## 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

times ranked

24

6102 citing authors

#	Article	IF	CITATIONS
1	Massive migration from the steppe was a source for Indo-European languages in Europe. Nature, 2015, 522, 207-211.	27.8	1,435
2	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
3	The genetic history of Ice Age Europe. Nature, 2016, 534, 200-205.	27.8	729
4	A Revised Timescale for Human Evolution Based on Ancient Mitochondrial Genomes. Current Biology, 2013, 23, 553-559.	3.9	540
5	The Beaker phenomenon and the genomic transformation of northwest Europe. Nature, 2018, 555, 190-196.	27.8	503
6	The genomic history of southeastern Europe. Nature, 2018, 555, 197-203.	27.8	479
7	Reconstructing Prehistoric African Population Structure. Cell, 2017, 171, 59-71.e21.	28.9	308
8	Pleistocene Mitochondrial Genomes Suggest a Single Major Dispersal of Non-Africans and a Late Glacial Population Turnover in Europe. Current Biology, 2016, 26, 827-833.	3.9	277
9	Ancient genomes revisit the ancestry of domestic and Przewalski's horses. Science, 2018, 360, 111-114.	12.6	241
10	Genetic origins of the Minoans and Mycenaeans. Nature, 2017, 548, 214-218.	27.8	203
11	Kinship-based social inequality in Bronze Age Europe. Science, 2019, 366, 731-734.	12.6	175
12	The genetic prehistory of the Baltic Sea region. Nature Communications, 2018, 9, 442.	12.8	151
13	Female exogamy and gene pool diversification at the transition from the Final Neolithic to the Early Bronze Age in central Europe. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10083-10088.	7.1	136
14	The Stone Age Plague and Its Persistence in Eurasia. Current Biology, 2017, 27, 3683-3691.e8.	3.9	125
15	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. Nature Ecology and Evolution, 2020, 4, 334-345.	7.8	95
16	A Molecular Approach to the Sexing of the Triple Burial at the Upper Paleolithic Site of DolnÃ-VÄ>stonice. PLoS ONE, 2016, 11, e0163019.	2.5	92
17	Reconciling material cultures in archaeology with genetic data: The nomenclature of clusters emerging from archaeogenomic analysis. Scientific Reports, 2018, 8, 13003.	3.3	69
18	Ancient DNA sheds light on the genetic origins of early Iron Age Philistines. Science Advances, 2019, 5, eaax0061.	10.3	64

## ALISSA MITTNIK

#	Article	IF	CITATION
19	Ten millennia of hepatitis B virus evolution. Science, 2021, 374, 182-188.	12.6	64
20	Stone Age <i>Yersinia pestis</i> genomes shed light on the early evolution, diversity, and ecology of plague. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2116722119.	7.1	31
21	Inferring genetic origins and phenotypic traits of George Bär, the architect of the Dresden Frauenkirche. Scientific Reports, 2018, 8, 2115.	3.3	11
22	Pedigree-based Bayesian modelling of radiocarbon dates. PLoS ONE, 2022, 17, e0270374.	2.5	2