Peter W H Holland

List of Publications by Year
in descending order

165 papers	15,966 citations	27035 58 h-index	20023 121 g-index
179 all docs	179 docs citations	179 times ranked	15055 citing authors

#	Article	IF	Citations
1	PRD-Class Homeobox Genes in Bovine Early Embryos: Function, Evolution, and Overlapping Roles. Molecular Biology and Evolution, 2022, 39, .	3.5	3
2	Compromised Function of the Pancreatic Transcription Factor PDX1 in a Lineage of Desert Rodents. Journal of Mammalian Evolution, 2021, 28, 965-977.	1.0	0
3	Evidence from oyster suggests an ancient role for Pdx in regulating insulin gene expression in animals. Nature Communications, 2021, 12, 3117.	5.8	10
4	Dynamic Molecular Evolution of Mammalian Homeobox Genes: Duplication, Loss, Divergence and Gene Conversion Sculpt PRDÂClass Repertoires. Journal of Molecular Evolution, 2021, 89, 396-414.	0.8	9
5	The genome sequence of the yellow-tail moth, Euproctis similisÂ(Fuessly, 1775). Wellcome Open Research, 2021, 6, 227.	0.9	0
6	Functional genomics of supergene-controlled behavior in the white-throated sparrow. Faculty Reviews, 2021, 10, 75.	1.7	0
7	The genome sequence of the white ermine, Spilosoma lubricipeda Linnaeus 1758. Wellcome Open Research, 2021, 6, 271.	0.9	0
8	The genome sequence of the peach blossom moth, Thyatira batis (Linnaeus, 1758). Wellcome Open Research, 2021, 6, 267.	0.9	0
9	The genome sequence of the sycamore, Acronicta aceris (Linnaeus, 1758). Wellcome Open Research, 2021, 6, 326.	0.9	1
10	The genome sequence of the lime hawk-moth, Mimas tiliae (Linnaeus, 1758). Wellcome Open Research, 2021, 6, 357.	0.9	2
11	The genome sequence of the spectacle, Abrostola tripartita Hufnagel, 1766. Wellcome Open Research, 2021, 6, 330.	0.9	3
12	Divergent genes in gerbils: prevalence, relation to GC-biased substitution, and phenotypic relevance. BMC Evolutionary Biology, 2020, 20, 134.	3.2	6
13	Mutation of amphioxus Pdx and Cdx demonstrates conserved roles for ParaHox genes in gut, anus and tail patterning. BMC Biology, 2020, 18, 68.	1.7	13
14	Jellyfish genomes reveal distinct homeobox gene clusters and conservation of small RNA processing. Nature Communications, 2020, 11, 3051.	5.8	47
15	Runaway GC Evolution in Gerbil Genomes. Molecular Biology and Evolution, 2020, 37, 2197-2210.	3.5	14
16	Widespread patterns of gene loss in the evolution of the animal kingdom. Nature Ecology and Evolution, 2020, 4, 519-523.	3.4	89
17	Gene profiling of head mesoderm in early zebrafish development: insights into the evolution of cranial mesoderm. EvoDevo, 2019, 10, 14.	1.3	21
18	Of eyes and embryos: subfunctionalization of the<i>CRX</i>homeobox gene in mammalian evolution. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20190830.	1.2	6

