

Diogo Hashimoto

List of Publications by Year in descending order

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72
papers

1,307
citations

361296

20
h-index

434063

31
g-index

74
all docs

74
docs citations

74
times ranked

1023
citing authors

#	ARTICLE	IF	CITATIONS
1	Delimiting the Origin of a B Chromosome by FISH Mapping, Chromosome Painting and DNA Sequence Analysis in <i>Astyanax paranae</i> (Teleostei, Characiformes). PLoS ONE, 2014, 9, e94896.	1.1	85
2	Interspecific fish hybrids in Brazil: management of genetic resources for sustainable use. Reviews in Aquaculture, 2012, 4, 108-118.	4.6	61
3	Identification of Bacterial Fish Pathogens in Brazil by Direct Colony PCR and 16S rRNA Gene Sequencing. Advances in Microbiology, 2015, 05, 409-424.	0.3	61
4	Chromosome Mapping of H1 Histone and 5S rRNA Gene Clusters in Three Species of <i>Astyanax</i> (Teleostei, Characiformes). Cytogenetic and Genome Research, 2011, 134, 64-71.	0.6	52
5	Chromosomal organization of repetitive DNA sequences in <i>Astyanax bockmanni</i> (Teleostei, Characiformes). Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 329-336.	0.5	51
6	Uncovering the Ancestry of B Chromosomes in <i>Moenkhausia sanctaefilomenae</i> (Teleostei, Characidae). PLoS ONE, 2016, 11, e0150573.	1.1	48
7	Molecular diagnostic methods for identifying Serrasalmid fish (Pacu, Pirapitinga, and Tambaqui) and their hybrids in the Brazilian aquaculture industry. Aquaculture, 2011, 321, 49-53.	1.7	42
8	High-throughput analysis unveils a highly shared satellite DNA library among three species of fish genus <i>Astyanax</i> . Scientific Reports, 2017, 7, 12726.	1.6	40
9	Satellite landscape analysis of <i>Megaleporinus macrocephalus</i> (Teleostei, Anostomidae) reveals intense accumulation of satellite sequences on the heteromorphic sex chromosome. Scientific Reports, 2019, 9, 5856.	1.6	40
10	Genetic Identification of F1 and Post-F1 Serrasalmid Juvenile Hybrids in Brazilian Aquaculture. PLoS ONE, 2014, 9, e89902.	1.1	34
11	The farming and husbandry of <i>Colossoma macropomum</i> : From Amazonian waters to sustainable production. Reviews in Aquaculture, 2022, 14, 993-1027.	4.6	33
12	Identification of the shark species <i>Rhizoprionodon landii</i> and <i>R. porosus</i> (Elasmobranchii, Carcharhinidae) by multiplex PCR and PCR-RFLP techniques. Molecular Ecology Resources, 2009, 9, 771-773.	2.2	32
13	Detection of post-F1 fish hybrids in broodstock using molecular markers: approaches for genetic management in aquaculture. Aquaculture Research, 2013, 44, 876-884.	0.9	30
14	Cytogenetic markers as diagnoses in the identification of the hybrid between <i>Piauí</i> (<i>Leporinus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 0.6 29	0.6	29
15	Identification of hybrids between Neotropical fish <i>Leporinus macrocephalus</i> and <i>Leporinus elongatus</i> by PCR-RFLP and multiplex-PCR: Tools for genetic monitoring in aquaculture. Aquaculture, 2010, 298, 346-349.	1.7	28
16	Detection of hybrids and genetic introgression in wild stocks of two catfish species (Siluriformes: <i>Leporinus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 0.9 28	0.9	28
17	Origin of B chromosomes in the genus <i>Astyanax</i> (Characiformes, Characidae) and the limits of chromosome painting. Molecular Genetics and Genomics, 2016, 291, 1407-1418.	1.0	28
18	Molecular identification of hybrids between Neotropical catfish species <i>Pseudoplatystoma corruscans</i> and <i>Pseudoplatystoma reticulatum</i> . Aquaculture Research, 2011, 42, 1890-1894.	0.9	27

