List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Mitogenomic diversity in Czechs and Slovaks. Forensic Science International: Genetics, 2022, 59, 102714.	1.6	0
2	Response to Wyckelsma etÂal.: Loss of α-actinin-3 during human evolution provides superior cold resilience and muscle heat generation. American Journal of Human Genetics, 2022, 109, 967-972.	2.6	4
3	Mitogenomics of modern Mongolic-speaking populations. Molecular Genetics and Genomics, 2021, , 1.	1.0	2
4	Complete mitogenome data for the Serbian population: the contribution to high-quality forensic databases. International Journal of Legal Medicine, 2020, 134, 1581-1590.	1.2	7
5	Insights into matrilineal genetic structure, differentiation and ancestry of Armenians based on complete mitogenome data. Molecular Genetics and Genomics, 2019, 294, 1547-1559.	1.0	9
6	Mitogenomic diversity and differentiation of the Buryats. Journal of Human Genetics, 2018, 63, 71-81.	1.1	10
7	Whole mitochondrial genome diversity in two Hungarian populations. Molecular Genetics and Genomics, 2018, 293, 1255-1263.	1.0	19
8	Mitochondrial super-haplogroup U diversity in Serbians. Annals of Human Biology, 2017, 44, 408-418.	0.4	16
9	Mitogenomic diversity in Russians and Poles. Forensic Science International: Genetics, 2017, 30, 51-56.	1.6	17
10	Eight Millennia of Matrilineal Genetic Continuity in the South Caucasus. Current Biology, 2017, 27, 2023-2028.e7.	1.8	37
11	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	13.7	360
12	Selective sweep on human amylase genes postdates the split with Neanderthals. Scientific Reports, 2016, 6, 37198.	1.6	67
13	East Eurasian ancestry in the middle of Europe: genetic footprints of Steppe nomads in the genomes of Belarusian Lipka Tatars. Scientific Reports, 2016, 6, 30197.	1.6	14
14	Y chromosome haplotype diversity in Mongolic-speaking populations and gene conversion at the duplicated STR DYS385a,b in haplogroup C3-M407. Journal of Human Genetics, 2016, 61, 491-496.	1.1	7
15	The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. PLoS Genetics, 2015, 11, e1005068.	1.5	149
16	Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884.	6.0	449
17	Colonization history of the sable <i>Martes zibellina</i> (Mammalia, Carnivora) on the marginal peninsula and islands of northeastern Eurasia. Journal of Mammalogy, 2015, 96, 172-184.	0.6	18
18	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	2.4	348

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19	Mitochondrial DNA perspective of Serbian genetic diversity. American Journal of Physical Anthropology, 2015, 156, 449-465.	2.1	15
20	A novel multiplex assay amplifying 13 Y-STRs characterized by rapid and moderate mutation rate. Forensic Science International: Genetics, 2015, 15, 49-55.	1.6	19
21	Simple and cost-effective 14-loci SNP assay designed for differentiation of European, East Asian and African samples. Forensic Science International: Genetics, 2015, 14, 42-49.	1.6	17
22	Genome-Wide Analysis of Cold Adaptation in Indigenous Siberian Populations. PLoS ONE, 2014, 9, e98076.	1.1	128
23	Western Eurasian ancestry in modern Siberians based on mitogenomic data. BMC Evolutionary Biology, 2014, 14, 217.	3.2	41
24	Phylogenetic relationships among Neoechinorhynchus species (Acanthocephala:) Tj ETQq0 0 0 rgBT /Overlock 10 2014, 63, 100-107.	Tf 50 547 0.6	Td (Neoech 31
25	A Selective Sweep on a Deleterious Mutation in CPT1A in Arctic Populations. American Journal of Human Genetics, 2014, 95, 584-589.	2.6	119
26	A mitogenomic phylogeny and genetic history of sable (Martes zibellina). Gene, 2014, 550, 56-67.	1.0	16
27	Y-chromosome diversity in the Kalmyks at the ethnical and tribal levels. Journal of Human Genetics, 2013, 58, 804-811.	1.1	18
28	Y-chromosome variation in Tajiks and Iranians. Annals of Human Biology, 2013, 40, 48-54.	0.4	6
29	Phylogeny and genetic history of the Siberian salamander (Salamandrella keyserlingii, Dybowski, 1870) inferred from complete mitochondrial genomes. Molecular Phylogenetics and Evolution, 2013, 67, 348-357.	1.2	17
30	Phylogeography of sable (Martes zibellina L. 1758) in the southeast portion of its range based on mitochondrial DNA variation: highlighting the evolutionary history of the sable. Acta Theriologica, 2013, 58, 139-148.	1.1	16
31	The History of Slavs Inferred from Complete Mitochondrial Genome Sequences. PLoS ONE, 2013, 8, e54360.	1.1	62
32	Complete Mitochondrial DNA Diversity in Iranians. PLoS ONE, 2013, 8, e80673.	1.1	93
33	The Y-chromosome C3* Star-Cluster Attributed to Genghis Khan's Descendants is Present at High Frequency in the Kerey Clan from Kazakhstan. Human Biology, 2012, 84, 79-89.	0.4	39
34	Complete Mitochondrial DNA Analysis of Eastern Eurasian Haplogroups Rarely Found in Populations of Northern Asia and Eastern Europe. PLoS ONE, 2012, 7, e32179.	1.1	57
35	On the Y-chromosome haplogroup C3c classification. Journal of Human Genetics, 2012, 57, 685-686.	1.1	5
36	Ancient links between Siberians and Native Americans revealed by subtyping the Y chromosome haplogroup Q1a. Journal of Human Genetics, 2011, 56, 583-588.	1.1	56