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19	The Interaction of Natural Selection and GC Skew May Drive the Fast Evolution of a Sand Rat Homeobox Gene. Molecular Biology and Evolution, 2019, 36, 1473-1480.	3.5	12
20	Reconstruction of the ancestral metazoan genome reveals an increase in genomic novelty. Nature Communications, 2018, 9, 1730.	5.8	101
21	Amphioxus functional genomics and the origins of vertebrate gene regulation. Nature, 2018, 564, 64-70.	13.7	224
22	Historical and current patterns of gene flow in the butterfly <i>Pararge aegeria</i>. Journal of Biogeography, 2018, 45, 1628-1639.	1.4	18
23	Mouse Obox and Crxos modulate preimplantation transcriptional profiles revealing similarity between paralogous mouse and human homeobox genes. EvoDevo, 2018, 9, 2.	1.3	13
24	Diversification of Hox Gene Clusters in Osteoglossomorph Fish in Comparison to Other Teleosts and the Spotted Gar Outgroup. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2017, 328, 638-644.	0.6	13
25	Lineage-specific rediploidization is a mechanism to explain time-lags between genome duplication and evolutionary diversification. Genome Biology, 2017, 18, 111.	3.8	136
26	New genes from old: asymmetric divergence of gene duplicates and the evolution of development. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20150480.	1.8	90
27	Novel and divergent genes in the evolution of placental mammals. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20171357.	1.2	23
28	A sister of <i>NANOG</i> regulates genes expressed in pre-implantation human development. Open Biology, 2017, 7, 170027.	1.5	13
29	Genome sequence of a diabetes-prone rodent reveals a mutation hotspot around the ParaHox gene cluster. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 7677-7682.	3.3	30
30	The dawn of amphioxus molecular biology - a personal perspective. International Journal of Developmental Biology, 2017, 61, 585-590.	0.3	4
31	Never Ending Analysis of a Century Old Evolutionary Debate: â€œUnringingâ€•the Urmetazoon Bell. Frontiers in Ecology and Evolution, 2016, 4, .	1.1	15
32	Evolutionary origin and functional divergence of totipotent cell homeobox genes in eutherian mammals. BMC Biology, 2016, 14, 45.	1.7	37
33	Diversity of human and mouse homeobox gene expression in development and adult tissues. BMC Developmental Biology, 2016, 16, 40.	2.1	20
34	Conservation, Duplication, and Divergence of Five Opsin Genes in Insect Evolution. Genome Biology and Evolution, 2016, 8, 579-587.	1.1	77
35	A single three-dimensional chromatin compartment in amphioxus indicates a stepwise evolution of vertebrate Hox bimodal regulation. Nature Genetics, 2016, 48, 336-341.	9.4	113
36	The spotted gar genome illuminates vertebrate evolution and facilitates human-teleost comparisons. Nature Genetics, 2016, 48, 427-437.	9.4	545

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37	Reinforcing the Egg-Timer: Recruitment of Novel Lophotrochozoa Homeobox Genes to Early and Late Development in the Pacific Oyster. Genome Biology and Evolution, 2015, 7, 677-688.	1.1	42
38	The Hox cluster microRNA miR-615: a case study of intronic microRNA evolution. EvoDevo, 2015, 6, 31.	1.3	16
39	A Burst of miRNA Innovation in the Early Evolution of Butterflies and Moths. Molecular Biology and Evolution, 2015, 32, 1161-1174.	3.5	30
40	Did homeobox gene duplications contribute to the Cambrian explosion?. Zoological Letters, 2015, 1, 1.	0.7	45
41	Scenarios for the making of vertebrates. Nature, 2015, 520, 450-455.	13.7	51
42	Cdx ParaHox genes acquired distinct developmental roles after gene duplication in vertebrate evolution. BMC Biology, 2015, 13, 56.	1.7	12
43	A Diversity of Conserved and Novel Ovarian MicroRNAs in the Speckled Wood (Pararge aegeria). PLoS ONE, 2015, 10, e0142243.	1.1	21
44	Strepsiptera, Phylogenomics and the Long Branch Attraction Problem. PLoS ONE, 2014, 9, e107709.	1.1	51
45	Ancient Expansion of the Hox Cluster in Lepidoptera Generated Four Homeobox Genes Implicated in Extra-Embryonic Tissue Formation. PLoS Genetics, 2014, 10, e1004698.	1.5	58
46	Genomic organisation of the seven ParaHox genes of coelacanths. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2014, 322, 352-358.	0.6	2
47	Enigmatic Orthology Relationships between Hox Clusters of the African Butterfly Fish and Other Teleosts Following Ancient Whole-Genome Duplication. Molecular Biology and Evolution, 2014, 31, 2592-2611.	3.5	37
48	Discovery and Classification of Homeobox Genes in Animal Genomes. Methods in Molecular Biology, 2014, 1196, 3-18.	0.4	4
49	A family of diatom-like silicon transporters in the siliceous loricate choanoflagellates. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20122543.	1.2	39
50	The genomes of four tapeworm species reveal adaptations to parasitism. Nature, 2013, 496, 57-63.	13.7	603
51	How are comparative genomics and the study of microRNAs changing our views on arthropod endocrinology and adaptations to the environment?. General and Comparative Endocrinology, 2013, 188, 16-22.	0.8	19
52	Evolution of homeobox genes. Wiley Interdisciplinary Reviews: Developmental Biology, 2013, 2, 31-45.	5.9	232
53	Extensive Chordate and Annelid Macrosynteny Reveals Ancestral Homeobox Gene Organization. Molecular Biology and Evolution, 2012, 29, 157-165.	3.5	53
54	A genome-wide view of transcription factor gene diversity in chordate evolution: less gene loss in amphioxus?. Briefings in Functional Genomics, 2012, 11, 177-186.	1.3	36

