

François Bonhomme

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6603217/publications.pdf>

Version: 2024-02-01

247
papers

14,631
citations

13865

67
h-index

28297

105
g-index

266
all docs

266
docs citations

266
times ranked

10919
citing authors

#	ARTICLE	IF	CITATIONS
1	The coupling hypothesis: why genome scans may fail to map local adaptation genes. <i>Molecular Ecology</i> , 2011, 20, 2044-2072.	3.9	456
2	Subspecific origin and haplotype diversity in the laboratory mouse. <i>Nature Genetics</i> , 2011, 43, 648-655.	21.4	439
3	DNA sequence variation of the mitochondrial control region among geographically and morphologically remote European brown trout <i>Salmo trutta</i> populations. <i>Molecular Ecology</i> , 1992, 1, 161-173.	3.9	392
4	European sea bass genome and its variation provide insights into adaptation to euryhalinity and speciation. <i>Nature Communications</i> , 2014, 5, 5770.	12.8	382
5	Biochemical diversity and evolution in the genus <i>Mus</i> . <i>Biochemical Genetics</i> , 1984, 22, 275-303.	1.7	345
6	A General Synthetic Procedure for Heteropolyniobates. <i>Science</i> , 2002, 297, 996-998.	12.6	310
7	Wild mice: an ever-increasing contribution to a popular mammalian model. <i>Trends in Genetics</i> , 2003, 19, 24-31.	6.7	301
8	Most classical <i>Mus musculus domesticus</i> laboratory mouse strains carry a <i>Mus musculus musculus</i> Y chromosome. <i>Nature</i> , 1985, 315, 70-72.	27.8	242
9	Introgression patterns in the mosaic hybrid zone between <i>Mytilus edulis</i> and <i>M. galloprovincialis</i> . <i>Molecular Ecology</i> , 2003, 12, 447-461.	3.9	223
10	Using neutral, selected, and hitchhiker loci to assess connectivity of marine populations in the genomic era. <i>Evolutionary Applications</i> , 2015, 8, 769-786.	3.1	223
11	High variance in reproductive success of the Pacific oyster (<i>Crassostrea gigas</i> , Thunberg) revealed by microsatellite-based parentage analysis of multifactorial crosses. <i>Aquaculture</i> , 2002, 204, 283-296.	3.5	200
12	Investigation of genetic linkage between myosin and actin genes using an interspecific mouse back-cross. <i>Nature</i> , 1985, 314, 181-183.	27.8	191
13	identix, a software to test for relatedness in a population using permutation methods. <i>Molecular Ecology Notes</i> , 2002, 2, 611-614.	1.7	186
14	Origin and radiation of the house mouse: mitochondrial DNA phylogeny. <i>Journal of Evolutionary Biology</i> , 1996, 9, 391-415.	1.7	169
15	Early Effect of Inbreeding as Revealed by Microsatellite Analyses on <i>Ostrea edulis</i> Larvae. <i>Genetics</i> , 1998, 148, 1893-1906.	2.9	165
16	Molecular phylogenies in the genus <i>Mus</i> : Comparative analysis of electrophoretic, scnDNA hybridization, and mtDNA RFLP data. <i>Biological Journal of the Linnean Society</i> , 1990, 41, 83-103.	1.6	164
17	Origins of Laboratory Mice Deduced from Restriction Patterns of Mitochondrial DNA. <i>Differentiation</i> , 1982, 22, 222-226.	1.9	154
18	Counterselection on sex chromosomes in the <i>Mus musculus</i> European hybrid zone. <i>Journal of Evolutionary Biology</i> , 1993, 6, 529-546.	1.7	153

#	ARTICLE	IF	CITATIONS
19	Do discrepancies between microsatellite and allozyme variation reveal differential selection between sea and lagoon in the sea bass (<i>Dicentrarchus labrax</i>)?. <i>Molecular Ecology</i> , 2000, 9, 457-467.	3.9	153
20	Fragmentation of sea bass populations in the western and eastern Mediterranean as revealed by microsatellite polymorphism. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2000, 267, 929-935.	2.6	151
21	Geographic Structure in the European Flat Oyster (<i>Ostrea edulis</i> L.) as Revealed by Microsatellite Polymorphism. , 2002, 93, 331-351.		141
22	Habitat preference and the marine-speciation paradox. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2003, 270, 1399-1406.	2.6	137
23	Synthesis, Structural Characterization, and Molecular Modeling of Dodecaniobate Keggin Chain Materials. <i>Inorganic Chemistry</i> , 2005, 44, 1774-1785.	4.0	136
24	[SiNb ₁₂ O ₄₀] ¹⁶⁻ and [GeNb ₁₂ O ₄₀] ¹⁶⁻ : Highly Charged Keggin Ions with Sticky Surfaces. <i>Angewandte Chemie - International Edition</i> , 2004, 43, 2787-2792.	13.8	135
25	Genetic study of the Atlantic/Mediterranean transition in sea bass (<i>Dicentrarchus labrax</i>). , 1999, 90, 591-596.		132
26	Increasing genomic information in bivalves through new EST collections in four species: Development of new genetic markers for environmental studies and genome evolution. <i>Gene</i> , 2008, 408, 27-36.	2.2	132
27	Genes for skeletal muscle myosin heavy chains are clustered and are not located on the same mouse chromosome as a cardiac myosin heavy chain gene.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1985, 82, 7183-7187.	7.1	126
28	V β 17 gene polymorphism in wild-derived mouse strains: Two amino acid substitutions in the V β 17 region greatly alter T cell receptor specificity. <i>Cell</i> , 1990, 63, 717-728.	28.9	126
29	An X-chromosome linked locus contributes to abnormal placental development in mouse interspecific hybrids. <i>Nature Genetics</i> , 1996, 12, 398-403.	21.4	126
30	Absence of Y-chromosome introgression across the hybrid zone between <i>Mus musculus domesticus</i> and <i>Mus musculus musculus</i> . <i>Genetical Research</i> , 1986, 48, 191-197.	0.9	125
31	ECOLOGICAL GENETICS IN THE NORTH ATLANTIC: ENVIRONMENTAL GRADIENTS AND ADAPTATION AT SPECIFIC LOCI. <i>Ecology</i> , 2008, 89, 591-107.	3.2	124
32	Solid-state Structures and Solution Behavior of Alkali Salts of the [Nb ₆ O ₁₉] ⁸⁻ Lindqvist Ion. <i>Journal of Cluster Science</i> , 2006, 17, 197-219.	3.3	122
33	Multivariate analysis of genetic exchanges between <i>Solea aegyptiaca</i> and <i>Solea senegalensis</i> (Teleosts.) Tj ETQq1 1,0,784314 rgBT /Over	1.6	120
34	Origin and radiation of the house mouse: clues from nuclear genes. <i>Journal of Evolutionary Biology</i> , 1996, 9, 519-539.	1.7	119
35	Molecular evidence linking hominid evolution to recent radiation of schistosomes (Platyhelminthes:) Tj ETQq1 1 0,784314 rgBT /Over	2.7	116
36	Small effective number of parents (Nb) inferred for a naturally spawned cohort of juvenile European flat oysters <i>Ostrea edulis</i> . <i>Marine Biology</i> , 2007, 150, 1173-1182.	1.5	116

