Population of Soay Sheep (<i>Ovis aries</i>). Genetics, 2016, 203, 583-598.	2.9	144
35	POSITIVE GENETIC CORRELATION BETWEEN PARASITE RESISTANCE AND BODY SIZE IN A FREE-LIVING UNGULATE POPULATION. Evolution; International Journal of Organic Evolution, 2001, 55, 2116-2125.	2.3	143
36	Predictors of reproductive cost in female Soay sheep. Journal of Animal Ecology, 2005, 74, 201-213.	2.8	139

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37	Repeated selection of morphometric traits in the Soay sheep on St Kilda. Journal of Animal Ecology, 1999, 68, 472-488.	2.8	134
38	THE DEMOGRAPHIC CONSEQUENCES OF RELEASING A POPULATION OF RED DEER FROM CULLING. Ecology, 2004, 85, 411-422.	3.2	134
39	Advancing breeding phenology in response to environmental change in a wild red deer population. Global Change Biology, 2011, 17, 2455-2469.	9.5	132
40	Bovine microsatellite loci are highly conserved in red deer (Cervus elaphus), sika deer (Cervus nippon) Tj ETQqO	0 0 rgBT / 1.7	Overlock 10 T
41	Bottlenecks, drift and differentiation: the population structure and demographic history of sika deer (Cervus nippon) in the Japanese archipelago. Molecular Ecology, 2001, 10, 1357-1370.	3.9	127
42	Patterns of body mass senescence and selective disappearance differ among three species of free-living ungulates. Ecology, 2011, 92, 1936-1947.	3.2	124
43	No evidence for major histocompatibility complex–dependent mating patterns in a free–living ruminant population. Proceedings of the Royal Society B: Biological Sciences, 1997, 264, 1813-1819.	2.6	123
44	ESTIMATING SELECTION ON NEONATAL TRAITS IN RED DEER USING ELASTICITY PATH ANALYSIS. Evolution; International Journal of Organic Evolution, 2003, 57, 2879-2892.	2.3	120
45	THE PREDICTION OF ADAPTIVE EVOLUTION: EMPIRICAL APPLICATION OF THE SECONDARY THEOREM OF SELECTION AND COMPARISON TO THE BREEDER'S EQUATION. Evolution; International Journal of Organic Evolution, 2012, 66, 2399-2410.	2.3	119
46	Lifelong leukocyte telomere dynamics and survival in a free-living mammal. Aging Cell, 2016, 15, 140-148.	6.7	118
47	Parasite-associated polymorphism in a cyclic ungulate population. Proceedings of the Royal Society B: Biological Sciences, 1993, 254, 7-13.	2.6	117
48	Compelling evidence that a single nucleotide substitution in TYRP1 is responsible for coat-colour polymorphism in a free-living population of Soay sheep. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 619-626.	2.6	116
49	LIVE FAST, DIE YOUNG: TRADE-OFFS BETWEEN FITNESS COMPONENTS AND SEXUALLY ANTAGONISTIC SELECTION ON WEAPONRY IN SOAY SHEEP. Evolution; International Journal of Organic Evolution, 2006, 60, 2168-2181.	2.3	114
50	Variable extent of hybridization between invasive sika (<i>Cervus nippon</i>) and native red deer (<i>C.) Tj ETQ</i>	q0	BT /Overlock 10
51	Detecting the True Extent of Introgression during Anthropogenic Hybridization. Trends in Ecology and Evolution, 2019, 34, 315-326.	8.7	105
52	Natural Selection on Individual Variation in Tolerance of Gastrointestinal Nematode Infection. PLoS Biology, 2014, 12, e1001917.	5.6	104
53	Environmental Heterogeneity Generates Fluctuating Selection on a Secondary Sexual Trait. Current Biology, 2008, 18, 751-757.	3.9	99
54	Comparing parentage inference software: reanalysis of a red deer pedigree. Molecular Ecology, 2010, 19, 1914-1928.	3.9	98

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55	Asynchrony of senescence among phenotypic traits in a wild mammal population. Experimental Gerontology, 2015, 71, 56-68.	2.8	92
56	Quantitative genetics of growth and cryptic evolution of body size in an island population. Evolutionary Ecology, 2007, 21, 337-356.	1.2	91
