

Geetha Annavi

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

199
citations

1478280

6
h-index

1372474

10
g-index

10
all docs

10
docs citations

10
times ranked

322
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic diversity and phylogenetic relationships of Malayan tapir (<i>Tapirus indicus</i>) populations in the Malay Peninsula based on mitochondrial DNA control region. <i>Biodiversity and Conservation</i> , 2021, 30, 2433-2449.	1.2	1
2	Social and Reproductive Behavior of Captive Malayan Tapirs™ (<i>Tapirus indicus</i>): Interactions with Maternal Experience and Environmental Conditions. <i>Scientific Reports</i> , 2020, 10, 4117.	1.6	2
3	Molecular sexing and preliminary assessment of population sex ratio of the endangered Malayan tapir (<i>Tapirus indicus</i>) in Peninsular Malaysia. <i>Scientific Reports</i> , 2020, 10, 3973.	1.6	1
4	Influence of Enclosure Conditions and Visitors on the Behavior of Captive Malayan Tapir (<i>Tapirus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 0 and Research Publications, 2018, 8, .	0.0	2
5	No Compensatory Relationship between the Innate and Adaptive Immune System in Wild-Living European Badgers. <i>PLoS ONE</i> , 2016, 11, e0163773.	1.1	8
6	MHC class II assortative mate choice in European badgers (<i>Meles meles</i>). <i>Molecular Ecology</i> , 2015, 24, 3138-3150.	2.0	40
7	Heterozygosity–fitness correlations in a wild mammal population: accounting for parental and environmental effects. <i>Ecology and Evolution</i> , 2014, 4, 2594-2609.	0.8	33
8	Pathogen burden, coinfection and major histocompatibility complex variability in the European badger (<i>Meles meles</i>). <i>Molecular Ecology</i> , 2014, 23, 5072-5088.	2.0	59
9	Neighbouring group composition and within group relatedness drive extra group paternity rate in the European badger (<i>Meles meles</i>). <i>Journal of Evolutionary Biology</i> , 2014, 27, 2191-2203.	0.8	43
10	Characterisation of twenty-one European badger (<i>Meles meles</i>) microsatellite loci facilitates the discrimination of second-order relatives. <i>Conservation Genetics Resources</i> , 2011, 3, 515-518.	0.4	10