

Marcel Keller

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2235745/publications.pdf>

Version: 2024-02-01

14
papers

480
citations

1040056

9
h-index

1199594

12
g-index

19
all docs

19
docs citations

19
times ranked

613
citing authors

#	ARTICLE	IF	CITATIONS
1	Performance and automation of ancient DNA capture with RNA hyRAD probes. <i>Molecular Ecology Resources</i> , 2022, 22, 891-907.	4.8	11
2	An invasive <i>Haemophilus influenzae</i> serotype b infection in an Anglo-Saxon plague victim. <i>Genome Biology</i> , 2022, 23, 22.	8.8	13
3	Dining in Tuva: Social correlates of diet and mobility in Southern Siberia during the 2nd–4th centuries <i>CE</i> . <i>American Journal of Biological Anthropology</i> , 2022, 178, 124-139.	1.1	4
4	Troubles in Tuva: Patterns of perimortem trauma in a nomadic community from Southern Siberia (second to fourth c. <i>CE</i>). <i>American Journal of Physical Anthropology</i> , 2021, 174, 3-19.	2.1	11
5	Beyond Plague Pits: Using Genetics to Identify Responses to Plague in Medieval Cambridgeshire. <i>European Journal of Archaeology</i> , 2021, 24, 496-518.	0.5	12
6	Human mobility at Tell Atchana (Alalakh), Hatay, Turkey during the 2nd millennium BC: Integration of isotopic and genomic evidence. <i>PLoS ONE</i> , 2021, 16, e0241883.	2.5	7
7	Ten millennia of hepatitis B virus evolution. <i>Science</i> , 2021, 374, 182-188.	12.6	64
8	Die Justinianische Pest. <i>Evangelische Theologie</i> , 2021, 81, 385-401.	0.0	0
9	A58–Epidemic dynamics of ancient disease outbreaks. <i>Virus Evolution</i> , 2019, 5, .	4.9	0
10	Phylogeography of the second plague pandemic revealed through analysis of historical <i>Yersinia pestis</i> genomes. <i>Nature Communications</i> , 2019, 10, 4470.	12.8	113
11	Ancient <i>Yersinia pestis</i> genomes from across Western Europe reveal early diversification during the First Pandemic (541–750). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12363-12372.	7.1	100
12	Digging up the plague: A diachronic comparison of aDNA confirmed plague burials and associated burial customs in Germany. <i>Prahistorische Zeitschrift</i> , 2018, 92, 405-427.	0.4	5
13	A High-Coverage <i>Yersinia pestis</i> Genome from a Sixth-Century Justinianic Plague Victim. <i>Molecular Biology and Evolution</i> , 2016, 33, 2911-2923.	8.9	109
14	United in death—related by blood? Genetic and archeometric analyses of skeletal remains from the neolithic earthwork bruchsal–eue. <i>American Journal of Physical Anthropology</i> , 2015, 157, 458-471.	2.1	17