Rem I Sukernik

List of Publications by Year
in descending order

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26 papers	6,028 citations	361413 20 h-index	552781 26 g-index
27 all docs	27 docs citations	27 times ranked	8594 citing authors

#	Article	IF	Citations
1	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206.	27.8	1,216
2	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
3	Natural selection shaped regional mtDNA variation in humans. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 171-176.	7.1	889
4	Reconstructing Native American population history. Nature, 2012, 488, 370-374.	27.8	699
5	Global diversity, population stratification, and selection of human copy-number variation. Science, 2015, 349, aab3761.	12.6	293
6	Adaptations to Climate-Mediated Selective Pressures in Humans. PLoS Genetics, 2011, 7, e1001375.	3.5	247
7	mtDNA Diversity in Chukchi and Siberian Eskimos: Implications for the Genetic History of Ancient Beringia and the Peopling of the New World. American Journal of Human Genetics, 1998, 63, 1473-1491.	6.2	209
8	The role of mtDNA background in disease expression: a new primary LHON mutation associated with Western Eurasian haplogroupÂJ. Human Genetics, 2002, 110, 130-138.	3.8	195
9	Mitochondrial DNA variation in Koryaks and Itel'men: Population replacement in the Okhotsk Sea-Bering Sea region during the neolithic. American Journal of Physical Anthropology, 1999, 108, 1-39.	2.1	186
10	Mitochondrial DNA Diversity in Indigenous Populations of the Southern Extent of Siberia, and the Origins of Native American Haplogroups. Annals of Human Genetics, 2005, 69, 67-89.	0.8	175
11	The Dual Origin and Siberian Affinities of Native American Y Chromosomes. American Journal of Human Genetics, 2002, 70, 192-206.	6.2	169
12	Mitochondrial Genome Diversity in Arctic Siberians, with Particular Reference to the EvolutionaryÂHistory of Beringia and Pleistocenic Peopling of the Americas. American Journal of Human Genetics, 2008, 82, 1084-1100.	6.2	109
13	Traces of Early Eurasians in the Mansi of Northwest Siberia Revealed by Mitochondrial DNA Analysis. American Journal of Human Genetics, 2002, 70, 1009-1014.	6.2	95
14	Novel mtDNA mutations and oxidative phosphorylation dysfunction in Russian LHON families. Human Genetics, 2001, 109, 33-39.	3.8	90
15	Analysis of Mitochondrial DNA Diversity in the Aleuts of the Commander Islands and Its Implications for the Genetic History of Beringia. American Journal of Human Genetics, 2002, 71, 415-421.	6.2	88
16	Origin and affinities of indigenous Siberian populations as revealed by HLA class II gene frequencies. Human Genetics, 2002, 110, 209-226.	3.8	37
17	Mitochondrial genome diversity in the tubalar, even, and ulchi: Contribution to prehistory of native siberians and their affinities to native americans. American Journal of Physical Anthropology, 2012, 148, 123-138.	2.1	34
18	Growth and nutritional status of the Evenki reindeer herders of Siberia. American Journal of Human Biology, 1994, 6, 339-350.	1.6	31

#	Article	IF	Citations
19	Mitochondrial genome diversity at the Bering Strait area highlights prehistoric human migrations from Siberia to northern North America. European Journal of Human Genetics, 2015, 23, 1399-1404.	2.8	25
20	Correlates of low serum lipid levels among the Evenki herders of Siberia. American Journal of Human Biology, 1994, 6, 329-338.	1.6	22
21	Reindeer Chukchi and Siberian Eskimos: Studies on blood groups, serum proteins, and red cell enzymes with regard to genetic heterogeneity. American Journal of Physical Anthropology, 1981, 55, 121-128.	2.1	14
22	Blood groups, serum proteins, and red cell enzymes in the Nganasans(Tavghi)-reindeer hunters from Taimir Peninsula. American Journal of Physical Anthropology, 1981, 56, 139-145.	2.1	11
23	Reply to Tarazona-Santos and Santos. American Journal of Human Genetics, 2002, 70, 1380-1381.	6.2	4
24	Mitochondrial genome diversity on the Central Siberian Plateau with particular reference to the prehistory of northernmost Eurasia. PLoS ONE, 2021, 16, e0244228.	2.5	4
25	Genetic legacy of cultures indigenous to the Northeast Asian coast in mitochondrial genomes of nearly extinct maritime tribes. BMC Evolutionary Biology, 2020, 20, 83.	3.2	3
26	Mitochondrial DNA variation in Koryaks and Itel'men: Population replacement in the Okhotsk Seaâ€“Bering Sea region during the neolithic. , 0, .		1