

Alissa Mittnik

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5728309/publications.pdf>

Version: 2024-02-01

22
papers

6,922
citations

361413

20
h-index

677142

22
g-index

24
all docs

24
docs citations

24
times ranked

6102
citing authors

#	ARTICLE	IF	CITATIONS
1	Stone Age <i>Yersinia pestis</i> genomes shed light on the early evolution, diversity, and ecology of plague. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2116722119.	7.1	31
2	Pedigree-based Bayesian modelling of radiocarbon dates. <i>PLoS ONE</i> , 2022, 17, e0270374.	2.5	2
3	Ten millennia of hepatitis B virus evolution. <i>Science</i> , 2021, 374, 182-188.	12.6	64
4	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. <i>Nature Ecology and Evolution</i> , 2020, 4, 334-345.	7.8	95
5	Ancient DNA sheds light on the genetic origins of early Iron Age Philistines. <i>Science Advances</i> , 2019, 5, eaax0061.	10.3	64
6	Kinship-based social inequality in Bronze Age Europe. <i>Science</i> , 2019, 366, 731-734.	12.6	175
7	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018, 555, 190-196.	27.8	503
8	The genomic history of southeastern Europe. <i>Nature</i> , 2018, 555, 197-203.	27.8	479
9	Ancient genomes revisit the ancestry of domestic and Przewalski's horses. <i>Science</i> , 2018, 360, 111-114.	12.6	241
10	The genetic prehistory of the Baltic Sea region. <i>Nature Communications</i> , 2018, 9, 442.	12.8	151
11	Inferring genetic origins and phenotypic traits of George Bähr, the architect of the Dresden Frauenkirche. <i>Scientific Reports</i> , 2018, 8, 2115.	3.3	11
12	Reconciling material cultures in archaeology with genetic data: The nomenclature of clusters emerging from archaeogenomic analysis. <i>Scientific Reports</i> , 2018, 8, 13003.	3.3	69
13	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 2017, 171, 59-71.e21.	28.9	308
14	Female exogamy and gene pool diversification at the transition from the Final Neolithic to the Early Bronze Age in central Europe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 10083-10088.	7.1	136
15	Genetic origins of the Minoans and Mycenaeans. <i>Nature</i> , 2017, 548, 214-218.	27.8	203
16	The Stone Age Plague and Its Persistence in Eurasia. <i>Current Biology</i> , 2017, 27, 3683-3691.e8.	3.9	125
17	The genetic history of Ice Age Europe. <i>Nature</i> , 2016, 534, 200-205.	27.8	729
18	Pleistocene Mitochondrial Genomes Suggest a Single Major Dispersal of Non-Africans and a Late Glacial Population Turnover in Europe. <i>Current Biology</i> , 2016, 26, 827-833.	3.9	277

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19	A Molecular Approach to the Sexing of the Triple Burial at the Upper Paleolithic Site of DolnĀ-VĀstovice. PLoS ONE, 2016, 11, e0163019.	2.5	92
20	Massive migration from the steppe was a source for Indo-European languages in Europe. Nature, 2015, 522, 207-211.	27.8	1,435
21	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
22	A Revised Timescale for Human Evolution Based on Ancient Mitochondrial Genomes. Current Biology, 2013, 23, 553-559.	3.9	540