57	Quantitative trait loci (QTL) mapping of resistance to strongyles and coccidia in the free-living Soay sheep (Ovis aries). International Journal for Parasitology, 2007, 37, 121-129.	3.1	87
58	Reproductive senescence in female <scp>S</scp> oay sheep: variation across traits and contributions of individual ageing and selective disappearance. Functional Ecology, 2013, 27, 184-195.	3.6	82
59	Genomic analysis reveals depression due to both individual and maternal inbreeding in a freeâ€living mammal population. Molecular Ecology, 2016, 25, 3152-3168.	3.9	79
60	Ageing in a variable habitat: environmental stress affects senescence in parasite resistance in St Kilda Soay sheep. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 3477-3485.	2.6	77
61	Density-dependent selection in a fluctuating ungulate population. Proceedings of the Royal Society B: Biological Sciences, 1996, 263, 31-38.	2.6	75
62	Heterogeneity of genetic architecture of body size traits in a freeâ€living population. Molecular Ecology, 2015, 24, 1810-1830.	3.9	72
63	Heritable variation in resistance to gastro-intestinal nematodes in an unmanaged mammal population. Proceedings of the Royal Society B: Biological Sciences, 1999, 266, 1283-1290.	2.6	71
64	Gastrointestinal nematode species burdens and host mortality in a feral sheep population. Parasitology, 2006, 133, 485-496.	1.5	71
65	Inbreeding depression in red deer calves. BMC Evolutionary Biology, 2011, 11, 318.	3.2	69
66	SHARED SPATIAL EFFECTS ON QUANTITATIVE GENETIC PARAMETERS: ACCOUNTING FOR SPATIAL AUTOCORRELATION AND HOME RANGE OVERLAP REDUCES ESTIMATES OF HERITABILITY IN WILD RED DEER. Evolution; International Journal of Organic Evolution, 2012, 66, 2411-2426.	2.3	69
67	Fluctuating optimum and temporally variable selection on breeding date in birds and mammals. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 31969-31978.	7.1	69
68	Genetic variance in fitness indicates rapid contemporary adaptive evolution in wild animals. Science, 2022, 376, 1012-1016.	12.6	69
69	SELECTION ON MOTHERS AND OFFSPRING: WHOSE PHENOTYPE IS IT AND DOES IT MATTER?. Evolution; International Journal of Organic Evolution, 2005, 59, 451-463.	2.3	68
70	Molecular analysis of a promiscuous, fluctuating mating system. Biological Journal of the Linnean Society, 1999, 68, 289-301.	1.6	67
71	Male mate choice influences female promiscuity in Soay sheep. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 365-373.	2.6	67
72	The use of marker-based relationship information to estimate the heritability of body weight in a natural population: a cautionary tale. Journal of Evolutionary Biology, 2002, 15, 92-99.	1.7	66

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#	Article	IF	CITATIONS
73	The maintenance of genetic polymorphism in small island populations: large mammals in the Hebrides. Philosophical Transactions of the Royal Society B: Biological Sciences, 1996, 351, 745-752.	4.0	65
74	A framework for power and sensitivity analyses for quantitative genetic studies of natural populations, and case studies in Soay sheep (<i>Ovis aries</i>). Journal of Evolutionary Biology, 2007, 20, 2309-2321.	1.7	62
75	Heterozygosity, inbreeding and neonatal traits in Soay sheep on St Kilda. Molecular Ecology, 2005, 14, 3383-3393.	3.9	61
76	Heritable, Heterogeneous, and Costly Resistance of Sheep against Nematodes and Potential Feedbacks to Epidemiological Dynamics. American Naturalist, 2014, 184, S58-S76.	2.1	60
77	No evidence for inbreeding avoidance in a great reed warbler population. Behavioral Ecology, 2007, 18, 157-164.	2.2	59
78	The Impact of Environmental Heterogeneity on Genetic Architecture in a Wild Population of Soay Sheep. Genetics, 2009, 181, 1639-1648.	2.9	58
79	Genetic architecture and lifetime dynamics of inbreeding depression in a wild mammal. Nature Communications, 2021, 12, 2972.	12.8	58