#	ARTICLE	IF	CITATIONS
37	Parallel genetic divergence among coastal marine ecotype pairs of European anchovy explained by differential introgression after secondary contact. <i>Molecular Ecology</i> , 2016, 25, 3187-3202.	3.9	113
38	Heterozygote deficiencies in small lacustrine populations of brook charr <i>Salvelinus Fontinalis</i> Mitchill (Pisces, Salmonidae): a test of alternative hypotheses. <i>Heredity</i> , 2002, 89, 27-35.	2.6	109
39	Inferences of selection and migration in the Danish house mouse hybrid zone. <i>Biological Journal of the Linnean Society</i> , 2005, 84, 593-616.	1.6	104
40	Phylogeography and postglacial expansion of <i>Mus musculus domesticus</i> inferred from mitochondrial DNA coalescent, from Iran to Europe. <i>Molecular Ecology</i> , 2008, 17, 627-641.	3.9	103
41	Thermal regime and host clade, rather than geography, drive Symbiodinium and bacterial assemblages in the scleractinian coral <i>Pocillopora damicornis</i> sensu lato. <i>Microbiome</i> , 2018, 6, 39.	11.1	100
42	The zone of sympatry and hybridization of <i>Mytilus edulis</i> and <i>M. galloprovincialis</i> , as described by intron length polymorphism at locus <i>mac-1</i> . <i>Heredity</i> , 2001, 86, 342-354.	2.6	99
43	Evidence for a mitochondrial lineage originating from the Arabian peninsula in the Madagascar house mouse (<i>Mus musculus</i>). <i>Heredity</i> , 2002, 89, 154-158.	2.6	97
44	The mouse <i>Igh-1 a</i> and <i>Igh-1 b</i> H chain constant regions are derived from two distinct isotypic genes. <i>Immunogenetics</i> , 1989, 29, 92-97.	2.4	96
45	ASSORTATIVE FERTILIZATION AND SELECTION AT LARVAL STAGE IN THE MUSSELS <i>MYTILUS EDULIS</i> AND <i>M. GALLOPROVINCIALIS</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 292-298.	2.3	94
46	Genetic differentiation of the house mouse around the Mediterranean basin: matrilineal footprints of early and late colonization. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2011, 278, 1034-1043.	2.6	94
47	Maintenance of genetic differentiation across a transition zone in the sea: discordance between nuclear and cytoplasmic markers. <i>Journal of Evolutionary Biology</i> , 2005, 18, 70-80.	1.7	93
48	Wormy mice in a hybrid zone: A genetic control of susceptibility to parasite infection. <i>Journal of Evolutionary Biology</i> , 1991, 4, 679-687.	1.7	92
49	The origins of the domestication of the olive tree. <i>Comptes Rendus - Biologies</i> , 2009, 332, 1059-1064.	0.2	90
50	An aqueous route to [TaO] ₂ and solid-state studies of isostructural niobium and tantalum oxide complexes. <i>Dalton Transactions</i> , 2007, , 4517.	3.3	86
51	The origin and remolding of genomic islands of differentiation in the European sea bass. <i>Nature Communications</i> , 2018, 9, 2518.	12.8	86
52	The polyphyletic origin of laboratory inbred mice and their rate of evolution. <i>Biological Journal of the Linnean Society</i> , 1987, 30, 51-58.	1.6	84
53	Lessepsian invasion without bottleneck: example of two rabbitfish species (<i>Siganus rivulatus</i> and <i>T. ETQq1</i>). <i>Overlooked</i> , 2018, 1.5, 84.	1.5	84
54	The application of microsatellite markers to breeding programmes in the sea bass, <i>Dicentrarchus labrax</i> . <i>Aquaculture</i> , 1998, 159, 303-316.	3.5	83

