Choongwon Jeong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8640712/publications.pdf

Version: 2024-02-01

44 papers

3,760 citations

218381 26 h-index 38 g-index

53 all docs 53 docs citations

53 times ranked

5635 citing authors

#	Article	IF	CITATIONS
1	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206.	13.7	1,216
2	Admixture facilitates genetic adaptations to high altitude in Tibet. Nature Communications, 2014, 5, 3281.	5.8	172
3	Long-term genetic stability and a high-altitude East Asian origin for the peoples of the high valleys of the Himalayan arc. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 7485-7490.	3.3	151
4	Pleistocene North African genomes link Near Eastern and sub-Saharan African human populations. Science, 2018, 360, 548-552.	6.0	142
5	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.	3.3	135
6	The genetic history of admixture across inner Eurasia. Nature Ecology and Evolution, 2019, 3, 966-976.	3.4	135
7	Ancient genomes from northern China suggest links between subsistence changes and human migration. Nature Communications, 2020, 11, 2700.	5.8	133
8	A Dynamic 6,000-Year Genetic History of Eurasia's Eastern Steppe. Cell, 2020, 183, 890-904.e29.	13.5	124
9	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. Nature, 2019, 570, 236-240.	13.7	118
10	Ancient human genome-wide data from a 3000-year interval in the Caucasus corresponds with eco-geographic regions. Nature Communications, 2019, 10, 590.	5.8	113
11	Ancient genome-wide DNA from France highlights the complexity of interactions between Mesolithic hunter-gatherers and Neolithic farmers. Science Advances, 2020, 6, eaaz5344.	4.7	92
12	Language continuity despite population replacement in Remote Oceania. Nature Ecology and Evolution, 2018, 2, 731-740.	3.4	91
13	Survival of Late Pleistocene Hunter-Gatherer Ancestry in the Iberian Peninsula. Current Biology, 2019, 29, 1169-1177.e7.	1.8	90
14	Ancient Fennoscandian genomes reveal origin and spread of Siberian ancestry in Europe. Nature Communications, 2018, 9, 5018.	5.8	86
15	Genomic History of Neolithic to Bronze Age Anatolia, Northern Levant, and Southern Caucasus. Cell, 2020, 181, 1158-1175.e28.	13.5	86
16	Late Pleistocene human genome suggests a local origin for the first farmers of central Anatolia. Nature Communications, 2019, 10, 1218.	5.8	74
17	Paleolithic to Bronze Age Siberians Reveal Connections with First Americans and across Eurasia. Cell, 2020, 181, 1232-1245.e20.	13.5	71
18	Adaptations to local environments in modern human populations. Current Opinion in Genetics and Development, 2014, 29, 1-8.	1.5	70

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19	The genomic origins of the Bronze Age Tarim Basin mummies. Nature, 2021, 599, 256-261.	13.7	65
20	Ancient DNA sheds light on the genetic origins of early Iron Age Philistines. Science Advances, 2019, 5, eaax0061.	4.7	64
21	The mosaic genome of indigenous African cattle as a unique genetic resource for African pastoralism. Nature Genetics, 2020, 52, 1099-1110.	9.4	61
22	Evidence for early dispersal of domestic sheep into Central Asia. Nature Human Behaviour, 2021, 5, 1169-1179.	6.2	50
23	Detecting past and ongoing natural selection among ethnically Tibetan women at high altitude in Nepal. PLoS Genetics, 2018, 14, e1007650.	1.5	43
24	Ancient genomic time transect from the Central Asian Steppe unravels the history of the Scythians. Science Advances, $2021, 7, .$	4.7	39
25	Genome of a middle Holocene hunter-gatherer from Wallacea. Nature, 2021, 596, 543-547.	13.7	35
26	CoproID predicts the source of coprolites and paleofeces using microbiome composition and host DNA content. PeerJ, 2020, 8, e9001.	0.9	32
27	Deep History of East Asian Populations Revealed Through Genetic Analysis of the Ainu. Genetics, 2016, 202, 261-272.	1.2	28
28	Ethnically Tibetan women in Nepal with low hemoglobin concentration have better reproductive outcomes. Evolution, Medicine and Public Health, 2017, 2017, 82-96.	1.1	28
29	Ancient genomes reveal origin and rapid trans-Eurasian migration of 7th century Avar elites. Cell, 2022, 185, 1402-1413.e21.	13.5	26
30	Ancient genomes from the Himalayas illuminate the genetic history of Tibetans and their Tibeto-Burman speaking neighbors. Nature Communications, 2022, 13, 1203.	5.8	25
31	Genetic structure in the Sherpa and neighboring Nepalese populations. BMC Genomics, 2017, 18, 102.	1.2	21
32	Mapping Variation in Cellular and Transcriptional Response to 1,25-Dihydroxyvitamin D3 in Peripheral Blood Mononuclear Cells. PLoS ONE, 2016, 11, e0159779.	1.1	18
33	The genomic landscape of Nepalese Tibeto-Burmans reveals new insights into the recent peopling of Southern Himalayas. Scientific Reports, 2017, 7, 15512.	1.6	15
34	A longitudinal cline characterizes the genetic structure of human populations in the Tibetan plateau. PLoS ONE, 2017, 12, e0175885.	1.1	15
35	Genomic and dietary discontinuities during the Mesolithic and Neolithic in Sicily. IScience, 2022, 25, 104244.	1.9	11
36	Early nomads of the Eastern Steppe and their tentative connections in the West. Evolutionary Human Sciences, 2020, 2, .	0.9	7

#	Article	IF	CITATIONS
37	Northeastern Asian and Jomon-related genetic structure in the Three Kingdoms period of Gimhae, Korea. Current Biology, 2022, 32, 3232-3244.e6.	1.8	6
38	Mitonuclear incompatibility as a hidden driver behind the genome ancestry of African admixed cattle. BMC Biology, 2022, 20, 20.	1.7	3
39	Ancient DNA Study., 2020, , 1-15.		1
40	A Population Genetic Perspective on Korean Prehistory. Korean Studies, 2019, , .	0.2	0
41	Current Trends in Ancient DNA Study. , 2020, , 1-16.		О
42	A Population Genetic Perspective on Korean Prehistory. Korean Studies, 2020, 44, 27-53.	0.2	0
43	Current Trends in Ancient DNA Study. , 2021, , 285-300.		0
44	Ancient DNA Study. , 2021, , 301-315.		0