Martine Robbeets

List of Publications by Year in descending order

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933447 642732 1,206 28 10 23 citations g-index h-index papers 32 32 32 1040 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Demography, trade and state power: a tripartite model of medieval farming/language dispersals in the Ryukyu Islands. Evolutionary Human Sciences, 2022, 4, .	1.7	4
2	Genomic insights into the formation of human populations in East Asia. Nature, 2021, 591, 413-419.	27.8	216
3	Bronze Age Globalisation and Eurasian Impacts on Later JÅmon Social Change. Journal of World Prehistory, 2021, 34, 121-158.	3.6	10
4	Ancient Y-DNA with reconstructed phylogeny provides insights into the demographic history of paternal haplogroup N1a2-F1360. Journal of Genetics and Genomics, 2021, 48, 1130-1133.	3.9	5
5	The ARCHIPELAGO Archaeological Isotope Database for the Japanese Islands. Journal of Open Archaeology Data, 2021, 9, .	0.8	8
6	Ancient genome analyses shed light on kinship organization and mating practice of Late Neolithic society in China. IScience, 2021, 24, 103352.	4.1	10
7	The genomic origins of the Bronze Age Tarim Basin mummies. Nature, 2021, 599, 256-261.	27.8	65
8	Triangulation supports agricultural spread of the Transeurasian languages. Nature, 2021, 599, 616-621.	27.8	58
9	Bayesian phylolinguistics infers the internal structure and the time-depth of the Turkic language family. Journal of Language Evolution, 2020, 5, 39-53.	2.2	10
10	Archaeolinguistic evidence for the farming/language dispersal of Koreanic. Evolutionary Human Sciences, 2020, 2, .	1.7	7
11	About millets and beans, words and genes. Evolutionary Human Sciences, 2020, 2, .	1.7	2
12	Bioarchaeological perspective on the expansion of Transeurasian languages in Neolithic Amur River basin. Evolutionary Human Sciences, 2020, 2, .	1.7	6
13	Ancient genomes from northern China suggest links between subsistence changes and human migration. Nature Communications, 2020, 11, 2700.	12.8	133
14	Millet agriculture dispersed from Northeast China to the Russian Far East: Integrating archaeology, genetics, and linguistics. Archaeological Research in Asia, 2020, 22, 100177.	0.7	21
15	Tracing population movements in ancient East Asia through the linguistics and archaeology of textile production. Evolutionary Human Sciences, 2020, 2, .	1.7	8
16	The homeland of Proto-Tungusic inferred from contemporary words and ancient genomes. Evolutionary Human Sciences, 2020, 2, .	1.7	7
17	The homelands of the individual Transeurasian proto-languages. , 2020, , 753-771.		69
18	The Transeurasian homeland: where, what, and when?., 2020, , 772-783.		68

#	Article	IF	CITATIONS
19	Ancient Genomes Reveal Yamnaya-Related Ancestry and a Potential Source of Indo-European Speakers in Iron Age Tianshan. Current Biology, 2019, 29, 2526-2532.e4.	3.9	64
20	The genetic history of admixture across inner Eurasia. Nature Ecology and Evolution, 2019, 3, 966-976.	7.8	135
21	Bronze Age population dynamics and the rise of dairy pastoralism on the eastern Eurasian steppe. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E11248-E11255.	7.1	135
22	Bayesian phylolinguistics reveals the internal structure of the Transeurasian family. Journal of Language Evolution, 2018, 3, 145-162.	1.7	18
23	The development of finiteness in the Transeurasian languages. Linguistics, 2017, 55, 489-523.	1.0	4
24	Austronesian influence and Transeurasian ancestry in Japanese. Language Dynamics and Change, 2017, 7, 210-251.	0.6	18
25	Chapter 9. Insubordination and the establishment of genealogical relationship across Eurasia. Typological Studies in Language, 2016, , 209-246.	1.2	2
26	A Bayesian approach to the classification of Tungusic languages. Diachronica, 0, , .	0.5	2
27	ChapterÂ5. The language of the Transeurasian farmers. , 0, , 93-121.		73
28	Chapter 1. Farming/Language Dispersal. , 0, , 1-23.		0