80	Sex differences in the consequences of maternal loss in a long-lived mammal, the red deer (Cervus) Tj ETQq0 0	0 rgBT /Ove 1.4	erlock 10 Tf 5
81	Multivariate immune defences and fitness in the wild: complex but ecologically important associations among plasma antibodies, health and survival. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20132931.	2.6	57
82	A High-Density Linkage Map Reveals Sexual Dimorphism in Recombination Landscapes in Red Deer (<i>Cervus elaphus</i>). G3: Genes, Genomes, Genetics, 2017, 7, 2859-2870.	1.8	57
83	Early life expenditure in sexual competition is associated with increased reproductive senescence in male red deer. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20140792.	2.6	56
84	Senescence in immunity against helminth parasites predicts adult mortality in a wild mammal. Science, 2019, 365, 1296-1298.	12.6	55
85	Introgression and the fate of domesticated genes in a wild mammal population. Molecular Ecology, 2013, 22, 4210-4221.	3.9	53
86	Trading offspring size for number in a variable environment: selection on reproductive investment in female Soay sheep. Journal of Animal Ecology, 2009, 78, 354-364.	2.8	52
87	Genetic Analysis of Life-History Constraint and Evolution in a Wild Ungulate Population. American Naturalist, 2012, 179, E97-E114.	2.1	52
88	The role of selection and evolution in changing parturition date in a red deer population. PLoS Biology, 2019, 17, e3000493.	5.6	52
89	ANTLER SIZE IN RED DEER: HERITABILITY AND SELECTION BUT NO EVOLUTION. Evolution; International Journal of Organic Evolution, 2002, 56, 1683.	2.3	49

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91	Red and sika deer in the British Isles, current management issues and management policy. Mammalian Biology, 2009, 74, 247-262.	1.5	45
92	Natural selection on a measure of parasite resistance varies across ages and environmental conditions in a wild mammal. Journal of Evolutionary Biology, 2011, 24, 1664-1676.	1.7	44
93	Use of genetic data for conservation management: the case of the Arabian oryx. Animal Conservation, 1999, 2, 269-278.	2.9	41
94	Constraints on plastic responses to climate variation in red deer. Biology Letters, 2005, 1, 457-460.	2.3	41
95	Detecting genes for variation in parasite burden and immunological traits in a wild population: testing the candidate gene approach. Molecular Ecology, 2013, 22, 757-773.	3.9	39
96	Sex differences in leucocyte telomere length in a freeâ€living mammal. Molecular Ecology, 2017, 26, 3230-3240.	3.9	38
97	Multiple spatial behaviours govern social network positions in a wild ungulate. Ecology Letters, 2021, 24, 676-686.	6.4	38
98	Gestation length variation in a wild ungulate. Functional Ecology, 2011, 25, 691-703.	3.6	37
99	A Multivariate Analysis of Genetic Constraints to Life History Evolution in a Wild Population of Red Deer. Genetics, 2014, 198, 1735-1749.	2.9	37
100	Genetic diversity and population structure of Scottish Highland red deer (Cervus elaphus) populations: a mitochondrial survey. Heredity, 2009, 102, 199-210.	2.6	36
101	Cortisol but not testosterone is repeatable and varies with reproductive effort in wild red deer stags. General and Comparative Endocrinology, 2015, 222, 62-68.	1.8	36
102	A Genomic Region Containing <i>REC8</i> and <i>RNF212B</i> Is Associated with Individual Recombination Rate Variation in a Wild Population of Red Deer (<i>Cervus elaphus</i>). G3: Genes, Genomes, Genetics, 2018, 8, 2265-2276.	1.8	36
103	Declining home range area predicts reduced lateâ€life survival in two wild ungulate populations. Ecology Letters, 2018, 21, 1001-1009.	6.4	35