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55	The oyster genome reveals stress adaptation and complexity of shell formation. Nature, 2012, 490, 49-54.	13.7	1,966
56	Evolution of the Alx homeobox gene family: parallel retention and independent loss of the vertebrate Alx3 gene. Evolution & Development, 2011, 13, 343-351.	1.1	37
57	<scp>HomeoDB2</scp>: functional expansion of a comparative homeobox gene database for evolutionary developmental biology. Evolution & Development, 2011, 13, 567-568.	1.1	110
58	The hypoxiaâ€inducible transcription factor pathway regulates oxygen sensing in the simplest animal, <i>Trichoplax adhaerens</i>. EMBO Reports, 2011, 12, 63-70.	2.0	210
59	The dynamics of vertebrate homeobox gene evolution: gain and loss of genes in mouse and human lineages. BMC Evolutionary Biology, 2011, 11, 169.	3.2	44
60	The origin and evolution of ARGFX homeobox loci in mammalian radiation. BMC Evolutionary Biology, 2010, 10, 182.	3.2	10
61	From genomes to morphology: a view from amphioxus. Acta Zoologica, 2010, 91, 81-86.	0.6	10
62	Ancient homeobox gene loss and the evolution of chordate brain and pharynx development: deductions from amphioxus gene expression. Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 3381-3389.	1.2	11
63	Parallel Retention of Pdx2 Genes in Cartilaginous Fish and Coelacanths. Molecular Biology and Evolution, 2010, 27, 2386-2391.	3.5	18
64	Degenerate evolution of the hedgehog gene in a hemichordate lineage. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7491-7494.	3.3	12
65	Comprehensive survey and classification of homeobox genes in the genome of amphioxus, Branchiostoma floridae. Development Genes and Evolution, 2008, 218, 579-590.	0.4	69
66	Asymmetry in a pterobranch hemichordate and the evolution of leftâ€“right patterning. Developmental Dynamics, 2008, 237, 3634-3639.	0.8	37
67	Developmental biology of pterobranch hemichordates: History and perspectives. Genesis, 2008, 46, 587-591.	0.8	35
68	The origins of graptolites and other pterobranchs: a journey from â€˜Polyzoaâ€™. Lethaia, 2008, 41, 303-316.	0.6	15
69	The amphioxus genome and the evolution of the chordate karyotype. Nature, 2008, 453, 1064-1071.	13.7	1,496
70	HomeoDB: a database of homeobox gene diversity. Evolution & Development, 2008, 10, 516-518.	1.1	75
71	Do cnidarians have a ParaHox cluster? Analysis of synteny around a <i>Nematostella</i> homeobox gene cluster. Evolution & Development, 2008, 10, 725-730.	1.1	33
72	Protein evolution of ANTP and PRD homeobox genes. BMC Evolutionary Biology, 2008, 8, 200.	3.2	7

