

Ancient human parallel lineages within North America

Science

360, 1024-1027

DOI: [10.1126/science.aar6851](https://doi.org/10.1126/science.aar6851)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Early human dispersals within the Americas. <i>Science</i> , 2018, 362, .	6.0	230
2	Native American Genomic Diversity through Ancient DNA. <i>Cell</i> , 2018, 175, 1173-1174.	13.5	1
3	Reconstructing the Deep Population History of Central and South America. <i>Cell</i> , 2018, 175, 1185-1197.e22.	13.5	259
4	Interred with Their DNA. <i>Cell</i> , 2018, 175, 299-301.	13.5	0
5	Pre-Clovis projectile points at the Debra L. Friedkin site, Texas—Implications for the Late Pleistocene peopling of the Americas. <i>Science Advances</i> , 2018, 4, eaat4505.	4.7	80
6	Ancient human genomes—keys to understanding our past. <i>Science</i> , 2018, 360, 964-965.	6.0	12
7	The Current Genomic Landscape of Western South America: Andes, Amazonia, and Pacific Coast. <i>Molecular Biology and Evolution</i> , 2019, 36, 2698-2713.	3.5	59
8	The presence and impact of reference bias on population genomic studies of prehistoric human populations. <i>PLoS Genetics</i> , 2019, 15, e1008302.	1.5	137
9	Late Upper Paleolithic occupation at Cooper’s Ferry, Idaho, USA, ~16,000 years ago. <i>Science</i> , 2019, 365, 891-897.	6.0	126
10	Analysis of the human Y-chromosome haplogroup Q characterizes ancient population movements in Eurasia and the Americas. <i>BMC Biology</i> , 2019, 17, 3.	1.7	36
11	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. <i>Nature</i> , 2019, 570, 236-240.	13.7	118
12	Mutation of a bHLH transcription factor allowed almond domestication. <i>Science</i> , 2019, 364, 1095-1098.	6.0	116
13	Confluences: Fluted Points in the Ice-Free Corridor. <i>PaleoAmerica</i> , 2019, 5, 143-156.	0.4	10
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15	Specialized sledge dogs accompanied Inuit dispersal across the North American Arctic. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191929.	1.2	38
16	Evaluating macroscopic sex estimation methods using genetically sexed archaeological material: The medieval skeletal collection from St John’s Divinity School, Cambridge. <i>American Journal of Physical Anthropology</i> , 2019, 168, 340-351.	2.1	35
17	Y Chromosome Sequences Reveal a Short Beringian Standstill, Rapid Expansion, and early Population structure of Native American Founders. <i>Current Biology</i> , 2019, 29, 149-157.e3.	1.8	94
18	Fladmark + 40: What Have We Learned about a Potential Pacific Coast Peopling of the Americas?. <i>American Antiquity</i> , 2020, 85, 1-21.	0.6	52

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19	Discerning Dispersals along the Pacific and Interior Corridors: Contributions of Geometric Morphometrics to the Peopling of the Americas. <i>PaleoAmerica</i> , 2020, 6, 109-130.	0.4	5
20	Population History and Gene Divergence in Native Mexicans Inferred from 76 Human Exomes. <i>Molecular Biology and Evolution</i> , 2020, 37, 994-1006.	3.5	43
21	The Impact of Ancient Genome Studies in Archaeology. <i>Annual Review of Anthropology</i> , 2020, 49, 277-298.	0.4	13
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30	Genomic insights into the early peopling of the Caribbean. <i>Science</i> , 2020, 369, 456-460.	6.0	44
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123	Peopling of the Caribbean. , 2024, , 292-300.		0

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