Levon Yepiskoposyan

List of Publications by Year
in descending order

42 papers	6,636 citations	279798 23 h-index	265206 42 g-index
48 all docs	48 docs citations	48 times ranked	8023 citing authors

#	Article	IF	Citations
1	Prevalence of uncoupling protein one genetic polymorphisms and their relationship with cardiovascular and metabolic health. PLoS ONE, 2022, 17, e0266386.	2.5	2
2	Origin and diffusion of human Y chromosome haplogroup J1-M267. Scientific Reports, 2021, 11, 6659.	3.3	26
3	Insights into matrilineal genetic structure, differentiation and ancestry of Armenians based on complete mitogenome data. Molecular Genetics and Genomics, 2019, 294, 1547-1559.	2.1	9
4	Ancient DNA shows high faunal diversity in the Lesser Caucasus during the Late Pleistocene. Quaternary Science Reviews, 2019, 219, 102-111.	3.0	5
5	The genetic history of admixture across inner Eurasia. Nature Ecology and Evolution, 2019, 3, 966-976.	7.8	135
6	Ancient pathogen <scp>DNA</scp> in human teeth and petrous bones. Ecology and Evolution, 2018, 8, 3534-3542.	1.9	38
7	137 ancient human genomes from across the Eurasian steppes. Nature, 2018, 557, 369-374.	27.8	325
8	Origin and spread of human mitochondrial DNA haplogroup U7. Scientific Reports, 2017, 7, 46044.	3.3	25
9	The genetic variation in the R1a clade among the Ashkenazi Levitesâ€™ Y chromosome. Scientific Reports, 2017, 7, 14969.	3.3	13
10	Eight Millennia of Matrilineal Genetic Continuity in the South Caucasus. Current Biology, 2017, 27, 2023-2028.e7.	3.9	37
11	Demographic cognitive patterns revealed from human genome. , 2017, , .		0
12	The mRNA expression levels of uncoupling proteins 1 and 2 in mononuclear cells from patients with metabolic disorders: obesity and type 2 diabetes mellitus. Postepy Higieny I Medycyny Doswiadczalnej, 2017, 71, 0-0.	0.1	4
13	Genetic Structure of the Armenian Population. Archivum Immunologiae Et Therapiae Experimentalis, 2016, 64, 113-116.	2.3	5
14	Azokh Cave Hominin Remains. Vertebrate Paleobiology and Paleoanthropology, 2016, , 103-116.	0.5	13
15	Introduction: Azokh Cave and the Transcaucasian Corridor. Vertebrate Paleobiology and Paleoanthropology, 2016, , 1-26.	0.5	9
16	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206.	27.8	1,216
17	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	27.8	360
18	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.	3.3	14

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19	Coevolution of genes and languages and high levels of population structure among the highland populations of Daghestan. Journal of Human Genetics, 2016, 61, 181-191.	2.3	27
20	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
21	Population genomics of Bronze Age Eurasia. Nature, 2015, 522, 167-172.	27.8	1,166
22	Extensive genome-wide autozygosity in the population isolates of Daghestan. European Journal of Human Genetics, 2015, 23, 1405-1412.	2.8	21
23	The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. PLoS Genetics, 2015, 11, e1005068.	3.5	149
24	Global diversity, population stratification, and selection of human copy-number variation. Science, 2015, 349, aab3761.	12.6	293
25	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
26	Early Divergent Strains of Yersinia pestis in Eurasia 5,000 Years Ago. Cell, 2015, 163, 571-582.	28.9	425
27	The phylogenetic and geographic structure of Y-chromosome haplogroup R1a. European Journal of Human Genetics, 2015, 23, 124-131.	2.8	122
28	Different waves and directions of Neolithic migrations in the Armenian Highland. Investigative Genetics, 2014, 5, 15.	3.3	12
29	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
30	No Evidence from Genome-Wide Data of a Khazar Origin for the Ashkenazi Jews. Human Biology, 2013, 85, 859-900.	0.2	68
31	No Evidence from Genome-wide Data of a Khazar Origin fo the Ashkenazi Jews. Human Biology, 2013, 85, 859.	0.2	30
32	Complete Mitochondrial DNA Diversity in Iranians. PLoS ONE, 2013, 8, e80673.	2.5	93
33	Paternal Lineage Analysis Supports an Armenian Rather Than a Central Asian Genetic Origin of the Hamshenis. Human Biology, 2012, 84, 405-422.	0.2	1
34	Neolithic patrilineal signals indicate that the Armenian plateau was repopulated by agriculturalists. European Journal of Human Genetics, 2012, 20, 313-320.	2.8	33
35	Distinguishing the co-ancestries of haplogroup G Y-chromosomes in the populations of Europe and the Caucasus. European Journal of Human Genetics, 2012, 20, 1275-1282.	2.8	74
36	The Caucasus as an Asymmetric Semipermeable Barrier to Ancient Human Migrations. Molecular Biology and Evolution, 2012, 29, 359-365.	8.9	161

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37	Regionalized autosomal STR profiles among Armenian groups suggest disparate genetic influences. American Journal of Physical Anthropology, 2011, 146, 171-178.	2.1	10
38	Infant mortality decline in Armenia: Why with uneven rates?. Economics and Human Biology, 2010, 8, 134-137.	1.7	7
39	PLEISTOCENE TO HOLOCENE STRATIGRAPHY OF AZOKH 1 CAVE, LESSER CAUCASUS. Irish Journal of Earth Sciences, 2010, 28, 75-91.	0.3	13
40	Population genetics of familial Mediterranean fever: a review. European Journal of Human Genetics, 2007, 15, 911-916.	2.8	54
41	Infant mortality in Armenia, 1992â€“2003. Economics and Human Biology, 2006, 4, 351-358.	1.7	8
42	Armenian Y chromosome haplotypes reveal strong regional structure within a single ethno-national group. Human Genetics, 2001, 109, 659-674.	3.8	58