Elvira Pocheshkhova

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6698262/publications.pdf

Version: 2024-02-01

12 1,269 11
papers citations h-index

13 13 2366
all docs docs citations times ranked citing authors

12

g-index

#	Article	IF	CITATIONS
1	The genetic history of admixture across inner Eurasia. Nature Ecology and Evolution, 2019, 3, 966-976.	7.8	135
2	Genetic differentiation between upland and lowland populations shapes the Y-chromosomal landscape of West Asia. Human Genetics, 2017, 136, 437-450.	3.8	17
3	Update of the GJB2/DFNB1 mutation spectrum in Russia: a founder Ingush mutation del(GJB2-D13S175) is the most frequent among other large deletions. Journal of Human Genetics, 2017, 62, 789-795.	2.3	29
4	Population biobanks: Organizational models and prospects of application in gene geography and personalized medicine. Russian Journal of Genetics, 2016, 52, 1227-1243.	0.6	26
5	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	27.8	360
6	Deep Phylogenetic Analysis of Haplogroup G1 Provides Estimates of SNP and STR Mutation Rates on the Human Y-Chromosome and Reveals Migrations of Iranic Speakers. PLoS ONE, 2015, 10, e0122968.	2.5	35
7	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
8	Parallel Evolution of Genes and Languages in the Caucasus Region. Molecular Biology and Evolution, 2011, 28, 2905-2920.	8.9	149
9	The russian gene pool: the gene geography of Alu insertions (ACE, APOA1, B65, PV92, TPA25). Molecular Biology, 2010, 44, 393-400.	1.3	3
10	Two Sources of the Russian Patrilineal Heritage in Their Eurasian Context. American Journal of Human Genetics, 2008, 82, 236-250.	6.2	122
11	Is Spatial Distribution of the HIV-1-resistant CCR5Δ32 Allele Formed by Ecological Factors?. Journal of Physiological Anthropology and Applied Human Science, 2005, 24, 375-382.	0.4	18
12	Apolipoprotein B 3′-VNTR polymorphism in Eastern European populations. European Journal of Human Genetics, 2003, 11, 444-451.	2.8	23