#	ARTICLE	IF	CITATIONS
55	Microsatellite polymorphism and genetic impact of restocking in Mediterranean brown trout (<i>Salmo trutta</i>) populations. <i>Journal of Heredity</i> , 2006, 97, 1-10.	2.6	83
56	A transcriptomic approach of salinity response in the euryhaline teleost, <i>Dicentrarchus labrax</i> . <i>Gene</i> , 2006, 379, 40-50.	2.2	83
57	Comparison between classical and Bayesian methods to investigate the history of olive cultivars using SSR-polymorphisms. <i>Plant Science</i> , 2008, 175, 524-532.	3.6	82
58	GENETIC DIFFERENTIATION AT NUCLEAR AND MITOCHONDRIAL LOCI AMONG LARGE WHITE-HEADED GULLS: SEX-BIASED INTERSPECIFIC GENE FLOW?. <i>Evolution; International Journal of Organic Evolution</i> , 2003, 57, 2865-2878.	2.3	81
59	Differential freshwater adaptation in juvenile sea-bass <i>Dicentrarchus labrax</i> : involvement of gills and urinary system. <i>Journal of Experimental Biology</i> , 2005, 208, 3859-3871.	1.7	80
60	Population subdivision and gene flow in Danish house mice. <i>Molecular Ecology</i> , 1995, 4, 311-320.	3.9	79
61	Lithium Polyniobates. A Lindqvist-Supported Lithium-Water Adamantane Cluster and Conversion of Hexaniobate to a Discrete Keggin Complex. <i>Crystal Growth and Design</i> , 2007, 7, 719-723.	3.0	77
62	A steep cline for mitochondrial DNA in Danish mice. <i>Genetical Research</i> , 1988, 52, 185-193.	0.9	76
63	Seventeen new exon-primed intron-crossing polymerase chain reaction amplifiable introns in fish. <i>Molecular Ecology Notes</i> , 2002, 2, 334-340.	1.7	76
64	Comparative Study of Inorganic Cluster-Surfactant Arrays. <i>Chemistry of Materials</i> , 2005, 17, 2885-2895.	6.7	75
65	Amplification of major histocompatibility complex class II gene diversity by intraexonic recombination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1991, 88, 453-457.	7.1	74
66	Microsatellite-associated heterosis in hatchery-propagated stocks of the shrimp <i>Penaeus stylirostris</i> . <i>Aquaculture</i> , 2000, 184, 203-219.	3.5	74
67	Synthesis, structure, and molecular modeling of a titanoniobate isopolyanion. <i>Journal of Solid State Chemistry</i> , 2003, 176, 111-119.	2.9	74
68	Adaptive Evolution and Effective Population Size in Wild House Mice. <i>Molecular Biology and Evolution</i> , 2012, 29, 2949-2955.	8.9	73
69	<i>Hst-3</i> : an X-linked hybrid sterility gene. <i>Genetical Research</i> , 1990, 56, 163-165.	0.9	72
70	Fluctuating asymmetry in the <i>Mus musculus</i> hybrid zone: a heterotic effect in disrupted co-adapted genomes. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1994, 258, 53-59.	2.6	69
71	Genetic structure of the common sole <i>Solea vulgaris</i> at different geographic scales. <i>Marine Biology</i> , 1995, 122, 361-375.	1.5	69
72	Analyse génétique de la zone d'hybridation entre les deux sous-espèces de souris <i>Mus musculus domesticus</i> et <i>Mus musculus musculus</i> en Bulgarie. <i>Genome</i> , 1988, 30, 427-437.	2.0	66

#	ARTICLE	IF	CITATIONS
73	Molecular phylogeny and plumage evolution in gulls (Larini). <i>Journal of Evolutionary Biology</i> , 2000, 13, 47-57.	1.7	65
74	Variable microsatellites in the Pacific Oyster <i>Crassostrea gigas</i> and other cupped oyster species. <i>Animal Genetics</i> , 2000, 31, 71-72.	1.7	65
75	Differential expression of the heat shock protein Hsp70 in natural populations of the tilapia, <i>Sarotherodon melanotheron</i> , acclimatised to a range of environmental salinities. <i>BMC Ecology</i> , 2010, 10, 11.	3.0	65
76	The musculus-type Y Chromosome of the laboratory mouse is of Asian origin. <i>Mammalian Genome</i> , 1992, 3, 84-91.	2.2	64
77	Fitness landscapes support the dominance theory of post-zygotic isolation in the mussels <i>Mytilus edulis</i> and <i>M. galloprovincialis</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 1253-1260.	2.6	63
78	Direct selection on allozymes is not required to explain heterogeneity among marker loci across a <i>Mytilus</i> hybrid zone. <i>Molecular Ecology</i> , 2003, 12, 2505-2510.	3.9	61
79	A genetic metapopulation model for reef fishes in oceanic islands: the case of the surgeonfish, <i>Acanthurus triostegus</i> . <i>Journal of Evolutionary Biology</i> , 1996, 9, 103-117.	1.7	59
80	Ion-Exchange Behavior of One-Dimensional Linked Dodecaniobate Keggin Ion Materials. <i>Chemistry of Materials</i> , 2008, 20, 2513-2521.	6.7	59
81	Eurasian house mouse (<i>Mus musculus</i> L.) differentiation at microsatellite loci identifies the Iranian plateau as a phylogeographic hotspot. <i>BMC Evolutionary Biology</i> , 2015, 15, 26.	3.2	59
82	Can habitat specialisation maintain a mosaic hybrid zone in marine bivalves?. <i>Marine Ecology - Progress Series</i> , 2002, 245, 157-170.	1.9	59
83	Screening for intron-length polymorphisms in penaeid shrimps using exon-primed intron-crossing (EPIC)-PCR. <i>Molecular Ecology</i> , 2000, 9, 233-235.	3.9	57
84	Some Evolutionary Arguments about what maintains the Pelagic Interval in Reef Fishes. <i>Environmental Biology of Fishes</i> , 2000, 59, 365-383.	1.0	56
85	Genetic structure at different spatial scales in the pearl oyster (<i>Pinctada margaritifera cumingii</i>) in French Polynesian lagoons: beware of sampling strategy and genetic patchiness. <i>Marine Biology</i> , 2008, 155, 147-157.	1.5	56
86	Genetic diversity, clonality and connectivity in the scleractinian coral <i>Pocillopora damicornis</i> : a multi-scale analysis in an insular, fragmented reef system. <i>Marine Biology</i> , 2014, 161, 531-541.	1.5	52
87	Genetical variation and polyphyletic origin in Japanese <i>Mus musculus</i> . <i>Heredity</i> , 1989, 63, 299-308.	2.6	51
88	Large discrepancies in differentiation of allozymes, nuclear and mitochondrial DNA loci in recently founded Pacific populations of the pearl oyster <i>Pinctada margaritifera</i> . <i>Journal of Evolutionary Biology</i> , 2003, 16, 388-398.	1.7	51
89	The Complex History of a Gene Proposed to Participate in a Sexual Isolation Mechanism in House Mice. <i>Molecular Biology and Evolution</i> , 2002, 19, 462-471.	8.9	50
90	Allelic constitution of the hemoglobin beta chain in wild populations of the house mouse, <i>Mus musculus</i> . <i>Biochemical Genetics</i> , 1985, 23, 975-986.	1.7	48