104	Phenotypic correlates of hybridisation between red and sika deer (genus <i>Cervus</i>). Journal of Animal Ecology, 2010, 79, 414-425.	2.8	34
105	Cellular and humoral immunity in a wild mammal: Variation with age & sex and association with overwinter survival. Ecology and Evolution, 2016, 6, 8695-8705.	1.9	34
106	Introgression of exotic <i>Cervus</i> (<i>nippon</i> and <i>canadensis</i>) into red deer (<i>Cervus) Tj ETQq0 0 2122-2134.</i>	0 rgBT /0 1.9	verlock 10 T 34
107	The Fine-Scale Landscape of Immunity and Parasitism in a Wild Ungulate Population. Integrative and Comparative Biology, 2019, 59, 1165-1175.	2.0	34

108Inbreeding depression by environment interactions in a free-living mammal population. Heredity, 2017,
118, 64-77.2.633

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#	Article	IF	CITATIONS
109	Seasonality of helminth infection in wild red deer varies between individuals and between parasite taxa. Parasitology, 2018, 145, 1410-1420.	1.5	33
110	Multiple pathways mediate the effects of climate change on maternal reproductive traits in a red deer population. Ecology, 2014, 95, 3124-3138.	3.2	31
111	Relative costs of offspring sex and offspring survival in a polygynous mammal. Biology Letters, 2016, 12, 20160417.	2.3	31
112	A survey of the hybridisation status of Cervus deer species on the island of Ireland. Conservation Genetics, 2014, 15, 823-835.	1.5	30
113	Survival costs of reproduction are mediated by parasite infection in wild Soay sheep. Ecology Letters, 2019, 22, 1203-1213.	6.4	30
114	Digital gene expression analysis of gastrointestinal helminth resistance in Scottish blackface lambs. Molecular Ecology, 2011, 20, 910-919.	3.9	29
115	Reproduction has different costs for immunity and parasitism in a wild mammal. Functional Ecology, 2020, 34, 229-239.	3.6	29
116	Heritable variation in telomere length predicts mortality in Soay sheep. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	29
117	Mutation load decreases with haplotype age in wild Soay sheep. Evolution Letters, 2021, 5, 187-195.	3.3	29
118	Increased genetic marker density reveals high levels of admixture between red deer and introduced Japanese sika in Kintyre, Scotland. Evolutionary Applications, 2020, 13, 432-441.	3.1	28
119	Evidence for Selection-by-Environment but Not Genotype-by-Environment Interactions for Fitness-Related Traits in a Wild Mammal Population. Genetics, 2018, 208, 349-364.	2.9	27
120	The genetic architecture of helminth-specific immune responses in a wild population of Soay sheep (Ovis aries). PLoS Genetics, 2019, 15, e1008461.	3.5	26
121	Investigating temporal changes in hybridization and introgression in a predominantly bimodal hybridizing population of invasive sika (<i>Cervus nippon</i>) and native red deer (<i>C. elaphus</i>) on the Kintyre Peninsula, Scotland. Molecular Ecology, 2010, 19, 910-924.	3.9	25
122	Natural Selection on Antihelminth Antibodies in a Wild Mammal Population. American Naturalist, 2018, 192, 745-760.	2.1	25
123	No evidence for parental age effects on offspring leukocyte telomere length in free-living Soay sheep. Scientific Reports, 2017, 7, 9991.	3.3	24
124	Variation in the prion protein gene (PRNP) sequence of wild deer in Great Britain and mainland Europe. Veterinary Research, 2019, 50, 59.	3.0	22
125	Temporal changes in key factors and key age groups influencing the population dynamics of female red deer. Journal of Animal Ecology, 2000, 69, 1099-1110.	2.8	21
126	Accounting for female space sharing in St. Kilda Soay sheep (<i>OvisÂaries</i>) results in little change in heritability estimates. Journal of Evolutionary Biology, 2017, 30, 96-111.	1.7	21

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127	Landscape-scale vegetation patterns influence small-scale grazing impacts. Biological Conservation, 2015, 192, 218-225.	4.1	20
