Aleksandr A Khokhlov

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7392679/publications.pdf

Version: 2024-02-01

1163117 1281871 4,488 13 8 citations h-index papers

g-index 14 14 14 4741 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Emergence and intensification of dairying in the Caucasus and Eurasian steppes. Nature Ecology and Evolution, 2022, 6, 813-822.	7.8	22
2	Tracing genetic connections of ancient Hungarians to the 6th–14th century populations of the Volga-Ural region. Human Molecular Genetics, 2022, 31, 3266-3280.	2.9	6
3	Ten millennia of hepatitis B virus evolution. Science, 2021, 374, 182-188.	12.6	64
4	To the Issue of Anthropological Contacts Between the Populations of the Srubnaya and Alakul Cultures of the Late Bronze Age in the Southern Urals and Western Kazakhstan Steppes. Vestnik Volgogradskogo Gosudarstvennogo Universiteta, Seriia 4: Istoriia, Regionovedenie, Mezhdunarodnye Otnosheniia, 2020, , 65-83.	0.1	2
5	The formation of human populations in South and Central Asia. Science, 2019, 365, .	12.6	383
6	Origins of the Physical Appearance of the Sintashta Circle of Cultures People of the Bronze Age: theoretical and practical aspects. Povolzhskaya Arkheologiya, 2019, 1, 59-71.	0.1	4
7	The oldest burials in the Orenburg region (concerning two graves of the burial mound near Labazy) Tj ETQq $1\ 1\ 0$	0.784314 r _{ 0.1	gBT /Overlock 0
8	Analysis of 3800 -year-old Yersinia pestis genomes suggests Bronze Age origin for bubonic plague. Nature Communications, 2018 , 9 , 2234 .	12.8	123
9	Extraordinary Burial of the Eneolithic Burial Ground Ekaterinovsky Cape. Povolzhskaya Arkheologiya, 2018, 3, 58-67.	0.1	0
10	Ancestry and demography and descendants of Iron Age nomads of the Eurasian Steppe. Nature Communications, 2017, 8, 14615.	12.8	96
11	Population genomics of Bronze Age Eurasia. Nature, 2015, 522, 167-172.	27.8	1,166
12	Massive migration from the steppe was a source for Indo-European languages in Europe. Nature, 2015, 522, 207-211.	27.8	1,435
13	Genome-wide patterns of selection in 230 ancient Eurasians. Nature, 2015, 528, 499-503.	27.8	1,160