#	ARTICLE	IF	CITATIONS
91	How to detect polymorphisms undergoing selection in marine fishes? A review of methods and case studies, including flatfishes. <i>Journal of Sea Research</i> , 2004, 51, 167-182.	1.6	48
92	Experimental and Theoretical Methods to Investigate Extraframework Species in a Layered Material of Dodecaniobate Anions. <i>Inorganic Chemistry</i> , 2007, 46, 2067-2079.	4.0	48
93	Tracking the Near Eastern origins and European dispersal of the western house mouse. <i>Scientific Reports</i> , 2020, 10, 8276.	3.3	47
94	Population structure of the common sole (<i>Solea solea</i>) in the Northeastern Atlantic and the Mediterranean Sea: revisiting the divide with EPIC markers. <i>Marine Biology</i> , 2007, 151, 327-341.	1.5	46
95	Evolution of pseudogenes in the immunoglobulin V H-gene family of the mouse. <i>Immunogenetics</i> , 1987, 26, 237-248.	2.4	45
96	Assessment of sea bass (<i>Dicentrarchus labrax</i> , L.) stock delimitation in the Bay of Biscay and the English Channel based on mark-recapture and genetic data. <i>Fisheries Research</i> , 2007, 83, 123-132.	1.7	45
97	Polymerase chain reaction-single strand conformation polymorphism analyses of nuclear and chloroplast DNA provide evidence for recombination, multiple introductions and nascent speciation in the <i>Caulerpa taxifolia</i> complex. <i>Molecular Ecology</i> , 2008, 11, 2317-2325.	3.9	45
98	Speciation in the Deep Sea: Multi-Locus Analysis of Divergence and Gene Flow between Two Hybridizing Species of Hydrothermal Vent Mussels. <i>PLoS ONE</i> , 2009, 4, e6485.	2.5	45
99	Variance in the reproductive success of flat oyster <i>Ostrea edulis</i> L. assessed by parentage analyses in natural and experimental conditions. <i>Genetical Research</i> , 2010, 92, 175-187.	0.9	45
100	Genetic structure of <i>Dascyllus aruanus</i> populations in French Polynesia. <i>Marine Biology</i> , 1993, 117, 665-674.	1.5	44
101	Genetic differentiation of deep-sea hydrothermal vent alvinellid populations (Annelida: Polychaeta) along the East Pacific Rise. <i>Heredity</i> , 1995, 74, 376-391.	2.6	44
102	House mouse phylogeography. , 2012, , 278-296.		44
103	Reduced Female Gene Flow in the European Flat Oyster <i>Ostrea edulis</i> . <i>Journal of Heredity</i> , 2004, 95, 510-516.	2.4	43
104	Spatio-temporal variation in the genetic composition of wild populations of pearl oyster (<i>Pinctada</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Ecology, 2004, 13, 2001-2007.	3.9	43
105	Charge density and electrostatic potential analyses in paracetamol. <i>Acta Crystallographica Section B: Structural Science</i> , 2009, 65, 363-374.	1.8	43
106	Evolutionary Patterns in Pearl Oysters of the Genus <i>Pinctada</i> (Bivalvia: Pteriidae). <i>Marine Biotechnology</i> , 2011, 13, 181-192.	2.4	43
107	Properties of Bias and Variance of Two Multiallelic Estimators of FST. <i>Theoretical Population Biology</i> , 2000, 57, 285-296.	1.1	41
108	Fluctuating Asymmetry in <i>Mus musculus</i> Subspecific Hybridization. , 1996, , 275-283.		41

#	ARTICLE	IF	CITATIONS
109	Molecular phylogeny of Fv1. <i>Mammalian Genome</i> , 1998, 9, 1049-1055.	2.2	40
110	Species-wide distribution of highly polymorphic minisatellite markers suggests past and present genetic exchanges among house mouse subspecies. <i>Genome Biology</i> , 2007, 8, R80.	9.6	39
111	Animal behaviour and cancer. <i>Animal Behaviour</i> , 2015, 101, 19-26.	1.9	39
112	SYSTEMATICS OF LARGE WHITE-HEADED GULLS: PATTERNS OF MITOCHONDRIAL DNA VARIATION IN WESTERN EUROPEAN TAXA. <i>Auk</i> , 2002, 119, 603.	1.4	39
113	Mitochondrial DNA analysis of the genetic relationships among populations of scad mackerel (<i>Scomber japonicus</i>) in the western Pacific. <i>Journal of Molecular Evolution</i> , 2002, 55, 699-707.	1.5	37
114	Polymorphism of metallothionein genes in the Pacific oyster <i>Crassostrea gigas</i> as a biomarker of response to metal exposure. <i>Biomarkers</i> , 2002, 7, 439-450.	1.9	37
115	Systematics of Large White-Headed Gulls: Patterns of Mitochondrial DNA Variation in Western European Taxa. <i>Auk</i> , 2002, 119, 603-620.	1.4	37
116	On the trail of Neolithic mice and men towards Transcaucasia: zooarchaeological clues from Nakhchivan (Azerbaijan). <i>Biological Journal of the Linnean Society</i> , 2013, 108, 917-928.	1.6	37
117	Conserved secondary structures in the ITS2 of trematode pre-rRNA. <i>FEBS Letters</i> , 1993, 316, 247-252.	2.8	36
118	Morphometric stepwise discriminant analysis of the five genetically determined European taxa of the genus <i>Mus</i> . <i>Biological Journal of the Linnean Society</i> , 1990, 41, 47-64.	1.6	35
119	Chromosomal introgression in house mice from the hybrid zone between <i>M. m. domesticus</i> and <i>M. m. musculus</i> in Denmark. <i>Biological Journal of the Linnean Society</i> , 1990, 41, 215-227.	1.6	35
120	Range-wide population structure of European sea bass <i>Dicentrarchus labrax</i> . <i>Biological Journal of the Linnean Society</i> , 2015, 116, 86-105.	1.6	35
121	Deleterious mutations in a hybrid zone: can mutational load decrease the barrier to gene flow?. <i>Genetical Research</i> , 2002, 80, 197-204.	0.9	34
122	Transcriptional responses of the black-chinned tilapia <i>Sarotherodon melanocheilus</i> to salinity extremes. <i>Marine Genomics</i> , 2008, 1, 37-46.	1.1	34
123	The south-eastern house mouse <i>Mus musculus castaneus</i> (Rodentia: Muridae) is a polytypic subspecies. <i>Biological Journal of the Linnean Society</i> , 2012, 107, 295-306.	1.6	34
124	Concerted evolution in the GAPDH family of retrotransposed pseudogenes. <i>Mammalian Genome</i> , 1993, 4, 695-703.	2.2	32
125	Genetic hitchhiking in a subdivided population of <i>Mytilus edulis</i> . <i>BMC Evolutionary Biology</i> , 2008, 8, 164.	3.2	31
126	Molecular phylogeny of the genus <i>Pseudoplatystoma</i> (Bleeker, 1862): Biogeographic and evolutionary implications. <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 588-594.	2.7	31