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73	The Urbilaterian Super-Hox cluster. Trends in Genetics, 2008, 24, 259-262.	2.9	43
74	Correlating Bayesian date estimates with climatic events and domestication using a bovine case study. Biology Letters, 2008, 4, 370-374.	1.0	70
75	The amphioxus genome illuminates vertebrate origins and cephalochordate biology. Genome Research, 2008, 18, 1100-1111.	2.4	456
76	Protochordates. Methods in Molecular Biology, 2008, 461, 563-566.	0.4	1
77	Wholemount In Situ Hybridization to Amphioxus Embryos. Methods in Molecular Biology, 2008, 461, 703-706.	0.4	6
78	The butterfly Danaus chrysippus (Lepidoptera: Nymphalidae) in Kenya is variably infected with respect to genotype and body size by a maternally transmitted male-killing endosymbiont (Spiroplasma). International Journal of Tropical Insect Science, 2007, 27, 62.	0.4	21
79	A Degenerate ParaHox Gene Cluster in a Degenerate Vertebrate. Molecular Biology and Evolution, 2007, 24, 2681-2686.	3.5	34
80	Annotation, nomenclature and evolution of four novel homeobox genes expressed in the human germ line. Gene, 2007, 387, 7-14.	1.0	41
81	Classification and nomenclature of all human homeobox genes. BMC Biology, 2007, 5, 47.	1.7	322
82	The origins of multicellularity: a multi-taxon genome initiative. Trends in Genetics, 2007, 23, 113-118.	2.9	201
83	Origin and evolution of a myxozoan worm. Integrative and Comparative Biology, 2007, 47, 752-758.	0.9	19
84	<i>Buddenbrockia</i> Is a Cnidarian Worm. Science, 2007, 317, 116-118.	6.0	151
85	A Gbx homeobox gene in amphioxus: Insights into ancestry of the ANTP class and evolution of the midbrain/hindbrain boundary. Developmental Biology, 2006, 295, 40-51.	0.9	98
86	Patterns of conservation and change in honey bee developmental genes. Genome Research, 2006, 16, 1376-1384.	2.4	139
87	An unusual choanoflagellate protein released by Hedgehog autocatalytic processing. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 401-407.	1.2	56
88	Breakup of a homeobox cluster after genome duplication in teleosts. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 10369-10372.	3.3	68
89	The evolution of homeobox genes: Implications for the study of brain development. Brain Research Bulletin, 2005, 66, 484-490.	1.4	75
90	No more than 14: the end of the amphioxus Hox cluster. International Journal of Biological Sciences, 2005, 1, 19-23.	2.6	63

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91	DEVELOPMENTAL BIOLOGY: Enhanced: The Ups and Downs of a Sea Anemone. Science, 2004, 304, 1255-1256.	6.0	7
92	Amphioxus and ascidian Dmbx homeobox genes give clues to the vertebrate origins of midbrain development. Development (Cambridge), 2004, 131, 3285-3294.	1.2	71
93	Polyploidy in vertebrate ancestry: Ohno and beyond. Biological Journal of the Linnean Society, 2004, 82, 425-430.	0.7	71
94	The Trox-2 Hox/ParaHox gene of Trichoplax (Placozoa) marks an epithelial boundary. Development Genes and Evolution, 2004, 214, 170-175.	0.4	110
95	An antecedent of the MHC-linked genomic region in amphioxus. Immunogenetics, 2004, 55, 782-784.	1.2	38
96	Eleven daughters of NANOGâ˜†. Genomics, 2004, 84, 229-238.	1.3	129
97	Phylogenomics of Eukaryotes: Impact of Missing Data on Large Alignments. Molecular Biology and Evolution, 2004, 21, 1740-1752.	3.5	371
98	More genes in vertebrates?. Journal of Structural and Functional Genomics, 2003, 3, 75-84.	1.2	59
99	The evolution of multicellularity and early animal genomes. Current Opinion in Genetics and Development, 2003, 13, 599-603.	1.5	34
100	Dispersal of NK homeobox gene clusters in amphioxus and humans. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 5292-5295.	3.3	81
101	More genes in vertebrates?. , 2003, , 75-84.		2
102	More genes in vertebrates?. Journal of Structural and Functional Genomics, 2003, 3, 75-84.	1.2	23
103	Were vertebrates octoploid?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2002, 357, 531-544.	1.8	200
104	Orphan Worm Finds a Home: Buddenbrockia is a Myxozoan. Molecular Biology and Evolution, 2002, 19, 968-971.	3.5	63
105	Bayesian Phylogenetic Analysis Supports Monophyly of Ambulacraria and of Cyclostomes. Zoological Science, 2002, 19, 593-599.	0.3	137
106	Exploiting genomics in evolutionary developmental biology. International Congress Series, 2002, 1246, 217-229.	0.2	0
107	Ciona. Current Biology, 2002, 12, R609.	1.8	2
108	An orphan PRD class homeobox gene expressed in mouse brain and limb development. Development Genes and Evolution, 2002, 212, 293-297.	0.4	11