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19	A Glimpse into the Satellite DNA Library in Characidae Fish (Teleostei, Characiformes). <i>Frontiers in Genetics</i> , 2017, 8, 103.	1.1	27
20	Hybridization and genetic introgression patterns between two South American catfish along their sympatric distribution range. <i>Hydrobiologia</i> , 2017, 788, 319-343.	1.0	23
21	Genetic identification of lamniform and carcharhiniform sharks using multiplex-PCR. <i>Conservation Genetics Resources</i> , 2010, 2, 31-35.	0.4	22
22	Genetic parameters for resistance to <i>Aeromonas hydrophila</i> in the Neotropical fish pacu (<i>Piaractus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.7	21
23	SNP discovery from liver transcriptome in the fish <i>Piaractus mesopotamicus</i> . <i>Conservation Genetics Resources</i> , 2016, 8, 109-114.	0.4	20
24	First report of a B chromosome in a natural population of <i>Astyanax altiparanae</i> (Characiformes,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.6	20
25	Assessing Genetic Diversity for a Pre-Breeding Program in <i>Piaractus mesopotamicus</i> by SNPs and SSRs. <i>Genes</i> , 2019, 10, 668.	1.0	19
26	Chromosome polymorphism of heterochromatin and nucleolar regions in two populations of the fish <i>Astyanax bockmanni</i> (Teleostei: Characiformes). <i>Neotropical Ichthyology</i> , 2010, 8, 861-866.	0.5	18
27	Cytogenetic Mapping of H1 Histone and Ribosomal RNA Genes in Hybrids between Catfish Species <i>Pseudoplatystoma corruscans</i> and <i>Pseudoplatystoma reticulatum</i>. <i>Cytogenetic and Genome Research</i> , 2013, 139, 102-106.	0.6	18
28	Motile <i>Aeromonas</i> septicemia in tambaqui <i>Colossoma macropomum</i> : Pathogenicity, lethality and new insights for control and disinfection in aquaculture. <i>Microbial Pathogenesis</i> , 2020, 149, 104512.	1.3	17
29	Chromosomal features of nucleolar dominance in hybrids between the Neotropical fish <i>Leporinus macrocephalus</i> and <i>Leporinus elongatus</i> (Characiformes, Anostomidae). <i>Genetica</i> , 2009, 137, 135-140.	0.5	16
30	Genetic markers for the identification of hybrids among catfish species of the family Pimelodidae. <i>Journal of Applied Ichthyology</i> , 2013, 29, 643-647.	0.3	16
31	Genetic (co)variation between resistance to <i>Aeromonas hydrophila</i> and growth in tambaqui (<i>Colossoma macropomum</i>). <i>Aquaculture</i> , 2020, 523, 735225.	1.7	16
32	Forensic identification of the guitarfish species <i>Rhinobatos horkelli</i>, <i>R. percellens</i> and <i>Zapteryx brevirostris</i> using multiplexâ€PCR. <i>Molecular Ecology Resources</i> , 2010, 10, 197-199.	2.2	15
33	Innovative molecular approach to the identification of <i>Colossoma macropomum</i> and its hybrids. <i>Anais Da Academia Brasileira De Ciencias</i> , 2012, 84, 517-526.	0.3	15
34	Survival of purebred and hybrid Serrasalminidae under low water temperature conditions. <i>Aquaculture</i> , 2018, 497, 97-102.	1.7	15
35	Repetitive DNA probe linked to sex chromosomes in hybrids between Neotropical fish <i>Leporinus macrocephalus</i> and <i>Leporinus elongatus</i>(Characiformes,) Tj ETQq1 1 0.784314rgBT /Over	0.7	14
36	Cytogenetic analysis of B chromosomes in one population of the fish <i>Moenkhausia sanctaefilomenae</i> (Steindachner, 1907) (Teleostei, Characiformes). <i>Comparative Cytogenetics</i> , 2012, 6, 141-151.	0.3	14