#	ARTICLE	IF	CITATIONS
127	Caractérisation biochimique du complexe d'espèces du genre <i>Lepus</i> en Espagne. <i>Mammalia</i> , 1986, 50, 495-506.	0.7	30
128	Spat collection of the pearl oyster (<i>Pinctada margaritifera cumingii</i>) in French Polynesia: an evaluation of the potential impact on genetic variability of wild and farmed populations after 20 years of commercial exploitation. <i>Aquaculture</i> , 2003, 219, 181-192.	3.5	30
129	Patterns of morphological evolution in the mandible of the house mouse <i>Mus musculus</i> (Rodentia: Tj ETQq1 1 0.784314 rgBT /Overlo	1.6	30
130	Genomic and geographic footprints of differential introgression between two divergent fish species (<i>Solea</i> spp.). <i>Heredity</i> , 2018, 121, 579-593.	2.6	30
131	Estimating total genic diversity in the house mouse. <i>Biochemical Genetics</i> , 1978, 16, 287-297.	1.7	29
132	Nuclear-DNA evidence that northeastern Atlantic <i>Mytilus trossulus</i> mussels carry <i>M. edulis</i> genes. <i>Journal of Molluscan Studies</i> , 1999, 65, 504-507.	1.2	29
133	No reduction in neutral variability of mitochondrial and nuclear genes for a Lessepsian migrant, <i>Upeneus moluccensis</i> . <i>Journal of Fish Biology</i> , 2005, 66, 865-870.	1.6	29
134	Genetic variation and phylogeography of free-living mouse species (genus <i>Mus</i>) in the Balkans and the Middle East. <i>Molecular Ecology</i> , 2007, 16, 4774-4788.	3.9	29
135	Isolation and polymorphism in mitochondrial DNA from <i>Schistosoma mansoni</i> . <i>Molecular and Biochemical Parasitology</i> , 1991, 47, 139-141.	1.1	28
136	Variations of a Y chromosome repeated sequence across subspecies of <i>Mus musculus</i> . <i>Heredity</i> , 1989, 63, 289-297.	2.6	27
137	Population genetic structure in a Robertsonian race of house mice: evidence from microsatellite polymorphism. <i>Heredity</i> , 1998, 80, 70-77.	2.6	27
138	Genetic and morphological differentiation between the two largest breeding colonies of Audouin's Gull <i>Larus audouinii</i> . <i>Ibis</i> , 2003, 145, 448-456.	1.9	27
139	Geographic clines and stepping-stone patterns detected along the East Pacific Rise in the vetigastropod <i>Lepetodrillus elevatus</i> reflect species crypticism. <i>Marine Biology</i> , 2008, 153, 545-563.	1.5	27
140	Serological survey of T-lymphocyte differentiation antigens in wild mice. <i>Immunogenetics</i> , 1985, 22, 211-218.	2.4	26
141	Two deeply divergent mitochondrial clades in the wild mouse <i>Mus macedonicus</i> reveal multiple glacial refuges south of Caucasus. <i>Heredity</i> , 2002, 89, 353-357.	2.6	26
142	Tribasic Lead Maleate and Lead Maleate: Synthesis and Structural and Spectroscopic Characterizations. <i>Inorganic Chemistry</i> , 2005, 44, 7394-7402.	4.0	26
143	Growth hormone and Prolactin-1 gene transcription in natural populations of the black-chinned tilapia <i>Sarotherodon melanocheilus</i> acclimatised to different salinities. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2007, 147, 541-549.	1.6	26
144	Electronic Properties of 3,3-Dimethyl-5,5-bis(1,2,4-triazine): Towards Design of Supramolecular Arrangements of N-Heterocyclic CuI Complexes. <i>Chemistry - A European Journal</i> , 2007, 13, 3414-3423.	3.3	26

#	ARTICLE	IF	CITATIONS
145	X-Y chromosome dissociation in wild derived <i>Mus musculus</i> subspecies, laboratory mice, and their F ₁ hybrids. <i>Cytogenetic and Genome Research</i> , 1982, 34, 241-252.	1.1	25
146	Differences between nuclear and mitochondrial introgressions of brown trout populations from a restocked main river and its unstocked tributary. <i>Biological Journal of the Linnean Society</i> , 1998, 63, 379-392.	1.6	25
147	Phylogenetic position and description of a new species of subgenus <i>Mus</i> (Rodentia, Mammalia) from Thailand. <i>Zoologica Scripta</i> , 2003, 32, 119-127.	1.7	25
148	Isolation by distance and Pleistocene expansion of the lowland populations of the white piranha <i>Serrasalmus rhombeus</i> . <i>Molecular Ecology</i> , 2007, 16, 2488-2503.	3.9	25
149	Gilthead sea bream (<i>Sparus auratus</i>) and European sea bass (<i>Dicentrarchus labrax</i>) expressed sequence tags: Characterization, tissue-specific expression and gene markers. <i>Marine Genomics</i> , 2010, 3, 179-191.	1.1	25
150	Whole exome sequencing of wild-derived inbred strains of mice improves power to link phenotype and genotype. <i>Mammalian Genome</i> , 2017, 28, 416-425.	2.2	25
151	Gene flow at major transitional areas in sea bass (<i>Dicentrarchus labrax</i>) and the possible emergence of a hybrid swarm. <i>Ecology and Evolution</i> , 2012, 2, 3061-3078.	1.9	24
152	Salinity-related variation in gene expression in wild populations of the black-chinned tilapia from various West African coastal marine, estuarine and freshwater habitats. <i>Estuarine, Coastal and Shelf Science</i> , 2011, 91, 102-109.	2.1	23
153	Microsatellite length variation in candidate genes correlates with habitat in the gilthead sea bream <i>Sparus aurata</i> . <i>Molecular Ecology</i> , 2012, 21, 5497-5511.	3.9	23
154	Cancer: an emergent property of disturbed resource-rich environments? Ecology meets personalized medicine. <i>Evolutionary Applications</i> , 2015, 8, 527-540.	3.1	23
155	Lambda light chain constant and variable gene complements in wild-derived inbred mouse strains. <i>European Journal of Immunology</i> , 1985, 15, 535-540.	2.9	22
156	Evolution of tropomyosin functional domains: Differential splicing and genomic constraints. <i>Journal of Molecular Evolution</i> , 1988, 27, 228-235.	1.8	22
157	Induction of a Melanoma-specific Antibody Response by a Monovalent, but not a Divalent, Synthetic GM2 Neoglycopeptide. <i>ChemMedChem</i> , 2009, 4, 582-587.	3.2	22
158	The generation of MHC class II gene polymorphism in the genus <i>Mus</i> . <i>Biological Journal of the Linnean Society</i> , 1990, 41, 141-161.	1.6	21
159	Structures génétiques comparées de trois espèces de rongeurs africains du genre <i>Mastomys</i> au Sénégal. <i>Genetica</i> , 1990, 81, 179-192.	1.1	21
160	Ga ₄ (PO ₄) ₄ ·N ₂ C ₇ H ₁₁ ·1/2H ₂ O: A Three-Dimensional Open Framework with 12-Membered Pores and Bowl-Shaped Building Units. <i>Chemistry of Materials</i> , 2001, 13, 2112-2117.	6.7	21
161	Evidence for male-biased effective sex ratio and recent step-by-step colonization in the bivalve <i>Pinctada mazatlanica</i> . <i>Journal of Evolutionary Biology</i> , 2003, 16, 790-796.	1.7	21
162	Prevalence and evolutionary origins of autoimmune susceptibility alleles in natural mouse populations. <i>Genes and Immunity</i> , 2008, 9, 61-68.	4.1	21

