Choongwon Jeong

List of Publications by Year
in descending order

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44 papers	3,760 citations	218381 26 h-index	315357 38 g-index
53 all docs	53 docs citations	53 times ranked	5635 citing authors

#	Article	IF	Citations
1	Mitonuclear incompatibility as a hidden driver behind the genome ancestry of African admixed cattle. BMC Biology, 2022, 20, 20.	1.7	3
2	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
3	Ancient genomes reveal origin and rapid trans-Eurasian migration of 7th century Avar elites. Cell, 2022, 185, 1402-1413.e21.	13.5	26
4	Genomic and dietary discontinuities during the Mesolithic and Neolithic in Sicily. IScience, 2022, 25, 104244.	1.9	11
5	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
6	Ancient genomic time transect from the Central Asian Steppe unravels the history of the Scythians. Science Advances, 2021, 7, .	4.7	39
7	Evidence for early dispersal of domestic sheep into Central Asia. Nature Human Behaviour, 2021, 5, 1169-1179.	6.2	50
8	Genome of a middle Holocene hunter-gatherer from Wallacea. Nature, 2021, 596, 543-547.	13.7	35
9	The genomic origins of the Bronze Age Tarim Basin mummies. Nature, 2021, 599, 256-261.	13.7	65
10	Current Trends in Ancient DNA Study. , 2021, , 285-300.		0
11	Ancient DNA Study. , 2021, , 301-315.		0
12	The mosaic genome of indigenous African cattle as a unique genetic resource for African pastoralism. Nature Genetics, 2020, 52, 1099-1110.	9.4	61
13	Current Trends in Ancient DNA Study. , 2020, , 1-16.		0
14	Ancient DNA Study. , 2020, , 1-15.		1
15	A Dynamic 6,000-Year Genetic History of Eurasiaâ€™s Eastern Steppe. Cell, 2020, 183, 890-904.e29.	13.5	124
16	Ancient genomes from northern China suggest links between subsistence changes and human migration. Nature Communications, 2020, 11, 2700.	5.8	133
17	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
18	A Population Genetic Perspective on Korean Prehistory. Korean Studies, 2020, 44, 27-53.	0.2	0

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19	Paleolithic to Bronze Age Siberians Reveal Connections with First Americans and across Eurasia. Cell, 2020, 181, 1232-1245.e20.	13.5	71
20	Genomic History of Neolithic to Bronze Age Anatolia, Northern Levant, and Southern Caucasus. Cell, 2020, 181, 1158-1175.e28.	13.5	86
21	Early nomads of the Eastern Steppe and their tentative connections in the West. Evolutionary Human Sciences, 2020, 2, .	0.9	7
22	CoproID predicts the source of coprolites and paleofeces using microbiome composition and host DNA content. PeerJ, 2020, 8, e9001.	0.9	32
23	Ancient DNA sheds light on the genetic origins of early Iron Age Philistines. Science Advances, 2019, 5, eaax0061.	4.7	64
24	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. Nature, 2019, 570, 236-240.	13.7	118
25	The genetic history of admixture across inner Eurasia. Nature Ecology and Evolution, 2019, 3, 966-976.	3.4	135
26	Late Pleistocene human genome suggests a local origin for the first farmers of central Anatolia. Nature Communications, 2019, 10, 1218.	5.8	74
27	Survival of Late Pleistocene Hunter-Gatherer Ancestry in the Iberian Peninsula. Current Biology, 2019, 29, 1169-1177.e7.	1.8	90
28	A Population Genetic Perspective on Korean Prehistory. Korean Studies, 2019, , .	0.2	0
29	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
30	Language continuity despite population replacement in Remote Oceania. Nature Ecology and Evolution, 2018, 2, 731-740.	3.4	91
31	Pleistocene North African genomes link Near Eastern and sub-Saharan African human populations. Science, 2018, 360, 548-552.	6.0	142
32	Ancient Fennoscandian genomes reveal origin and spread of Siberian ancestry in Europe. Nature Communications, 2018, 9, 5018.	5.8	86
33	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
34	Detecting past and ongoing natural selection among ethnically Tibetan women at high altitude in Nepal. PLoS Genetics, 2018, 14, e1007650.	1.5	43
35	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
36	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

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37	Genetic structure in the Sherpa and neighboring Nepalese populations. BMC Genomics, 2017, 18, 102.	1.2	21
38	A longitudinal cline characterizes the genetic structure of human populations in the Tibetan plateau. PLoS ONE, 2017, 12, e0175885.	1.1	15
39	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
40	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206.	13.7	1,216
41	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
42	Deep History of East Asian Populations Revealed Through Genetic Analysis of the Ainu. Genetics, 2016, 202, 261-272.	1.2	28
43	Adaptations to local environments in modern human populations. Current Opinion in Genetics and Development, 2014, 29, 1-8.	1.5	70
44	Admixture facilitates genetic adaptations to high altitude in Tibet. Nature Communications, 2014, 5, 3281.	5.8	172