

Elmira Mohandesan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11228254/publications.pdf>

Version: 2024-02-01

12
papers

564
citations

1040056

9
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

719
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic signatures of domestication in Old World camels. <i>Communications Biology</i> , 2020, 3, 316.	4.4	32
2	Combined hybridization capture and shotgun sequencing for ancient <sc>DNA</sc> analysis of extinct wild and domestic dromedary camel. <i>Molecular Ecology Resources</i> , 2017, 17, 300-313.	4.8	25
3	Mitogenome Sequencing in the Genus <i>Camelus</i> Reveals Evidence for Purifying Selection and Long-term Divergence between Wild and Domestic Bactrian Camels. <i>Scientific Reports</i> , 2017, 7, 9970.	3.3	45
4	The major histocompatibility complex in Old World camelids and low polymorphism of its class II genes. <i>BMC Genomics</i> , 2016, 17, 167.	2.8	39
5	Ancient and modern DNA reveal dynamics of domestication and cross-continental dispersal of the dromedary. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6707-6712.	7.1	141
6	The de novo genome assembly and annotation of a female domestic dromedary of North African origin. <i>Molecular Ecology Resources</i> , 2016, 16, 314-324.	4.8	106
7	A Sunken Ship of the Desert at the River Danube in Tulln, Austria. <i>PLoS ONE</i> , 2015, 10, e0121235.	2.5	6
8	Diagnostic single nucleotide polymorphism markers to identify hybridization between dromedary and Bactrian camels. <i>Conservation Genetics Resources</i> , 2015, 7, 329-332.	0.8	5
9	Complete mitochondrial genomes of Tuatara endemic to different islands of New Zealand. <i>Mitochondrial DNA</i> , 2015, 26, 25-26.	0.6	1
10	Molecular and morphological evolution in tuatara are decoupled. <i>Trends in Genetics</i> , 2009, 25, 16-18.	6.7	21
11	Rapid molecular evolution in a living fossil. <i>Trends in Genetics</i> , 2008, 24, 106-109.	6.7	60
12	New developments in ancient genomics. <i>Trends in Ecology and Evolution</i> , 2008, 23, 386-393.	8.7	83