Tomasz StrzaÅ‚a

List of Publications by Year
in descending order

31 papers	169 citations	1162889 8 h-index	1281743 11 g-index
31 all docs	31 docs citations	31 times ranked	252 citing authors

#	Article	IF	Citations
1	Genetic diversity and relationship between cultivated, weedy and wild rye species as revealed by chloroplast and mitochondrial DNA non-coding regions analysis. PLoS ONE, 2019, 14, e0213023.	1.1	21
2	Agents of swimmerâ€™s itchâ€”dangerous minority in the Digenea invasion of Lymnaeidae in water bodies and the first report of Trichobilharzia regenti in Poland. Parasitology Research, 2018, 117, 3695-3704.	0.6	13
3	Morphological similarity and molecular divergence ofTrochulus striolatusandT. montanus, and their relationship to sympatric congeners (Gastropoda: Pulmonata: Hygromiidae). Systematics and Biodiversity, 2014, 12, 366-384.	0.5	11
4	Gene Polymorphisms of Novel Immunotolerant Molecule BTLA: Distribution of Alleles, Genotypes and Haplotypes in Polish Caucasian Population. Archivum Immunologiae Et Therapiae Experimentalis, 2015, 63, 73-78.	1.0	11
5	Phylogeography and postglacial colonization of Central Europe by Anguis fragilis and Anguis colchica. Amphibia - Reptilia, 2017, 38, 562-569.	0.1	11
6	Complete mitochondrial genome of endangered Socorro Conure (Aratinga brevipes) â€“ taxonomic position of the species and its relationship with Green Conure. Mitochondrial DNA, 2014, 25, 365-367.	0.6	10
7	Reintroduction of the European Capercaillie from the Capercaillie Breeding Centre in WisÅ‚a Forest District: Genetic Assessments of Captive and Reintroduced Populations. PLoS ONE, 2015, 10, e0145433.	1.1	10
8	Complete mitochondrial genome of Blue-crowned Parakeet (Aratinga acuticaudata)â€”phylogenetic position of the species among parrots group called Conures. Mitochondrial DNA, 2013, 24, 336-338.	0.6	9
9	Ongoing Speciation and Gene Flow between Taxonomically Challenging Trochulus Species Complex (Gastropoda: Hygromiidae). PLoS ONE, 2017, 12, e0170460.	1.1	8
10	Complete mitochondrial genome of endangered Maroon-fronted Parrot (Rhynchopsitta terrisi) â€“ conspecific relation of the species with Thick-billed Parrot (Rhynchopsitta pachyrhyncha). Mitochondrial DNA, 2014, 25, 424-426.	0.6	7
11	Low genetic variability of the edible dormouse (Glis glis) in Stołowe Mountains National Park (Poland)â€”preliminary results. Mammal Research, 2016, 61, 409-415.	0.6	7
12	Cepaea spp. as a source of Brachylaima mesostoma (Digenea: Brachylaimidae) and Brachylecithum sp. (Digenea: Dicrocoeliidae) larvae in Poland. Parasitology Research, 2020, 119, 145-152.	0.6	7
13	Phylogeny of Amazona barbadensis and the Yellow-Headed Amazon Complex (Aves: Psittacidae): A New Look at South American Parrot Evolution. PLoS ONE, 2014, 9, e97228.	1.1	6
14	Do aquatic barriers reduce male-mediated gene flow in a hybrid zone of the common shrew (Sorex) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.5	5
15	A microsatellite study in the ÅÄ™gucki MÅ‚yn/Popielno hybrid zone reveals no genetic differentiation between two chromosome races of the common shrew (Sorex araneus). Acta Theriologica, 2011, 56, 117-122.	1.1	5
16	Genetic structuring of the common shrew,Sorex araneus(Soricomorpha: Soricidae) in the Polish Sudetes may suggest ways of northwards colonization. Hereditas, 2012, 149, 197-206.	0.5	4
17	The first complete mitochondrial genome ofPyrrhurasp. â€“ question about conspecificity in the light of hybridization betweenPyrrhura molinaeandPyrrhura rupicolaspecies. Mitochondrial DNA, 2016, 27, 471-473.	0.6	4
18	Female-Male and Female-Female Social Interactions of Captive Kept Capercaillie (Tetrao Urogallus) and Its Consequences in Planning Breeding Programs. Animals, 2020, 10, 583.	1.0	4

#	Article	IF	Citations
19	Variation of the Common Shrew (Sorex Araneus L.) Dentition. Zoologica Poloniae: the Journal of Polish Zoological Society, 2008, 53, 49-56.	0.2	3
20	Complete mitochondrial genome of Mitred Conure (Psittacara mitratus): its comparison with mitogenome of Socorro Conure (Psittacara brevipes). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 3363-3364.	0.7	3
21	Salt mine microorganisms used for the biotransformation of chlorolactones. PLoS ONE, 2018, 13, e0197384.	1.1	3
22	Cytogenetic Examination of South American Tapirs, Tapirus Terrestris (Perissodactyla, Tapiridae), from the Wroclaw Zoological Garden. Vestnik Zoologii, 2015, 49, 529-536.	0.7	1
23	The first complete genome of â€˜trueâ€™ Aratinga genus in comparison to mitogenomes of other parrots from Arini tribe. Mitochondrial DNA Part B: Resources, 2016, 1, 853-855.	0.2	1
24	Complete mitochondrial genome of Red-throated Conure (Psittacara rubritorquis): its comparison with mitogenome of Socorro Conure (Psittacara brevipes). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 3354-3355.	0.7	1
25	Complete mitochondrial genome of Blue-and-yellow Macaw (Ara ararauna): the species morphologically similar to Blue-throated Macaw (Ara glaucogularis). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2017, 28, 307-308.	0.7	1
26	Complete mitochondrial genome of golden conure (Guaruba guarouba). Mitochondrial DNA Part B: Resources, 2017, 2, 33-34.	0.2	1
27	Characteristics of capercaillie (Tetrao urogallus) semen analysed with flow cytometry combined with fertility results. Reproduction in Domestic Animals, 2020, 55, 984-991.	0.6	1
28	Mitogenomes of Accipitriformes and Cathartiformes Were Subjected to Ancestral and Recent Duplications Followed by Gradual Degeneration. Genome Biology and Evolution, 2021, 13, .	1.1	1
29	Complete mitochondrial genome of the endemic legless lizard Anguis cephallonica Werner, 1894 and its comparison with mitogenome of Anguis fragilis Linnaeus, 1758. Mitochondrial DNA Part B: Resources, 2016, 1, 83-85.	0.2	0
30	Complete mitochondrial genome of the Eastern slow worm, Anguis colchica (Nordmann, 1840). Mitochondrial DNA Part B: Resources, 2017, 2, 67-68.	0.2	0
31	Complete mitochondrial genome of the Italian slow-worm Anguis veronensis Pollini, 1818, and its comparison with mitogenomes of other Anguis species. Mitochondrial DNA Part B: Resources, 2017, 2, 71-72.	0.2	0