

# Nathan Nakatsuka

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4833257/publications.pdf>

Version: 2024-02-01

13  
papers

1,287  
citations

840585

11  
h-index

1125617

13  
g-index

17  
all docs

17  
docs citations

17  
times ranked

3002  
citing authors

#	ARTICLE	IF	CITATIONS
1	The formation of human populations in South and Central Asia. <i>Science</i> , 2019, 365, .	6.0	383
2	Reconstructing the Deep Population History of Central and South America. <i>Cell</i> , 2018, 175, 1185-1197.e22.	13.5	259
3	Factors Influencing the Degradation of Archival Formalin-Fixed Paraffin-Embedded Tissue Sections. <i>Journal of Histochemistry and Cytochemistry</i> , 2011, 59, 356-365.	1.3	180
4	The promise of discovering population-specific disease-associated genes in South Asia. <i>Nature Genetics</i> , 2017, 49, 1403-1407.	9.4	129
5	A Paleogenomic Reconstruction of the Deep Population History of the Andes. <i>Cell</i> , 2020, 181, 1131-1145.e21.	13.5	69
6	An Ancient Harappan Genome Lacks Ancestry from Steppe Pastoralists or Iranian Farmers. <i>Cell</i> , 2019, 179, 729-735.e10.	13.5	62
7	Ethics of DNA research on human remains: five globally applicable guidelines. <i>Nature</i> , 2021, 599, 41-46.	13.7	49
8	CLYBL is a polymorphic human enzyme with malate synthase and Î <sup>2</sup> -methylmalate synthase activity. <i>Human Molecular Genetics</i> , 2014, 23, 2313-2323.	1.4	29
9	ContamLD: estimation of ancient nuclear DNA contamination using breakdown of linkage disequilibrium. <i>Genome Biology</i> , 2020, 21, 199.	3.8	29
10	Ancient genomes in South Patagonia reveal population movements associated with technological shifts and geography. <i>Nature Communications</i> , 2020, 11, 3868.	5.8	28
11	Integration of ancient DNA with transdisciplinary dataset finds strong support for Inca resettlement in the south Peruvian coast. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 18359-18368.	3.3	21
12	Two genetic variants explain the association of European ancestry with multiple sclerosis risk in African-Americans. <i>Scientific Reports</i> , 2020, 10, 16902.	1.6	10
13	Ancient genomes reveal long-range influence of the pre-Columbian culture and site of Tiwanaku. <i>Science Advances</i> , 2021, 7, eabg7261.	4.7	8