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37	Characterization of lipid metabolism genes and the influence of fatty acid supplementation in the hepatic lipid metabolism of dusky grouper (<i>Epinephelus marginatus</i>). <i>Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology</i> , 2018, 219-220, 1-9.	0.8	13
38	Cytogenetic characterization of distinct B chromosomes in a population of the fish <i>Astyanax bockmanni</i> (Teleostei, Characiformes). <i>Caryologia</i> , 2012, 65, 229-233.	0.2	12
39	Microsatellites Associated with Growth Performance and Analysis of Resistance to <i>Aeromonas hydrophila</i> in Tambaqui <i>Colossoma macropomum</i> . <i>Frontiers in Genetics</i> , 2018, 9, 3.	1.1	12
40	Genetic Characterization of the Fish <i>Piaractus brachipomus</i> by Microsatellites Derived from Transcriptome Sequencing. <i>Frontiers in Genetics</i> , 2018, 9, 46.	1.1	12
41	Development of a multi-species SNP array for serrasalmid fish <i>Colossoma macropomum</i> and <i>Piaractus mesopotamicus</i> . <i>Scientific Reports</i> , 2021, 11, 19289.	1.6	12
42	Identification and characterization of polymorphic microsatellite loci in the blue shark <i>Prionace glauca</i> , and cross-species amplification in other shark species. <i>Journal of Fish Biology</i> , 2012, 80, 2643-2646.	0.7	11
43	Molecular identification of intergenus crosses involving catfish hybrids: risks for aquaculture production. <i>Neotropical Ichthyology</i> , 2016, 14, .	0.5	11
44	Development of microsatellite markers using next-generation sequencing for the fish <i>Colossoma macropomum</i> . <i>Molecular Biology Reports</i> , 2018, 45, 9-18.	1.0	11
45	Development and characterization of microsatellite loci in <i>Phractocephalus hemiliopterus</i> (Siluriformes: Pimelodidae) and their cross-species amplification in six related species. <i>Conservation Genetics Resources</i> , 2012, 4, 499-501.	0.4	9
46	Organization and Chromosomal Distribution of Histone Genes and Transposable Elements in the Genome of <i>Astyanax bockmanni</i> (Teleostei, Characiformes). <i>Cytogenetic and Genome Research</i> , 2015, 146, 311-318.	0.6	9
47	New Insights into Karyotypic Relationships Among Populations of <i>Astyanax bockmanni</i> (Teleostei,) Tj ETQq1 1 0.784314 rgBT ₉ /Overlook	0.5	9
48	Isolation and characterization of microsatellite loci in the Neotropical fish <i>Astyanax altiparanae</i> (Teleostei: Characiformes) and cross-species amplification. <i>Journal of Genetics</i> , 2014, 93, 24-27.	0.4	8
49	Development of a SNP linkage map and genome-wide association study for resistance to <i>Aeromonas hydrophila</i> in pacu (<i>Piaractus mesopotamicus</i>). <i>BMC Genomics</i> , 2020, 21, 672.	1.2	8
50	Transcriptome Profiling of Pacu (<i>Piaractus mesopotamicus</i>) Challenged With Pathogenic <i>Aeromonas hydrophila</i> : Inference on Immune Gene Response. <i>Frontiers in Genetics</i> , 2020, 11, 604.	1.1	8
51	Genotype by environment interaction and genetic parameters for growth traits in the Neotropical fish pacu (<i>Piaractus mesopotamicus</i>). <i>Aquaculture</i> , 2021, 530, 735933.	1.7	8
52	Long-term persistence of supernumerary B chromosomes in multiple species of <i>Astyanax</i> fish. <i>BMC Biology</i> , 2021, 19, 52.	1.7	8
53	Quantitative genetic variation for resistance to the parasite <i>Ichthyophthirius multifiliis</i> in the Neotropical fish tambaqui (<i>Colossoma macropomum</i>). <i>Aquaculture Reports</i> , 2020, 17, 100338.	0.7	7
54	Imputation of genetic composition for missing pedigree data in Serrasalminae using morphometric data. <i>Scientia Agricola</i> , 2017, 74, 443-449.	0.6	6