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109	Ciona intestinalis ParaHox genes: evolution of Hox/ParaHox cluster integrity, developmental mode, and temporal colinearity. Molecular Phylogenetics and Evolution, 2002, 24, 412-417.	1.2	92
110	The Mnx homeobox gene class defined by HB9 , MNR2 and amphioxus AmphiMnx. Development Genes and Evolution, 2001, 211, 103-107.	0.4	54
111	Beyond the Hox: how widespread is homeobox gene clustering?. Journal of Anatomy, 2001, 199, 13-23.	0.9	77
112	Ancient origin of the Hox gene cluster. Nature Reviews Genetics, 2001, 2, 33-38.	7.7	233
113	Hsp70 sequences indicate that choanoflagellates are closely related to animals. Current Biology, 2001, 11, 967-970.	1.8	72
114	Beyond the Hox: how widespread is homeobox gene clustering?. Journal of Anatomy, 2001, 199, 13-23.	0.9	32
115	An Amphioxus Emx Homeobox Gene Reveals Duplication During Vertebrate Evolution. Molecular Biology and Evolution, 2000, 17, 1520-1528.	3.5	28
116	The amphioxus Hox cluster: deuterostome posterior flexibility andHox14. Evolution & Development, 2000, 2, 284-293.	1.1	156
117	Conservation and elaboration of Hox gene regulation during evolution of the vertebrate head. Nature, 2000, 408, 854-857.	13.7	167
118	Evidence for 14 homeobox gene clusters in human genome ancestry. Current Biology, 2000, 10, 1059-1062.	1.8	177
119	Rare genomic changes as a tool for phylogenetics. Trends in Ecology and Evolution, 2000, 15, 454-459.	4.2	616
120	The future of evolutionary developmental biology. Nature, 1999, 402, C41-C44.	13.7	81
121	Dicyemids are higher animals. Nature, 1999, 401, 762-762.	13.7	51
122	Amphioxus type I keratin cDNA and the evolution of intermediate filament genes. , 1999, 285, 50-56.		20
123	Colinear and Segmental Expression of Amphioxus Hox Genes. Developmental Biology, 1999, 213, 131-141.	0.9	147
124	Introduction: Gene duplication in development and evolution. Seminars in Cell and Developmental Biology, 1999, 10, 515-516.	2.3	5
125	Gene duplication: Past, present and future. Seminars in Cell and Developmental Biology, 1999, 10, 541-547.	2.3	170
126	The ParaHox gene cluster is an evolutionary sister of the Hox gene cluster. Nature, 1998, 392, 920-922.	13.7	445

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127	Major Transitions in Animal Evolution: A Developmental Genetic Perspective. American Zoologist, 1998, 38, 829-842.	0.7	67
128	Neural Tube Is Partially Dorsalized by Overexpression ofHrPax-37:The Ascidian Homologue ofPax-3andPax-7. Developmental Biology, 1997, 187, 240-252.	0.9	109
129	Cloning and analysis of an HMG gene from the lamprey Lampetra fluviatilis: gene duplication in vertebrate evolution. Gene, 1997, 184, 99-105.	1.0	34
130	Vertebrate evolution: Something fishy about Hox genes. Current Biology, 1997, 7, R570-R572.	1.8	44
131	Evolution of 28S Ribosomal DNA in Chaetognaths: Duplicate Genes and Molecular Phylogeny. Journal of Molecular Evolution, 1997, 44, 135-144.	0.8	77
132	HoxGenes and Chordate Evolution. Developmental Biology, 1996, 173, 382-395.	0.9	433
133	Old head on young shoulders. Nature, 1996, 383, 490-490.	13.7	111
134	Origin of patterning in neural tubes. Nature, 1996, 384, 123-123.	13.7	78
135	Conservation, Duplication, and Divergence of Developmental Genes During Chordate Evolution. Animal Biology, 1995, 46, 47-67.	0.4	69
136	Archetypal organization of the amphioxus Hox gene cluster. Nature, 1994, 370, 563-566.	13.7	550
137	Gene duplications and the origins of vertebrate development. Development (Cambridge), 1994, 1994, 125-133.	1.2	610
138	Development of the zootype. Nature, 1993, 363, 307-308.	13.7	4
139	Molecular evolution and diversification of the vestimentiferan tube worms. Journal of the Marine Biological Association of the United Kingdom, 1993, 73, 437-452.	0.4	55
140	Mice and flies head to head. Nature, 1992, 358, 627-628.	13.7	46
141	Problems and paradigms: Hoemeobox genes in vertebrate evolution. BioEssays, 1992, 14, 267-273.	1.2	131
142	Cloning and evolutionary analysis of msh-like homeobox genes from mouse, zebrafish and ascidian. Gene, 1991, 98, 253-257.	1.0	123
143	Cloning of fish zinc-finger genes related to Krox-20 and Krox-24. Biochemical and Biophysical Research Communications, 1991, 179, 1220-1224.	1.0	13
144	Cloning of segment polarity gene homologues from the unsegmented brachiopodTerebratulina retusa(Linnaeus). FEBS Letters, 1991, 291, 211-213.	1.3	15

