

# Horolma Pamjav

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3218792/publications.pdf>

Version: 2024-02-01

20  
papers

469  
citations

1039406

9  
h-index

794141

19  
g-index

20  
all docs

20  
docs citations

20  
times ranked

657  
citing authors

#	ARTICLE	IF	CITATIONS
1	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. <i>Forensic Science International: Genetics</i> , 2014, 12, 12-23.	1.6	214
2	A Y-chromosomal comparison of the Madjars (Kazakhstan) and the Magyars (Hungary). <i>American Journal of Physical Anthropology</i> , 2009, 139, 305-310.	2.1	56
3	Genetic structure of the paternal lineage of the Roma People. <i>American Journal of Physical Anthropology</i> , 2011, 145, 21-29.	2.1	30
4	A comparative phylogenetic study of genetics and folk music. <i>Molecular Genetics and Genomics</i> , 2012, 287, 337-349.	1.0	30
5	Testing Central and Inner Asian admixture among contemporary Hungarians. <i>Forensic Science International: Genetics</i> , 2015, 15, 121-126.	1.6	22
6	Brief communication: New Y-chromosome binary markers improve phylogenetic resolution within haplogroup R1a1. <i>American Journal of Physical Anthropology</i> , 2012, 149, 611-615.	2.1	21
7	A study of the Bodrogköz population in north-eastern Hungary by Y chromosomal haplotypes and haplogroups. <i>Molecular Genetics and Genomics</i> , 2017, 292, 883-894.	1.0	21
8	Hungarian population data of four X-linked markers: DXS8378, DXS7132, HPRTB, and DXS7423. <i>International Journal of Legal Medicine</i> , 2006, 121, 74-77.	1.2	19
9	Y-SNP L1034: limited genetic link between Mansi and Hungarian-speaking populations. <i>Molecular Genetics and Genomics</i> , 2015, 290, 377-386.	1.0	15
10	Genetic history of Bashkirian Mari and Southern Mansi ethnic groups in the Ural region. <i>Molecular Genetics and Genomics</i> , 2019, 294, 919-930.	1.0	9
11	Autosomal STR variations reveal genetic heterogeneity in the Mon-Khmer speaking group of Northern Thailand. <i>Forensic Science International: Genetics</i> , 2017, 27, 92-99.	1.6	7
12	New clustering methods for population comparison on paternal lineages. <i>Molecular Genetics and Genomics</i> , 2015, 290, 767-784.	1.0	5
13	Identification of World War II bone remains found in Ukraine using classical anthropological and mitochondrial DNA results. <i>International Journal of Legal Medicine</i> , 2020, 134, 487-489.	1.2	5
14	A simultaneous search for footprints of early human migration processes using the genetic and folk music data in Eurasia. <i>Molecular Genetics and Genomics</i> , 2019, 294, 941-962.	1.0	4
15	mtDNA analysis of 174 Eurasian populations using a new iterative rank correlation method. <i>Molecular Genetics and Genomics</i> , 2016, 291, 493-509.	1.0	3
16	A new self-learning computational method for footprints of early human migration processes. <i>Molecular Genetics and Genomics</i> , 2018, 293, 1579-1594.	1.0	3
17	Paternal genetic history of the Yong population in northern Thailand revealed by Y-chromosomal haplotypes and haplogroups. <i>Molecular Genetics and Genomics</i> , 2020, 295, 579-589.	1.0	3
18	Statistical and population genetics issues of two Hungarian datasets from the aspect of DNA evidence interpretation. <i>Forensic Science International: Genetics</i> , 2015, 19, 18-21.	1.6	1

#	ARTICLE	IF	CITATIONS
19	DNA identification of relics of Blessed Stephen Sándor beatified by Pope Francis 2013. <i>Biologia Futura</i> , 2020, 71, 131-136.	0.6	1
20	A populációgenetika újabb eredménye: genetikai eredetkutatás. <i>Belvárosi Szemle</i> , 2019, 67, .	0.0	0