

Tomasz StrzaÅ,a

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Mitogenomes of Accipitriformes and Cathartiformes Were Subjected to Ancestral and Recent Duplications Followed by Gradual Degeneration. <i>Genome Biology and Evolution</i> , 2021, 13, .	1.1	1
2	<i>Cepaea</i> spp. as a source of <i>Brachylaima mesostoma</i> (Digenea: Brachylaimidae) and <i>Brachylecithum</i> sp. (Digenea: Dicrocoeliidae) larvae in Poland. <i>Parasitology Research</i> , 2020, 119, 145-152.	0.6	7
3	Characteristics of capercaillie (<i>Tetrao urogallus</i>) semen analysed with flow cytometry combined with fertility results. <i>Reproduction in Domestic Animals</i> , 2020, 55, 984-991.	0.6	1
4	Female-Male and Female-Female Social Interactions of Captive Kept Capercaillie (<i>Tetrao Urogallus</i>) and Its Consequences in Planning Breeding Programs. <i>Animals</i> , 2020, 10, 583.	1.0	4
5	Genetic diversity and relationship between cultivated, weedy and wild rye species as revealed by chloroplast and mitochondrial DNA non-coding regions analysis. <i>PLoS ONE</i> , 2019, 14, e0213023.	1.1	21
6	Agents of swimmerâ€™s itchâ€™ dangerous minority in the Digenea invasion of Lymnaeidae in water bodies and the first report of <i>Trichobilharzia regenti</i> in Poland. <i>Parasitology Research</i> , 2018, 117, 3695-3704.	0.6	13
7	Salt mine microorganisms used for the biotransformation of chlorolactones. <i>PLoS ONE</i> , 2018, 13, e0197384.	1.1	3
8	Complete mitochondrial genome of Blue-and-yellow Macaw (<i>Ara ararauna</i>): the species morphologically similar to Blue-throated Macaw (<i>Ara glaucogularis</i>). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 307-308.	0.7	1
9	Complete mitochondrial genome of the Eastern slow worm, <i>Anguis colchica</i> (Nordmann, 1840). <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 67-68.	0.2	0
10	Complete mitochondrial genome of the Italian slow-worm <i>Anguis veronensis</i> Pollini, 1818, and its comparison with mitogenomes of other <i>Anguis</i> species. <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 71-72.	0.2	0
11	Complete mitochondrial genome of golden conure (<i>Guaruba guarouba</i>). <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 33-34.	0.2	1
12	Phylogeography and postglacial colonization of Central Europe by <i>Anguis fragilis</i> and <i>Anguis colchica</i> . <i>Amphibia - Reptilia</i> , 2017, 38, 562-569.	0.1	11
13	Ongoing Speciation and Gene Flow between Taxonomically Challenging <i>Trochulus</i> Species Complex (Gastropoda: Hygromiidae). <i>PLoS ONE</i> , 2017, 12, e0170460.	1.1	8
14	Complete mitochondrial genome of the endemic legless lizard <i>Anguis cephallonica</i> Werner, 1894 and its comparison with mitogenome of <i>Anguis fragilis</i> Linnaeus, 1758. <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 83-85.	0.2	0
15	The first complete genome of â€™ <i>Aratinga</i> genus in comparison to mitogenomes of other parrots from <i>Arini</i> tribe. <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 853-855.	0.2	1
16	Low genetic variability of the edible dormouse (<i>Glis glis</i>) in Stolowe Mountains National Park (Poland)â€™ preliminary results. <i>Mammal Research</i> , 2016, 61, 409-415.	0.6	7
17	The first complete mitochondrial genome of <i>Pyrrhura</i> sp. â€“ question about conspecificity in the light of hybridization between <i>Pyrrhura molinae</i> and <i>Pyrrhura rupicola</i> species. <i>Mitochondrial DNA</i> , 2016, 27, 471-473.	0.6	4
18	Complete mitochondrial genome of Red-throated Conure (<i>Psittacara rubritorquis</i>): its comparison with mitogenome of Socorro Conure (<i>Psittacara brevipes</i>). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016, 27, 3354-3355.	0.7	1

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19	Complete mitochondrial genome of Mitred Conure (<i>Psittacara mitratus</i>): its comparison with mitogenome of Socorro Conure (<i>Psittacara brevipes</i>). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016, 27, 3363-3364.	0.7	3
20	Cytogenetic Examination of South American Tapirs, <i>Tapirus Terrestris</i> (<i>Perissodactyla</i> , <i>Tapiridae</i>), from the Wrocław Zoological Garden. <i>Vestnik Zoologii</i> , 2015, 49, 529-536.	0.7	1
21	Gene Polymorphisms of Novel Immunotolerant Molecule BTLA: Distribution of Alleles, Genotypes and Haplotypes in Polish Caucasian Population. <i>Archivum Immunologiae Et Therapiae Experimentalis</i> , 2015, 63, 73-78.	1.0	11
22	Reintroduction of the European Capercaillie from the Capercaillie Breeding Centre in WisÅa Forest District: Genetic Assessments of Captive and Reintroduced Populations. <i>PLoS ONE</i> , 2015, 10, e0145433.	1.1	10
23	Phylogeny of <i>Amazona barbadensis</i> and the Yellow-Headed Amazon Complex (Aves: <i>Psittacidae</i>): A New Look at South American Parrot Evolution. <i>PLoS ONE</i> , 2014, 9, e97228.	1.1	6
24	Complete mitochondrial genome of endangered Maroon-fronted Parrot (<i>Rhynchopsitta terrisi</i>) – conspecific relation of the species with Thick-billed Parrot (<i>Rhynchopsitta pachyrhyncha</i>). <i>Mitochondrial DNA</i> , 2014, 25, 424-426.	0.6	7
25	Morphological similarity and molecular divergence of <i>Trochulus striolatus</i> and <i>T. montanus</i> , and their relationship to sympatric congeners (Gastropoda: Pulmonata: Hygromiidae). <i>Systematics and Biodiversity</i> , 2014, 12, 366-384.	0.5	11
26	Complete mitochondrial genome of endangered Socorro Conure (<i>Aratinga brevipes</i>) – taxonomic position of the species and its relationship with Green Conure. <i>Mitochondrial DNA</i> , 2014, 25, 365-367.	0.6	10
27	Complete mitochondrial genome of Blue-crowned Parakeet (<i>Aratinga acuticaudata</i>) – phylogenetic position of the species among parrots group called Conures. <i>Mitochondrial DNA</i> , 2013, 24, 336-338.	0.6	9
28	Genetic structuring of the common shrew, <i>Sorex araneus</i> (Soricomorpha: Soricidae) in the Polish Sudetes may suggest ways of northwards colonization. <i>Hereditas</i> , 2012, 149, 197-206.	0.5	4
29	Do aquatic barriers reduce male-mediated gene flow in a hybrid zone of the common shrew (<i>Sorex</i>)? <i>Journal of Biogeography</i> , 2011, 38, 117-122.	0.5	5
30	A microsatellite study in the Gucki/Młocznik/Popielno hybrid zone reveals no genetic differentiation between two chromosome races of the common shrew (<i>Sorex araneus</i>). <i>Acta Theriologica</i> , 2011, 56, 117-122.	1.1	5
31	Variation of the Common Shrew (<i>Sorex Araneus</i> L.) Dentition. <i>Zoologica Poloniae: the Journal of Polish Zoological Society</i> , 2008, 53, 49-56.	0.2	3