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55	General and specific combining ability in Serrasalminae. <i>Aquaculture Research</i> , 2019, 50, 717.	0.9	6
56	Patterns of the innate immune response in tambaqui <i>Colossoma macropomum</i> : Modulation of gene expression in haemorrhagic septicaemia caused by <i>Aeromonas hydrophila</i> . <i>Microbial Pathogenesis</i> , 2021, 150, 104638.	1.3	6
57	Triploidy in tambaqui <i>Colossoma macropomum</i> identified by chromosomes of fish larvae. <i>Journal of Aquaculture & Marine Biology</i> , 2020, 9, 65-69.	0.2	6
58	Performance of tambacu hybrid ($\hat{\text{A}}^{\text{TM}}$, <i>Piaractus mesopotamicus</i> x $\hat{\text{A}}^{\text{TM}}$ <i>Colossoma macropomum</i>) and its parental pacu (<i>Piaractus mesopotamicus</i>) evaluated in cages under different feeding programmes. <i>Aquaculture Reports</i> , 2020, 17, 100355.	0.7	5
59	Genomic selection signatures in farmed <i>Colossoma macropomum</i> from tropical and subtropical regions in South America. <i>Evolutionary Applications</i> , 2022, 15, 679-693.	1.5	5
60	Linkage map for chromosome-level genome anchoring and genome-wide association study for resistance to <i>Aeromonas hydrophila</i> in <i>Colossoma macropomum</i> . <i>Aquaculture</i> , 2022, 560, 738462.	1.7	5
61	Molecular and morphological approaches for species delimitation and hybridization investigations of two <i>Cichla</i> species. <i>Iheringia - Serie Zoologia</i> , 2017, 107, .	0.5	4
62	Genetic Applications in the Conservation of Neotropical Freshwater Fish. , 0, , .		4
63	Cytogenetic markers as a tool for characterization of hybrids of <i>Astyanax Baird & Girard</i> , 1854 and <i>Hyphessobrycon Eigenmann</i> , 1907. <i>Comparative Cytogenetics</i> , 2020, 14, 231-242.	0.3	4
64	Organization and Distribution of Repetitive DNA Classes in the <i>Cichla kelberi</i> and <i>Cichla piquiti</i> Genome. <i>Cytologia</i> , 2017, 82, 193-197.	0.2	3
65	ETosis in tambaqui <i>Colossoma macropomum</i> : A programmed cell death pathway and approach of leukocytes immune response. <i>Microbial Pathogenesis</i> , 2021, 155, 104918.	1.3	3
66	Use of genetic markers to identify the illegal trade of billfish in the second largest fishing warehouse of Latin America. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2020, 30, 1251-1254.	0.9	2
67	Haplotypes traceability and genetic variability of the breeding population of pacu (<i>Piaractus</i>) Tj ETQq1 1 0.784314 $\frac{rgBT}{0.6}$ / Overlock 10		2
68	Morpho-molecular identification, pathogenicity for <i>Piaractus mesopotamicus</i> , and antimicrobial susceptibility of a virulent <i>Flavobacterium columnare</i> isolated from Nile tilapia cultured in Brazil. <i>Aquaculture</i> , 2022, 560, 738486.	1.7	2
69	Sex-Dependent Inheritance of B Chromosomes in <i>Psalidodon paranae</i> (Teleostei, Characiformes) Revealed by Directed Crossings. <i>Zebrafish</i> , 2021, 18, 363-368.	0.5	1
70	First Description of supernumerary Chromosomes in <i>Ictalurus punctatus</i> Rafinesque 1818 Reveals Active Ribosomal Genes in the B Complement. <i>Folia Biologica</i> , 2016, 64, 245-252.	0.1	0
71	DNA Barcode as an effective tool in the identification of billfishes (Scombroidei, Teleostei) from exported specimens. <i>Forensic Science International Animals and Environments</i> , 2021, 1, 100028.	0.3	0
72	Use of Molecular Genetic Methods to Reduce the Risk of Incorrect Identification of Fish Strains in Brazilian Aquaculture. <i>Frontiers in Genetics</i> , 2021, 12, 720736.	1.1	0