128	Marker-dependent associations among oxidative stress, growth and survival during early life in a wild mammal. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20161407.	2.6	20
129	Sharing and reporting benefits from biodiversity research. Molecular Ecology, 2021, 30, 1103-1107.	3.9	19
130	Vitamin D status predicts reproductive fitness in a wild sheep population. Scientific Reports, 2016, 6, 18986.	3.3	18
131	RADâ€sequencing for estimating genomic relatedness matrixâ€based heritability in the wild: A case study in roe deer. Molecular Ecology Resources, 2019, 19, 1205-1217.	4.8	18
132	Re-establishment of nematode infra-community and host survivorship in wild Soay sheep following anthelmintic treatment. Veterinary Parasitology, 2009, 161, 47-52.	1.8	17
133	Heritability and cross-sex genetic correlations of early-life circulating testosterone levels in a wild mammal. Biology Letters, 2014, 10, 20140685.	2.3	17
134	Quantification and decomposition of environment-selection relationships. Evolution; International Journal of Organic Evolution, 2018, 72, 851-866.	2.3	17
135	Characterisation of major histocompatibility complex class IIa haplotypes in an island sheep population. Immunogenetics, 2019, 71, 383-393.	2.4	17
136	Molecular analysis of a promiscuous, fluctuating mating system. Biological Journal of the Linnean Society, 1999, 68, 289-301.	1.6	17
137	A panel of microsatellites developed for meerkats (Suricata suricatta) by cross-species amplification and species-specific cloning. Molecular Ecology Notes, 2001, 1, 83-85.	1.7	16
138	VARIANCES AND COVARIANCES OF PHENOLOGICAL TRAITS IN A WILD MAMMAL POPULATION. Evolution; International Journal of Organic Evolution, 2011, 65, 788-801.	2.3	16
139	Joint associations of blood plasma proteins with overwinter survival of a large mammal. Ecology Letters, 2017, 20, 175-183.	6.4	16
140	Fitness Costs of Parasites Explain Multiple Life-History Trade-Offs in a Wild Mammal. American Naturalist, 2021, 197, 324-335.	2.1	16
141	Reâ€mating across years and intralineage polygyny are associated with greater than expected levels of inbreeding in wild red deer. Journal of Evolutionary Biology, 2012, 25, 2457-2469.	1.7	15
142	The Impact of Past Introductions on an Iconic and Economically Important Species, the Red Deer of Scotland. Journal of Heredity, 2013, 104, 14-22.	2.4	15
143	Sex differences in relationships between habitat use and reproductive performance in Soay sheep (<i>Ovis aries</i>). Ecology Letters, 2016, 19, 171-179.	6.4	15
144	Fragmentation and Translocation Distort the Genetic Landscape of Ungulates: Red Deer in the Netherlands. Frontiers in Ecology and Evolution, 2020, 8, .	2.2	15

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145	Genomic prediction in the wild: A case study in Soay sheep. Molecular Ecology, 2022, 31, 6541-6555.	3.9	14
146	The genetic architecture of maternal effects across ontogeny in the red deer. Evolution; International Journal of Organic Evolution, 2020, 74, 1378-1391.	2.3	13
147	Integrated population models poorly estimate the demographic contribution of immigration. Methods in Ecology and Evolution, 2021, 12, 1899-1910.	5.2	13
148	Stable isotopes reveal the importance of seabirds and marine foods in the diet of St Kilda field mice. Scientific Reports, 2020, 10, 6088.	3.3	12
149	Cumulative weather effects can impact across the whole life cycle. Global Change Biology, 2019, 25, 3282-3293.	9.5	11
150	Variation in earlyâ€life testosterone within a wild population of red deer. Functional Ecology, 2014, 28, 1224-1234.	3.6	10
151	Between-population differences in the genetic and maternal components of body mass in roe deer. BMC Evolutionary Biology, 2018, 18, 39.	3.2	10