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37	Mitochondrial haplogroup N1a phylogeography, with implication to the origin of European farmers. BMC Evolutionary Biology, 2010, 10, 304.	3.2	26
38	Similarities and distinctions in Y chromosome gene pool of Western Slavs. American Journal of Physical Anthropology, 2010, 142, 540-548.	2.1	27
39	Phylogeography and molecular adaptation of Siberian salamander Salamandrella keyserlingii based on mitochondrial DNA variation. Molecular Phylogenetics and Evolution, 2010, 56, 562-571.	1.2	18
40	Phylogeography of the Y hromosome haplogroup C in northern Eurasia. Annals of Human Genetics, 2010, 74, 539-546.	0.3	45
41	The Peopling of Europe from the Mitochondrial Haplogroup U5 Perspective. PLoS ONE, 2010, 5, e10285.	1.1	89
42	Origin and Post-Glacial Dispersal of Mitochondrial DNA Haplogroups C and D in Northern Asia. PLoS ONE, 2010, 5, e15214.	1.1	106
43	Mitogenomic Diversity in Tatars from the Volga-Ural Region of Russia. Molecular Biology and Evolution, 2010, 27, 2220-2226.	3.5	47
44	On the origin of Y-chromosome haplogroup N1b. European Journal of Human Genetics, 2009, 17, 1540-1541.	1.4	3
45	Developing STR databases on structured populations: The native South Siberian population versus the Russian population. Forensic Science International: Genetics, 2009, 3, e111-e116.	1.6	22
46	Reconstructing the phylogeny of African mitochondrial DNA lineages in Slavs. European Journal of Human Genetics, 2008, 16, 1091-1096.	1.4	14
47	Mitochondrial DNA Variability in Slovaks, with Application to the Roma Origin. Annals of Human Genetics, 2008, 72, 228-240.	0.3	43
48	Mitochondrial Haplogroup U2d Phylogeny and Distribution. Human Biology, 2008, 80, 565-571.	0.4	10
49	Mitochondrial DNA Phylogeny in Eastern and Western Slavs. Molecular Biology and Evolution, 2008, 25, 1651-1658.	3.5	84
50	Complex interactions of the Eastern and Western Slavic populations with other European groups as revealed by mitochondrial DNA analysis. Forensic Science International: Genetics, 2007, 1, 141-147.	1.6	60
51	Phylogeographic Analysis of Mitochondrial DNA in Northern Asian Populations. American Journal of Human Genetics, 2007, 81, 1025-1041.	2.6	183
52	The variation of 15 autosomal microsatellite DNA loci in five indigenous populations of South Siberia. Molecular Biology, 2007, 41, 531-538.	0.4	4
53	Distribution of the male lineages of Genghis Khan's descendants in northern Eurasian populations. Russian Journal of Genetics, 2007, 43, 334-337.	0.2	21
54	Y-chromosome haplogroup N dispersals from south Siberia to Europe. Journal of Human Genetics, 2007, 52, 763-770.	1.1	65

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55	Mitochondrial DNA Variability in the Czech Population, with Application to the Ethnic History of Slavs. Human Biology, 2006, 78, 681-695.	0.4	29
56	Mitochondrial DNA Diversity in the Polish Roma. Annals of Human Genetics, 2006, 70, 195-206.	0.3	34
57	Allelic and haplotypic frequencies at 11 Y-STR loci in Buryats from South-East Siberia. Forensic Science International, 2006, 164, 271-275.	1.3	20
58	Contrasting patterns of Y-chromosome variation in South Siberian populations from Baikal and Altai-Sayan regions. Human Genetics, 2006, 118, 591-604.	1.8	70
59	Complete Mitochondrial Genome and Phylogeny of Pleistocene MammothMammuthus primigenius. PLoS Biology, 2006, 4, e73.	2.6	107
60	Differentiation of Mitochondrial DNA and Y Chromosomes in Russian Populations. Human Biology, 2004, 76, 877-900.	0.4	67
61	Disuniting Uniformity: A Pied Cladistic Canvas of mtDNA Haplogroup H in Eurasia. Molecular Biology and Evolution, 2004, 21, 2012-2021.	3.5	170
62	Diversity of Mitochondrial DNA Lineages in South Siberia. Annals of Human Genetics, 2003, 67, 391-411.	0.3	115
63	Mitochondrial DNA Variability in Bosnians and Slovenians. Annals of Human Genetics, 2003, 67, 412-425.	0.3	68
64	Origin and Diffusion of mtDNA Haplogroup X. American Journal of Human Genetics, 2003, 73, 1178-1190.	2.6	148
65	Analysis of phylogenetically reconstructed mutational spectra in human mitochondrial DNA control region. Human Genetics, 2002, 111, 46-53.	1.8	67
66	Title is missing!. Russian Journal of Genetics, 2002, 38, 1098-1103.	0.2	3
67	Title is missing!. Russian Journal of Genetics, 2002, 38, 1196-1202.	0.2	8
68	Mitochondrial DNA Variation in Russian Populations of Stavropol Krai, Orel and Saratov Oblasts. Russian Journal of Genetics, 2002, 38, 1298-1303.	0.2	3
69	Mitochondrial DNA variability in Poles and Russians. Annals of Human Genetics, 2002, 66, 261-83.	0.3	63
70	The Presence of Mitochondrial Haplogroup X in Altaians from South Siberia. American Journal of Human Genetics, 2001, 69, 237-241.	2.6	67
71	Mitochondrial DNA variability in Russians and Ukrainians: Implication to the origin of the Eastern Slavs. Annals of Human Genetics, 2001, 65, 63-78.	0.3	79