#	ARTICLE	IF	CITATIONS
163	Genomic sequences and genetic differentiation at associated tandem repeat markers in growth hormone, somatotactin and insulin-like growth factor-1 genes of the sea bass, <i>Dicentrarchus labrax</i> . <i>Aquatic Living Resources</i> , 2010, 23, 285-296.	1.2	21
164	Very high genetic fragmentation in a large marine fish, the meagre <i>Argyrosomus regius</i> (Sciaenidae, Perciformes) : impact of reproductive migration, oceanographic barriers and ecological factors. <i>Aquatic Living Resources</i> , 2012, 25, 173-183.	1.2	21
165	The spatial scale of dispersal revealed by admixture tracts. <i>Evolutionary Applications</i> , 2019, 12, 1743-1756.	3.1	21
166	Comparaison génétique des tortues vertes (<i>Chelonia mydas</i>) des Océans Atlantique, Indien et Pacifique. <i>Genetica</i> , 1987, 74, 89-94.	1.1	20
167	Low genetic variability in a widely distributed and abundant clupeid species, <i>Sardinella aurita</i> . New empirical results and interpretations. <i>Journal of Fish Biology</i> , 1998, 52, 861-878.	1.6	20
168	Impact de l'élevage sur la structure génétique des populations méditerranéennes de <i>Dicentrarchus labrax</i> . <i>Aquatic Living Resources</i> , 2005, 18, 71-76.	1.2	20
169	A change of expression in the conserved signaling gene MKK7 is associated with a selective sweep in the western house mouse <i>Mus musculus domesticus</i> . <i>Journal of Evolutionary Biology</i> , 2006, 19, 1486-1496.	1.7	20
170	The contribution of ancient admixture to reproductive isolation between European sea bass lineages. <i>Evolution Letters</i> , 2020, 4, 226-242.	3.3	20
171	The Evolution of House Mice. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 1993, 24, 119-152.	6.7	20
172	Molecular and morphological relationships between two closely related species, <i>Turbinaria ornata</i> and <i>T. conoides</i> (Sargassaceae, Phaeophyceae). <i>Biochemical Systematics and Ecology</i> , 2007, 35, 91-98.	1.3	19
173	Expression of GM1 and GD1a in Liver of Wild Mice 1. <i>Journal of Biochemistry</i> , 1984, 95, 7-12.	1.7	18
174	Serological survey of complement factor H in common laboratory and wild mice: a new third allotype. <i>Immunogenetics</i> , 1989, 29, 148-154.	2.4	18
175	Genetic structure of the common sole (<i>Solea solea</i>) in the Bay of Biscay: Nurseries as units of selection?. <i>Estuarine, Coastal and Shelf Science</i> , 2008, 78, 316-326.	2.1	18
176	A Comparison of Rarefaction and Bayesian Methods for Predicting the Allelic Richness of Future Samples on the Basis of Currently Available Samples. <i>Journal of Heredity</i> , 2006, 97, 483-492.	2.4	17
177	<i>Lgals6</i> , a 2-Million-Year-Old Gene in Mice: A Case of Positive Darwinian Selection and Presence/Absence Polymorphism. <i>Genetics</i> , 2008, 178, 1533-1545.	2.9	17
178	Can Peto's paradox be used as the null hypothesis to identify the role of evolution in natural resistance to cancer? A critical review. <i>BMC Cancer</i> , 2015, 15, 792.	2.6	17
179	Lack of mitochondrial differentiation between Red sea and Mediterranean populations of the <i>Lessepsian rabbitfish</i> , <i>Siganus rivulatus</i> (Perciformes: Siganidae). <i>Scientia Marina</i> , 2003, 67, 215-217.	0.6	17
180	Allozymes, mtDNA and microsatellites study introgression in a stocked trout population in France. <i>Reviews in Fish Biology and Fisheries</i> , 2000, 10, 281-292.	4.9	16

