Miroslava Derenko

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3281221/publications.pdf

Version: 2024-02-01

71 papers

4,275 citations

33 h-index 63 g-index

72 all docs

 $\begin{array}{c} 72 \\ \text{docs citations} \end{array}$

72 times ranked 4886 citing authors

#	Article	IF	CITATIONS
1	Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884.	6.0	449
2	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	13.7	360
3	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	2.4	348
4	Phylogeographic Analysis of Mitochondrial DNA in Northern Asian Populations. American Journal of Human Genetics, 2007, 81, 1025-1041.	2.6	183
5	Disuniting Uniformity: A Pied Cladistic Canvas of mtDNA Haplogroup H in Eurasia. Molecular Biology and Evolution, 2004, 21, 2012-2021.	3 . 5	170
6	The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. PLoS Genetics, 2015, 11, e1005068.	1.5	149
7	Origin and Diffusion of mtDNA Haplogroup X. American Journal of Human Genetics, 2003, 73, 1178-1190.	2.6	148
8	Genome-Wide Analysis of Cold Adaptation in Indigenous Siberian Populations. PLoS ONE, 2014, 9, e98076.	1.1	128
9	A Selective Sweep on a Deleterious Mutation in CPT1A in Arctic Populations. American Journal of Human Genetics, 2014, 95, 584-589.	2.6	119
10	Diversity of Mitochondrial DNA Lineages in South Siberia. Annals of Human Genetics, 2003, 67, 391-411.	0.3	115
11	Complete Mitochondrial Genome and Phylogeny of Pleistocene MammothMammuthus primigenius. PLoS Biology, 2006, 4, e73.	2.6	107
12	Origin and Post-Glacial Dispersal of Mitochondrial DNA Haplogroups C and D in Northern Asia. PLoS ONE, 2010, 5, e15214.	1.1	106
13	Complete Mitochondrial DNA Diversity in Iranians. PLoS ONE, 2013, 8, e80673.	1.1	93
14	The Peopling of Europe from the Mitochondrial Haplogroup U5 Perspective. PLoS ONE, 2010, 5, e10285.	1.1	89
15	Mitochondrial DNA Phylogeny in Eastern and Western Slavs. Molecular Biology and Evolution, 2008, 25, 1651-1658.	3.5	84
16	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
17	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
18	Mitochondrial DNA Variability in Bosnians and Slovenians. Annals of Human Genetics, 2003, 67, 412-425.	0.3	68

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19	The Presence of Mitochondrial Haplogroup X in Altaians from South Siberia. American Journal of Human Genetics, 2001, 69, 237-241.	2.6	67
20	Analysis of phylogenetically reconstructed mutational spectra in human mitochondrial DNA control region. Human Genetics, 2002, 111, 46-53.	1.8	67
21	Differentiation of Mitochondrial DNA and Y Chromosomes in Russian Populations. Human Biology, 2004, 76, 877-900.	0.4	67
22	Selective sweep on human amylase genes postdates the split with Neanderthals. Scientific Reports, 2016, 6, 37198.	1.6	67
23	Y-chromosome haplogroup N dispersals from south Siberia to Europe. Journal of Human Genetics, 2007, 52, 763-770.	1.1	65
24	Mitochondrial DNA variability in Poles and Russians. Annals of Human Genetics, 2002, 66, 261-83.	0.3	63
25	The History of Slavs Inferred from Complete Mitochondrial Genome Sequences. PLoS ONE, 2013, 8, e54360.	1.1	62
26	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
27	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
28	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
29	Mitogenomic Diversity in Tatars from the Volga-Ural Region of Russia. Molecular Biology and Evolution, 2010, 27, 2220-2226.	3 . 5	47
30	Phylogeography of the Yâ€chromosome haplogroup C in northern Eurasia. Annals of Human Genetics, 2010, 74, 539-546.	0.3	45
31	Mitochondrial DNA Variability in Slovaks, with Application to the Roma Origin. Annals of Human Genetics, 2008, 72, 228-240.	0.3	43
32	Western Eurasian ancestry in modern Siberians based on mitogenomic data. BMC Evolutionary Biology, 2014, 14, 217.	3.2	41
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	Eight Millennia of Matrilineal Genetic Continuity in the South Caucasus. Current Biology, 2017, 27, 2023-2028.e7.	1.8	37
35	Mitochondrial DNA Diversity in the Polish Roma. Annals of Human Genetics, 2006, 70, 195-206.	0.3	34
36	Phylogenetic relationships among Neoechinorhynchus species (Acanthocephala:) Tj ETQq0 0 0 rgBT /Overlock 10	O Tf 50 67 To .6	Td (Neoechin 31

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2014, 63, 100-107.

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37	Mitochondrial DNA Variability in the Czech Population, with Application to the Ethnic History of Slavs. Human Biology, 2006, 78, 681-695.	0.4	29
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	Mitochondrial haplogroup N1a phylogeography, with implication to the origin of European farmers. BMC Evolutionary Biology, 2010, 10, 304.	3.2	26
40	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
41	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
42	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
43	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
44	Whole mitochondrial genome diversity in two Hungarian populations. Molecular Genetics and Genomics, 2018, 293, 1255-1263.	1.0	19
45	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
46	Y-chromosome diversity in the Kalmyks at the ethnical and tribal levels. Journal of Human Genetics, 2013, 58, 804-811.	1.1	18
47	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
48	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
49	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
50	Mitogenomic diversity in Russians and Poles. Forensic Science International: Genetics, 2017, 30, 51-56.	1.6	17
51	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
52	A mitogenomic phylogeny and genetic history of sable (Martes zibellina). Gene, 2014, 550, 56-67.	1.0	16
53	Mitochondrial super-haplogroup U diversity in Serbians. Annals of Human Biology, 2017, 44, 408-418.	0.4	16
54	Mitochondrial DNA perspective of Serbian genetic diversity. American Journal of Physical Anthropology, 2015, 156, 449-465.	2.1	15

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55	Reconstructing the phylogeny of African mitochondrial DNA lineages in Slavs. European Journal of Human Genetics, 2008, 16, 1091-1096.	1.4	14
56	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
57	Mitochondrial Haplogroup U2d Phylogeny and Distribution. Human Biology, 2008, 80, 565-571.	0.4	10
58	Mitogenomic diversity and differentiation of the Buryats. Journal of Human Genetics, 2018, 63, 71-81.	1.1	10
59	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
60	Title is missing!. Russian Journal of Genetics, 2002, 38, 1196-1202.	0.2	8
61	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
62	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
63	Y-chromosome variation in Tajiks and Iranians. Annals of Human Biology, 2013, 40, 48-54.	0.4	6
64	On the Y-chromosome haplogroup C3c classification. Journal of Human Genetics, 2012, 57, 685-686.	1.1	5
65	The variation of 15 autosomal microsatellite DNA loci in five indigenous populations of South Siberia. Molecular Biology, 2007, 41, 531-538.	0.4	4
66	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
67	Title is missing!. Russian Journal of Genetics, 2002, 38, 1098-1103.	0.2	3
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	On the origin of Y-chromosome haplogroup N1b. European Journal of Human Genetics, 2009, 17, 1540-1541.	1.4	3
70	Mitogenomics of modern Mongolic-speaking populations. Molecular Genetics and Genomics, 2021, , 1.	1.0	2
71	Mitogenomic diversity in Czechs and Slovaks. Forensic Science International: Genetics, 2022, 59, 102714.	1.6	O