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145	A novel avian W chromosome DNA repeat sequence in the lesser black-backed gull (Larus fuscus). Chromosoma, 1990, 99, 243-250.	1.0	34
146	Conservation ofengrailed-like homeobox sequences during vertebrate evolution. FEBS Letters, 1990, 277, 250-252.	1.3	48
147	Pursuing the functions of vertebrate homeobox genes: Progress and prospects. Trends in Neurosciences, 1989, 12, 206-209.	4.2	7
148	Homeobox genes and the vertebrate head. Development (Cambridge), 1988, 103, 17-24.	1.2	25
149	Phylogenetic distribution of Antennapedia-like homoeo boxes. Nature, 1986, 321, 251-253.	13.7	85
150	The genome sequence of the poplar hawk-moth, Laothoe populi Â(Linnaeus, 1758). Wellcome Open Research, 0, 6, 237.	0.9	2
151	The genome sequence of the snout, Hypena proboscidalisÂ(Linnaeus, 1758). Wellcome Open Research, 0, 6, 236.	0.9	0
152	The genome sequence of Svenssonâ€™s copper underwing, Amphipyra berbera Rungs, 1949. Wellcome Open Research, 0, 6, 314.	0.9	0
153	The genome sequence of the buff-tip, Phalera bucephala (Linnaeus, 1758). Wellcome Open Research, 0, 7, 28.	0.9	1
154	The genome sequence of the square-spot rustic, Xestia xanthographa (Schiffermuller, 1775). Wellcome Open Research, 0, 7, 37.	0.9	2
155	The genome sequence of the iron prominent, Notodonta dromedarius (Linnaeus, 1767). Wellcome Open Research, 0, 6, 341.	0.9	0
156	The genome sequence of the angle shades moth, Phlogophora meticulosa (Linnaeus, 1758). Wellcome Open Research, 0, 7, 89.	0.9	0
157	The genome sequence of the silver Y moth, Autographa gamma (Linnaeus, 1758). Wellcome Open Research, 0, 7, 100.	0.9	1
158	The genome sequence of the large yellow underwing, Noctua pronuba (Linnaeus, 1758). Wellcome Open Research, 0, 7, 119.	0.9	0
159	The genome sequence of the Clifden nonpareil, Catocala fraxini (Linnaeus, 1758). Wellcome Open Research, 0, 7, 129.	0.9	0
160	The genome sequence of the broad-bordered yellow underwing, Noctua fimbriata (Schreber, 1759). Wellcome Open Research, 0, 6, 345.	0.9	1
161	The genome sequence of the yellow-tail moth, Euproctis similisÂ(Fuessly, 1775). Wellcome Open Research, 0, 6, 227.	0.9	1
162	The genome sequence of the swallow prominent, Pheosia tremula (Clerck, 1759). Wellcome Open Research, 0, 6, 335.	0.9	1

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163	The genome sequence of the bramble shoot moth, Notocelia uddmanniana (Linnaeus, 1758). Wellcome Open Research, 0, 6, 348.	0.9	0
164	The genome sequence of the dun-bar pinion, Cosmia trapezina (Linnaeus, 1758). Wellcome Open Research, 0, 7, 189.	0.9	2
165	The genome sequence of the clay, Mythimna ferrago (Fabricius, 1787). Wellcome Open Research, 0, 7, 177.	0.9	2