

# Lei Yang

## List of Publications by Year in descending order

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27  
papers

1,040  
citations

567144

15  
h-index

580701

25  
g-index

27  
all docs

27  
docs citations

27  
times ranked

1043  
citing authors

#	ARTICLE	IF	CITATIONS
1	Deciphering reticulate evolution of the largest group of polyploid vertebrates, the subfamily cyprininae (Teleostei: Cypriniformes). <i>Molecular Phylogenetics and Evolution</i> , 2022, 166, 107323.	1.2	12
2	Molecular phylogeny and divergence of major clades in the <i>Puntius</i> complex (Teleostei: Cypriniformes). <i>Molecular Phylogenetics and Evolution</i> , 2021, 77, 107323.	0.7	9
3	DNA from Tooth Embedded in Man's Foot Resolves Quarter-Century-Old Shark Bite Mystery. <i>Wilderness and Environmental Medicine</i> , 2019, 30, 335-337.	0.4	3
4	Phylogenetic relationships of Cypriniformes and plasticity of pharyngeal teeth in the adaptive radiation of cyprinids. <i>Science China Life Sciences</i> , 2019, 62, 553-565.	2.3	29
5	Phylogeny of the manta and devilrays (Chondrichthyes: mobulidae), with an updated taxonomic arrangement for the family. <i>Zoological Journal of the Linnean Society</i> , 2018, 182, 50-75.	1.0	113
6	Phylogenetic relationships and classification of the Holarctic family Leuciscidae (Cypriniformes: Cyprinidae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 85, 107323.	1.2	98
7	Historical introgression drives pervasive mitochondrial admixture between two species of pelagic sharks. <i>Molecular Phylogenetics and Evolution</i> , 2017, 110, 122-126.	1.2	24
8	Complete mitochondrial genome of the blacknose shark <i>Carcharhinus acronotus</i> (Elasmobranchii: Carcharhinidae). <i>Mitochondrial DNA Part B</i> , 2017, 22, 107323.	0.6	4
9	Long-PCR based next generation sequencing of the whole mitochondrial genome of the peacock skate <i>Pavoraja nitida</i> (Elasmobranchii: Arhynchobatidae). <i>Mitochondrial DNA</i> , 2016, 27, 943-944.	0.6	4
10	Determination of complete mitochondrial genome sequence from the holotype of the southern Mandarin dogfish <i>Cirrhigaleus australis</i> (Elasmobranchii: Squalidae). <i>Mitochondrial DNA</i> , 2016, 27, 593-594.	0.6	3
11	Mitogenomic sequence and phylogenetic placement of the Hurtle's whipray <i>Himantura hortlei</i> (Elasmobranchii: Dasyatidae). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016, 27, 2437-2439.	0.7	1
12	Phylogeny and polyploidy: Resolving the classification of cyprinine fishes (Teleostei: Cypriniformes). <i>Molecular Phylogenetics and Evolution</i> , 2015, 85, 97-116.	1.2	218
13	Rediscovery of the Threatened River Sharks, <i>Glyphis garricki</i> and <i>G. glyphis</i> , in Papua New Guinea. <i>PLoS ONE</i> , 2015, 10, e0140075.	1.1	39
14	Stability versus diversity of the dentition during evolutionary radiation in cyprinine fish. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20132688.	1.2	12
15	Limits and phylogenetic relationships of East Asian fishes in the subfamily Oxygastrinae (Teleostei: Cypriniformes). <i>Molecular Phylogenetics and Evolution</i> , 2014, 90, 107323.	0.2	23
16	Molecular phylogeny of the cyprinid tribe Labeonini (Teleostei: Cypriniformes). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 362-379.	1.2	66
17	Phylogenetic placements of the barbin genera <i>Discherodontus</i> , <i>Chagunius</i> , and <i>Hypselobarbus</i> in the subfamily Cyprininae (Teleostei: Cypriniformes) and their relationships with other barbines. <i>Zootaxa</i> , 2012, 3586, 26.	0.2	16
18	GONORYNCHIFORMES AND OSTARIOPHYSAN RELATIONSHIPS: A COMPREHENSIVE REVIEW - Edited by T. Grande, F. J. Poyato-Ariza and R. Diogo. <i>Journal of Fish Biology</i> , 2011, 78, 1277-1278.	0.7	0

#	ARTICLE	IF	CITATIONS
19	Phylogeny of the gudgeons (Teleostei: Cyprinidae: Gobioninae). <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 103-124.	1.2	81
20	Phylogenetic relationships, subdivision, and biogeography of the cyprinid tribe Labeonini (sensu Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 7 structures in the labeonin classification. <i>Molecular Phylogenetics and Evolution</i> , 2010, 54, 254-265.	1.2	44
21	Molecular phylogeny of the fishes traditionally referred to Cyprinini sensu stricto (Teleostei: Tj ETQq1 1 0.784314 rgBT /Overlock 10 0.7 37	0.7	37
22	Threatened fishes of the world: <i>Cranoglanis boudierus</i> (Richardson, 1846) (Cranoglanididae). <i>Environmental Biology of Fishes</i> , 2009, 84, 157-158.	0.4	1
23	Threatened fishes of the world: <i>Procypris rabaudi</i> (Tchang, 1930) (Cyprinidae). <i>Environmental Biology of Fishes</i> , 2009, 84, 275-276.	0.4	10
24	Reconstructing the phylogenetic relationships of the earth's most diverse clade of freshwater fishes—order Cypriniformes (Actinopterygii: Ostariophysi): A case study using multiple nuclear loci and the mitochondrial genome. <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 500-514.	1.2	129
25	Population genetic structure and geographical differentiation of the Chinese catfish <i>Hemibagrus macropterus</i> (Siluriformes, Bagridae): Evidence for altered drainage patterns. <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 405-411.	1.2	29
26	Phylogeography of the freshwater catfish <i>Hemibagrus guttatus</i> (Siluriformes, Bagridae): Implications for South China biogeography and influence of sea-level changes. <i>Molecular Phylogenetics and Evolution</i> , 2008, 49, 393-398.	1.2	29
27	Taxonomic Identification of Two Poorly Known Lantern Shark Species Based on Mitochondrial DNA From Wet-Collection Paratypes. <i>Frontiers in Ecology and Evolution</i> , 0, 10, .	1.1	6