#	ARTICLE	IF	CITATIONS
181	Two ammonium templated gallophosphates: synthesis and structure determination from powder diffraction data of 2D and 3D-GAPON. <i>Microporous and Mesoporous Materials</i> , 2002, 53, 87-96.	4.4	16
182	Mitochondrial and Nuclear DNA Analysis of Genetic Heterogeneity Among Recruitment Cohorts of the European Flat Oyster <i>Ostrea edulis</i> . <i>Biological Bulletin</i> , 2009, 217, 233-241.	1.8	16
183	Expanding hybrid zone between <i>Solea aegyptiaca</i> and <i>Solea senegalensis</i> : genetic evidence over two decades. <i>Molecular Ecology</i> , 2011, 20, 1717-1728.	3.9	16
184	Genetic polymorphism of κ 1 and κ 3 immunoglobulin light chains in the <i>Mus</i> subgenus. <i>European Journal of Immunology</i> , 1983, 13, 312-317.	2.9	15
185	Mouse biodiversity in the genomic era. <i>Cytogenetic and Genome Research</i> , 2004, 105, 385-394.	1.1	15
186	Inferring gene flow in coral reef fishes from different molecular markers: which loci to trust?. <i>Heredity</i> , 2007, 99, 331-339.	2.6	15
187	Genetic population structure of the commercially most important demersal fish in the Southwest Atlantic: The whitemouth croaker (<i>Micropogonias furnieri</i>). <i>Fisheries Research</i> , 2015, 167, 333-337.	1.7	15
188	Mouse subspecies differentiation and H-2 polymorphism. <i>Biological Journal of the Linnean Society</i> , 1990, 41, 125-139.	1.6	14
189	Genetic structure of the feral cat (<i>Felis catus</i> L.) introduced 50½ years ago to a sub-Antarctic island. <i>Polar Biology</i> , 2005, 28, 268-275.	1.2	14
190	Evidence for a slightly deleterious effect of intron polymorphisms at the EF1 α gene in the deep-sea hydrothermal vent bivalve <i>Bathymodiolus</i> . <i>Gene</i> , 2007, 406, 99-107.	2.2	14
191	The effect of environmental salinity on the proteome of the sea bass (<i>Dicentrarchus labrax</i> L.). <i>Animal Genetics</i> , 2007, 38, 601-608.	1.7	14
192	Existence of two widespread semi-isolated genetic entities within Mediterranean anchovies. <i>Marine Biology</i> , 2014, 161, 1063-1071.	1.5	14
193	New evidence for trans-species evolution of the H-2 class I polymorphism. <i>Immunogenetics</i> , 1989, 30, 89-98.	2.4	13
194	Three polymorphic microsatellites in the shrimp <i>Penaeus stylirostris</i> . <i>Animal Genetics</i> , 1999, 30, 234-235.	1.7	13
195	Recent expansion of Northeast Atlantic and Mediterranean populations of <i>Melicertus</i> (<i>Penaeus</i>) <i>kerathurus</i> (Crustacea: Decapoda). <i>Fisheries Science</i> , 2009, 75, 1089-1095.	1.6	13
196	From the laboratory to the wild: salinity-based genetic differentiation of the European sea bass (<i>Dicentrarchus labrax</i>) using gene-associated and gene-independent microsatellite markers. <i>Marine Biology</i> , 2015, 162, 515-538.	1.5	13
197	Editorial Dedicated population genomics for the silent world: the specific questions of marine population genetics. <i>Environmental Epigenetics</i> , 2016, 62, 545-550.	1.8	13
198	ASSORTATIVE FERTILIZATION AND SELECTION AT LARVAL STAGE IN THE MUSSELS <i>MYTILUS EDULIS</i> AND <i>M. GALLOPROVINCIALIS</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 292.	2.3	12

#	ARTICLE	IF	CITATIONS
199	New anonymous nuclear DNA markers for the pearl oyster <i>Pinctada margaritifera</i> and other <i>Pinctada</i> species. <i>Molecular Ecology Notes</i> , 2002, 2, 220-222.	1.7	12
200	Population genetic structure of <i>Penaeus merguensis</i> in Thailand based on nuclear DNA variation. <i>Journal of Experimental Marine Biology and Ecology</i> , 2004, 311, 63-78.	1.5	12
201	Cyprus as an ancient hub for house mice and humans. <i>Journal of Biogeography</i> , 2018, 45, 2619-2630.	3.0	12
202	Population structure and inbreeding in wild house mice (<i>Mus musculus</i>) at different geographic scales. <i>Heredity</i> , 2022, 129, 183-194.	2.6	12
203	Piperazine templated 3D fluorogallophosphate: synthesis and characterization of Ga ₅ (PO ₄) ₅ F ₄ ·2[N ₂ C ₄ H ₁₂]. <i>Microporous and Mesoporous Materials</i> , 2001, 47, 185-194.	4.4	11
204	Out of Africa: demographic and colonization history of the Algerian mouse (<i>Mus spretus</i> Lataste). <i>Heredity</i> , 2019, 122, 150-171.	2.6	11
205	Bidirectional Introgression between <i>Mus musculus domesticus</i> and <i>Mus spretus</i> . <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	11
206	Characterization of newly isolated monoclonal antibodies against MHC of a Japanese wild mouse. <i>Immunogenetics</i> , 1986, 24, 361-367.	2.4	10
207	Species polyphyly and mtDNA introgression among three <i>Serrasalmus</i> sister-species. <i>Molecular Phylogenetics and Evolution</i> , 2008, 46, 375-381.	2.7	10
208	Gene activation cascade triggered by a single photoperiodic cycle inducing flowering in <i>Sinapis alba</i> . <i>Plant Journal</i> , 2009, 59, 962-973.	5.7	10
209	The guardians of inherited oncogenic vulnerabilities. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 1-6.	2.3	10
210	Polymorphism of heavy chain immunoglobulin isotypes in the <i>Mus</i> subgenus. <i>Immunogenetics</i> , 1984, 20, 577-581.	2.4	9
211	Characterization of a centromeric marker on mouse Chromosome 11 and its introgression in a <i>domesticus/musculus</i> hybrid zone. <i>Mammalian Genome</i> , 2004, 15, 924-934.	2.2	9
212	Gene expression plasticity and frontloading promote thermotolerance in <i>Pocillopora</i> corals. , 0, 2, .		9
213	Active hydrothermal vents in the Woodlark Basin may act as dispersing centres for hydrothermal fauna. <i>Communications Earth & Environment</i> , 2022, 3, .	6.8	9
214	Fitness of early life stages in F1 interspecific hybrids between <i>Dicentrarchus labrax</i> and <i>D. punctatus</i> . <i>Aquatic Living Resources</i> , 2012, 25, 67-75.	1.2	8
215	Within-Generation Polygenic Selection Shapes Fitness-Related Traits across Environments in Juvenile Sea Bream. <i>Genes</i> , 2020, 11, 398.	2.4	8
216	New preparation by sublimation at low pressure of glycine and physicochemical study. <i>Journal of Alloys and Compounds</i> , 2008, 458, 595-601.	5.5	7

