

# Levon Yepiskoposyan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11934352/publications.pdf>

Version: 2024-02-01

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	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. <i>Nature</i> , 2016, 538, 201-206.	27.8	1,216
2	Ancient human genomes suggest three ancestral populations for present-day Europeans. <i>Nature</i> , 2014, 513, 409-413.	27.8	1,179
3	Population genomics of Bronze Age Eurasia. <i>Nature</i> , 2015, 522, 167-172.	27.8	1,166
4	Early Divergent Strains of <i>Yersinia pestis</i> in Eurasia 5,000 Years Ago. <i>Cell</i> , 2015, 163, 571-582.	28.9	425
5	Genomic analyses inform on migration events during the peopling of Eurasia. <i>Nature</i> , 2016, 538, 238-242.	27.8	360
6	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. <i>Genome Research</i> , 2015, 25, 459-466.	5.5	348
7	137 ancient human genomes from across the Eurasian steppes. <i>Nature</i> , 2018, 557, 369-374.	27.8	325
8	Global diversity, population stratification, and selection of human copy-number variation. <i>Science</i> , 2015, 349, aab3761.	12.6	293
9	The Caucasus as an Asymmetric Semipermeable Barrier to Ancient Human Migrations. <i>Molecular Biology and Evolution</i> , 2012, 29, 359-365.	8.9	161
10	The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. <i>PLoS Genetics</i> , 2015, 11, e1005068.	3.5	149
11	The genetic history of admixture across inner Eurasia. <i>Nature Ecology and Evolution</i> , 2019, 3, 966-976.	7.8	135
12	The phylogenetic and geographic structure of Y-chromosome haplogroup R1a. <i>European Journal of Human Genetics</i> , 2015, 23, 124-131.	2.8	122
13	Complete Mitochondrial DNA Diversity in Iranians. <i>PLoS ONE</i> , 2013, 8, e80673.	2.5	93
14	Distinguishing the co-ancestries of haplogroup G Y-chromosomes in the populations of Europe and the Caucasus. <i>European Journal of Human Genetics</i> , 2012, 20, 1275-1282.	2.8	74
15	No Evidence from Genome-Wide Data of a Khazar Origin for the Ashkenazi Jews. <i>Human Biology</i> , 2013, 85, 859-900.	0.2	68
16	Armenian Y chromosome haplotypes reveal strong regional structure within a single ethno-national group. <i>Human Genetics</i> , 2001, 109, 659-674.	3.8	58
17	Population genetics of familial Mediterranean fever: a review. <i>European Journal of Human Genetics</i> , 2007, 15, 911-916.	2.8	54
18	Ancient pathogen DNA in human teeth and petrous bones. <i>Ecology and Evolution</i> , 2018, 8, 3534-3542.	1.9	38

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19	Eight Millennia of Matrilineal Genetic Continuity in the South Caucasus. <i>Current Biology</i> , 2017, 27, 2023-2028.e7.	3.9	37
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. <i>PLoS ONE</i> , 2015, 10, e0122968.	2.5	35
21	Neolithic patrilineal signals indicate that the Armenian plateau was repopulated by agriculturalists. <i>European Journal of Human Genetics</i> , 2012, 20, 313-320.	2.8	33
22	No Evidence from Genome-wide Data of a Khazar Origin fo the Ashkenazi Jews. <i>Human Biology</i> , 2013, 85, 859.	0.2	30
23	Coevolution of genes and languages and high levels of population structure among the highland populations of Daghestan. <i>Journal of Human Genetics</i> , 2016, 61, 181-191.	2.3	27
24	Origin and diffusion of human Y chromosome haplogroup J1-M267. <i>Scientific Reports</i> , 2021, 11, 6659.	3.3	26
25	Origin and spread of human mitochondrial DNA haplogroup U7. <i>Scientific Reports</i> , 2017, 7, 46044.	3.3	25
26	Extensive genome-wide autozygosity in the population isolates of Daghestan. <i>European Journal of Human Genetics</i> , 2015, 23, 1405-1412.	2.8	21
27	East Eurasian ancestry in the middle of Europe: genetic footprints of Steppe nomads in the genomes of Belarusian Lipka Tatars. <i>Scientific Reports</i> , 2016, 6, 30197.	3.3	14
28	Azokh Cave Hominin Remains. <i>Vertebrate Paleobiology and Paleoanthropology</i> , 2016, , 103-116.	0.5	13
29	The genetic variation in the R1a clade among the Ashkenazi Levitesâ€™™ Y chromosome. <i>Scientific Reports</i> , 2017, 7, 14969.	3.3	13
30	PLEISTOCENE TO HOLOCENE STRATIGRAPHY OF AZOKH 1 CAVE, LESSER CAUCASUS. <i>Irish Journal of Earth Sciences</i> , 2010, 28, 75-91.	0.3	13
31	Different waves and directions of Neolithic migrations in the Armenian Highland. <i>Investigative Genetics</i> , 2014, 5, 15.	3.3	12
32	Regionalized autosomal STR profiles among Armenian groups suggest disparate genetic influences. <i>American Journal of Physical Anthropology</i> , 2011, 146, 171-178.	2.1	10
33	Introduction: Azokh Cave and the Transcaucasian Corridor. <i>Vertebrate Paleobiology and Paleoanthropology</i> , 2016, , 1-26.	0.5	9
34	Insights into matrilineal genetic structure, differentiation and ancestry of Armenians based on complete mitogenome data. <i>Molecular Genetics and Genomics</i> , 2019, 294, 1547-1559.	2.1	9
35	Infant mortality in Armenia, 1992â€™“2003. <i>Economics and Human Biology</i> , 2006, 4, 351-358.	1.7	8
36	Infant mortality decline in Armenia: Why with uneven rates?. <i>Economics and Human Biology</i> , 2010, 8, 134-137.	1.7	7

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37	Genetic Structure of the Armenian Population. <i>Archivum Immunologiae Et Therapiae Experimentalis</i> , 2016, 64, 113-116.	2.3	5
38	Ancient DNA shows high faunal diversity in the Lesser Caucasus during the Late Pleistocene. <i>Quaternary Science Reviews</i> , 2019, 219, 102-111.	3.0	5
39	The mRNA expression levels of uncoupling proteins 1 and 2 in mononuclear cells from patients with metabolic disorders: obesity and type 2 diabetes mellitus. <i>Postepy Higieny I Medycyny Doswiadczalnej</i> , 2017, 71, 0-0.	0.1	4
40	Prevalence of uncoupling protein one genetic polymorphisms and their relationship with cardiovascular and metabolic health. <i>PLoS ONE</i> , 2022, 17, e0266386.	2.5	2
41	Paternal Lineage Analysis Supports an Armenian Rather Than a Central Asian Genetic Origin of the Hamshenis. <i>Human Biology</i> , 2012, 84, 405-422.	0.2	1
42	Demographic cognitive patterns revealed from human genome. , 2017, , .		0