Ben Fabian Krause-Kyora

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2124295/publications.pdf

Version: 2024-02-01

41 papers 1,890 citations

471509 17 h-index 330143 37 g-index

50 all docs

50 docs citations

50 times ranked

3576 citing authors

#	Article	IF	CITATIONS
1	Palaeogenomic analysis of black rat (Rattus rattus) reveals multiple European introductions associated with human economic history. Nature Communications, 2022, 13, 2399.	12.8	12
2	Phylogeography in an "oyster―shell provides first insights into the genetic structure of an extinct Ostrea edulis population. Scientific Reports, 2021, 11, 2307.	3.3	3
3	Genome-wide study of a Neolithic Wartberg grave community reveals distinct HLA variation and hunter-gatherer ancestry. Communications Biology, 2021, 4, 113.	4.4	20
4	Analysis of Genomic DNA from Medieval Plague Victims Suggests Long-Term Effect of <i>Yersinia pestis</i> on Human Immunity Genes. Molecular Biology and Evolution, 2021, 38, 4059-4076.	8.9	29
5	Mass burial genomics reveals outbreak of enteric paratyphoid fever in the Late Medieval trade city Lýbeck. IScience, 2021, 24, 102419.	4.1	9
6	A 5,000-year-old hunter-gatherer already plagued by Yersinia pestis. Cell Reports, 2021, 35, 109278.	6.4	42
7	Ancient DNA Study in Medieval Europeans Shows an Association Between HLA-DRB1*03 and Paratyphoid Fever. Frontiers in Immunology, 2021, 12, 691475.	4.8	3
8	Toward an Investigation of Diversity and Cultivation of Rye (Secale cereale ssp. cereale L.) in Germany: Methodological Insights and First Results from Early Modern Plant Material. Agronomy, 2021, 11, 2451.	3.0	6
9	Mitochondrial DNA of domesticated sheep confirms pastoralist component of Afanasievo subsistence economy in the Altai Mountains (3300–2900ÂcalÂBC). Archaeological Research in Asia, 2020, 24, 100232.	0.7	18
10	HIGH-PRECISION BAYESIAN CHRONOLOGICAL MODELING ON A CALIBRATION PLATEAU: THE NIEDERTIEFENBACH GALLERY GRAVE. Radiocarbon, 2020, 62, 1261-1284.	1.8	14
11	Yersinia pestis strains from Latvia show depletion of the pla virulence gene at the end of the second plague pandemic. Scientific Reports, 2020, 10, 14628.	3.3	25
12	Comparison of target enrichment strategies for ancient pathogen DNA. BioTechniques, 2020, 69, 455-459.	1.8	17
13	Targeted analysis of polymorphic loci from low-coverage shotgun sequence data allows accurate genotyping of HLA genes in historical human populations. Scientific Reports, 2020, 10, 7339.	3.3	6
14	Gene-flow from steppe individuals into Cucuteni-Trypillia associated populations indicates long-standing contacts and gradual admixture. Scientific Reports, 2020, 10, 4253.	3.3	15
15	Two burials in a unique freshwater shell midden: insights into transformations of Stone Age hunter-fisher daily life in Latvia. Archaeological and Anthropological Sciences, 2020, 12, 1.	1.8	13
16	High mitochondrial diversity of domesticated goats persisted among Bronze and Iron Age pastoralists in the Inner Asian Mountain Corridor. PLoS ONE, 2020, 15, e0233333.	2.5	19
17	Title is missing!. , 2020, 15, e0233333.		O
18	Title is missing!. , 2020, 15, e0233333.		0

#	Article	IF	CITATIONS
19	Title is missing!. , 2020, 15, e0233333.		O
20	Title is missing!. , 2020, 15, e0233333.		0
21	Infectious diseases and Neolithic transformations: Evaluating biological and archaeological proxies in the German loess zone between 5500 and 2500 BCE. Holocene, 2019, 29, 1545-1557.	1.7	19
22	Coming and going – Historical distributions of the European oyster Ostrea edulisÂLinnaeus, 1758 and the introduced slipper limpet Crepidula fornicataÂLinnaeus, 1758 in the North Sea. PLoS ONE, 2019, 14, e0224249.	2.5	11
23	Radiocarbon dating and isotope analysis on the purported Aurignacian skeletal remains from Fontana Nuova (Ragusa, Italy). PLoS ONE, 2019, 14, e0213173.	2.5	16
24	The genomic history of southeastern Europe. Nature, 2018, 555, 197-203.	27.8	479
25	Ancient DNA study reveals HLA susceptibility locus for leprosy in medieval Europeans. Nature Communications, 2018, 9, 1569.	12.8	67
26	Neolithic and medieval virus genomes reveal complex evolution of hepatitis B. ELife, 2018, 7, .	6.0	101
27	Draft Genome Sequence of Riemerella anatipestifer Isolate 17CS0503. Genome Announcements, 2018, 6, .	0.8	1
28	The Iceman's Last Meal Consisted of Fat, Wild Meat, and Cereals. Current Biology, 2018, 28, 2348-2355.e9.	3.9	39
29	Ancient genomes reveal a high diversity of Mycobacterium leprae in medieval Europe. PLoS Pathogens, 2018, 14, e1006997.	4.7	98
30	Insights into early pig domestication provided by ancient DNA analysis. Scientific Reports, 2017, 7, 44550.	3.3	19
31	Identification and characterization of two functional variants in the human longevity gene FOXO3. Nature Communications, 2017, 8, 2063.	12.8	69
32	Niedertiefenbach. Ein Galeriegrab der spÄtneolithischen Wartberggruppe sĽdwestlich von Niedertiefenbach (Landkreis Limburg-Weilburg, Hessen). Prahistorische Zeitschrift, 2016, 91, .	0.4	4
33	The 5300-year-old <i>Helicobacter pylori</i> genome of the Iceman. Science, 2016, 351, 162-165.	12.6	200
34	Exploring the complexity of domestication: a response to Rowley-Conwy and Zeder. World Archaeology, 2014, 46, 825-834.	1.1	15
35	Collective burials among agro-pastoral societies in later Neolithic Germany: perspectives from ancient DNA. Journal of Archaeological Science, 2014, 51, 174-180.	2.4	22
36	Ancient DNA insights from the Middle Neolithic in Germany. Archaeological and Anthropological Sciences, 2013, 6, 199.	1.8	5

#	Article	IF	CITATIONS
37	Genome-Wide Comparison of Medieval and Modern <i>Mycobacterium leprae</i> . Science, 2013, 341, 179-183.	12.6	313
38	Use of domesticated pigs by Mesolithic hunter-gatherers in northwestern Europe. Nature Communications, 2013, 4, 2348.	12.8	93
39	STR-typing of ancient skeletal remains: which multiplex-PCR kit is the best?. Croatian Medical Journal, 2012, 53, 416-422.	0.7	13
40	Emerging genetic patterns of the european neolithic: Perspectives from a late neolithic bell beaker burial site in Germany. American Journal of Physical Anthropology, 2012, 148, 571-579.	2.1	47
41	Sequencing of mitochondrial DNA and the problem of human specificity. Forensic Science International: Genetics Supplement Series, 2009, 2, 95-96.	0.3	0