#	ARTICLE	IF	CITATIONS
217	Introggressive hybridization and morphological transgression in the contact zone between two Mediterranean <i>Solea</i> species. <i>Ecology and Evolution</i> , 2017, 7, 1394-1402.	1.9	7
218	Subtle limits to connectivity revealed by outlier loci within two divergent metapopulations of the deep-sea hydrothermal gastropod <i>Ipremeria nautili</i> . <i>Molecular Ecology</i> , 2022, 31, 2796-2813.	3.9	7
219	Origin of the Laboratory Mouse and Related Subspecies. , 2004, , 3-13.		6
220	Physico-Chemical Characterization of .GAMMA.-Amino n-Butyric Acid Nanoparticles. <i>Chemical and Pharmaceutical Bulletin</i> , 2011, 59, 703-709.	1.3	6
221	Genetic structure of a vulnerable species, the freshwater blenny (<i>Salaria fluviatilis</i>). <i>Conservation Genetics</i> , 2015, 16, 571-581.	1.5	6
222	Microspatial genetic heterogeneity and gene flow in stray cats (<i>Felis catus</i> L.): a comparison of coat colour and microsatellite loci. <i>Molecular Ecology</i> , 2003, 12, 1669-1674.	3.9	5
223	Habitat-related allelic variation revealed by an anonymous DNA locus in reef-dwelling <i>Turbinaria ornata</i> (<i>Fucales</i> , <i>Phaeophyceae</i>). <i>Botanica Marina</i> , 2010, 53, 189-192.	1.2	5
224	Systematics of European coastal anchovies (genus <i>Engraulis</i> Cuvier). <i>Journal of Fish Biology</i> , 2022, 100, 594-600.	1.6	5
225	Inter-Specific Genetic Exchange Despite Strong Divergence in Deep-Sea Hydrothermal Vent Gastropods of the Genus <i>Alviniconcha</i> . <i>Genes</i> , 2022, 13, 985.	2.4	5
226	Restriction fragment length polymorphism and evolution of the mouse immunoglobulin constant region gamma loci. <i>Immunogenetics</i> , 1993, 38, 184-92.	2.4	4
227	GENETIC DIFFERENTIATION AT NUCLEAR AND MITOCHONDRIAL LOCI AMONG LARGE WHITE-HEADED GULLS: SEX-BIASED INTERSPECIFIC GENE FLOW?. <i>Evolution; International Journal of Organic Evolution</i> , 2003, 57, 2865.	2.3	4
228	Adaptive evolution of interferon- β in Glire lineage and evidence for a recent selective sweep in <i>Mus. m. domesticus</i> . <i>Genes and Immunity</i> , 2009, 10, 297-308.	4.1	4
229	Serologic variations in kappa light chains of various <i>Mus</i> species correlate with differences in gene structure. <i>Molecular Immunology</i> , 1986, 23, 831-838.	2.2	3
230	Evolutionary origins of retroposon lineages of Mhc class II Ab alleles. <i>Immunogenetics</i> , 1996, 43, 115-24.	2.4	3
231	Isolation and use of microsatellite loci in <i>Melicertus kerathurus</i> (Crustacea, Penaeidae). <i>Aquatic Living Resources</i> , 2010, 23, 103-107.	1.2	3
232	Populations and Pathways: Genomic Approaches to Understanding Population Structure and Environmental Adaptation. , 2010, , 73-118.		3
233	Population genetic structure in a Robertsonian race of house mice: evidence from microsatellite polymorphism. <i>Heredity</i> , 1998, 80, 70-77.	2.6	3
234	Hybridation naturelle entre deux sous-espèces de souris domestique, <i>Mus musculus domesticus</i> et <i>Mus musculus castaneus</i> , près du lac Casitas (Californie). <i>Genome</i> , 1998, 41, 104-110.	2.0	3

#	ARTICLE	IF	CITATIONS
235	â€œNewâ€•allotypes of the murine Î³ 1 immunoglobulins. <i>Molecular Immunology</i> , 1989, 26, 551-555.	2.2	2
236	Genomic resources for the aquaculture of European sea bass. <i>Aquaculture</i> , 2007, 272, S316-S317.	3.5	2
237	Genetic differentiation of European anchovy (<i>Engraulis encrasicolus</i>) along the Moroccan coast reveals a phylogeographic break around the 25th parallel North. <i>Marine Biology Research</i> , 2017, 13, 342-350.	0.7	2
238	Fitness difference between cryptic salinity-related phenotypes of sea bass (&em> <i>Dicentrarchus</i>) Tj ETQq0 0 0 ggBT /Overlock 10 Tf	0.6	2
239	Recent foundation of Mexican populations of pearl oysters (<i>Pteria sterna</i>) revealed by lack of genetic variation on two mitochondrial genes. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2005, 85, 363-366.	0.8	1
240	Local Mutagenic Impact of Insertions of LTR Retrotransposons on the Mouse Genome. <i>Journal of Molecular Evolution</i> , 2006, 63, 662-675.	1.8	1
241	Origins and Phylogenetic Relationships of the Laboratory Mouse. , 2012, , 3-20.		1
242	In memoriam Jamshid Darvish. <i>Mammalia</i> , 2018, 82, 521-528.	0.7	1
243	Response to ?estimating total genic diversity: Problems with the method of Bonhomme and Selander? by Jon D. Goguen and Lin Chao. <i>Biochemical Genetics</i> , 1980, 18, 337-338.	1.7	0
244	Mouse Inbred Strains, Origins of. , 1998, , 1771-1774.		0
245	Regulation of gene expression by polymorphism at non-coding regions? Prolactin and growth hormone genes in sea bass (<i>Dicentrarchus labrax</i>). <i>Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology</i> , 2008, 150, S204.	1.8	0
246	<i>Mus musculus</i> â†. , 2017, , .		0
247	GÃ©nÃ©tique et Ã©volution du gÃ©nome mitochondrial des MÃ©tazoaires. <i>Genetique, Selection, Evolution</i> , 1986, 18, 73-98.	0.0	0