152	Consistent withinâ€individual plasticity is sufficient to explain temperature responses in red deer reproductive traits. Journal of Evolutionary Biology, 2019, 32, 1194-1206.	1.7	10
153	Genomic analysis reveals a polygenic architecture of antler morphology in wild red deer (<i>Cervus) Tj ETQq1 1</i>	0.784314	rgBT/Overloc
154	Inbreeding depression and the probability of racing in the Thoroughbred horse. Proceedings of the Royal Society B: Biological Sciences, 2022, 289, .	2.6	10
155	Osteoarthritis of the temporo-mandibular joint in free-living Soay sheep on St Kilda. Veterinary Journal, 2015, 203, 120-125.	1.7	9
156	Locusâ€specific introgression in young hybrid swarms: Drift may dominate selection. Molecular Ecology, 2021, 30, 2104-2115.	3.9	9
157	Maternally derived anti-helminth antibodies predict offspring survival in a wild mammal. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20201931.	2.6	9
158	The sheep of St Kilda. , 2003, , 17-51.		8
159	Population dynamics in Soay sheep. , 2003, , 52-88.		8
160	Using genomic prediction to detect microevolutionary change of a quantitative trait. Proceedings of the Royal Society B: Biological Sciences, 2022, 289, 20220330.	2.6	8
161	Exposure to viral and bacterial pathogens among Soay sheep (<i>Ovis aries</i>) of the St Kilda archipelago. Epidemiology and Infection, 2016, 144, 1879-1888.	2.1	7
162	A candidate gene approach to study nematode resistance traits in naturally infected sheep. Veterinary Parasitology, 2017, 243, 71-74.	1.8	7

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#	Article	IF	CITATIONS
163	The genome sequence of the red deer, Cervus elaphus Linnaeus 1758. Wellcome Open Research, 0, 6, 336.	1.8	7
164	Microsatellite variation in Rufous Hummingbirds (Selasphorus rufus) and evidence for a weakly structured population. Journal of Ornithology, 2013, 154, 1029-1037.	1.1	6
165	Maternal longevity and offspring sex in wild ungulates. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20181968.	2.6	6
166	The association between female reproductive performance and leukocyte telomere length in wild Soay sheep. Molecular Ecology, 2022, 31, 6184-6196.	3.9	6
167	Contemporary selection on MHC genes in a freeâ€ŀiving ruminant population. Ecology Letters, 2022, 25, 828-838.	6.4	6
168	Functionally distinct T-helper cell phenotypes predict resistance to different types of parasites in a wild mammal. Scientific Reports, 2022, 12, 3197.	3.3	6
169	Estimating selection on the act of inbreeding in a population with strong inbreeding depression. Journal of Evolutionary Biology, 2018, 31, 1815-1827.	1.7	5
170	Patterns of MHCâ€dependent sexual selection in a freeâ€living population of sheep. Molecular Ecology, 2021, 30, 6733-6742.	3.9	4
171	Habitat impact assessment detects spatially driven patterns of grazing impacts in habitat mosaics but overestimates damage. Journal for Nature Conservation, 2018, 45, 20-29.	1.8	3
172	Within-trio tests provide little support for post-copulatory selection on major histocompatibility complex haplotypes in a free-living population. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20202862.	2.6	3
173	MHC class IIa haplotypes derived by high-throughput SNP screening in an isolated sheep population. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	3
174	Vitamin D status is heritable and under environmentâ€dependent selection in the wild. Molecular Ecology, 2022, 31, 4607-4621.	3.9	3
175	Mating patterns and male breeding success. , 2003, , 166-189.		2
176	Physiological proteins in resource-limited herbivores experiencing a population die-off. Die Naturwissenschaften, 2017, 104, 68.	1.6	2
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