

Ana Goios

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

682
citations

759233

12
h-index

580821

25
g-index

27
all docs

27
docs citations

27
times ranked

1094
citing authors

#	ARTICLE	IF	CITATIONS
1	High-resolution mtDNA evidence for the late-glacial resettlement of Europe from an Iberian refugium. <i>Genome Research</i> , 2005, 15, 19-24.	5.5	137
2	mtDNA phylogeny and evolution of laboratory mouse strains. <i>Genome Research</i> , 2007, 17, 293-298.	5.5	96
3	The global epidemiology of chikungunya from 1999 to 2020: A systematic literature review to inform the development and introduction of vaccines. <i>PLoS Neglected Tropical Diseases</i> , 2022, 16, e0010069.	3.0	65
4	Reconstructing the Indian Origin and Dispersal of the European Roma: A Maternal Genetic Perspective. <i>PLoS ONE</i> , 2011, 6, e15988.	2.5	61
5	Post-last glacial maximum expansion from Iberia to North Africa revealed by fine characterization of mtDNA H haplogroup in Tunisia. <i>American Journal of Physical Anthropology</i> , 2009, 139, 253-260.	2.1	54
6	The GHEP-EMPOP collaboration on mtDNA population data—A new resource for forensic casework. <i>Forensic Science International: Genetics</i> , 2011, 5, 146-151.	3.1	41
7	Human mtDNA haplogroups and reduced male fertility: real association or hidden population substructuring. <i>Journal of Developmental and Physical Disabilities</i> , 2005, 28, 241-247.	3.6	29
8	Specificity of mtDNA-directed PCR—influence of Nuclear MTDNA insertion (NUMT) contamination in routine samples and techniques. <i>International Journal of Legal Medicine</i> , 2008, 122, 341-345.	2.2	28
9	Y-chromosomal STR haplotypes in three ethnic groups and one cosmopolitan population from Tunisia. <i>Forensic Science International</i> , 2005, 152, 95-99.	2.2	26
10	Admixture and Genetic Diversity Distribution Patterns of Non-Recombining Lineages of Native American Ancestry in Colombian Populations. <i>PLoS ONE</i> , 2015, 10, e0120155.	2.5	22
11	Species identification in forensic samples using the SPInDel approach: A GHEP-ISFG inter-laboratory collaborative exercise. <i>Forensic Science International: Genetics</i> , 2017, 28, 219-224.	3.1	19
12	Evaluating the forensic informativeness of mtDNA haplogroup H sub-typing on a Eurasian scale. <i>Forensic Science International</i> , 2006, 159, 43-50.	2.2	17
13	Sequence Comparison Alignment-Free Approach Based on Suffix Tree and <i>L</i> -Words Frequency. <i>Scientific World Journal</i> , The, 2012, 2012, 1-4.	2.1	13
14	Identifying NUMT contamination in mtDNA analyses. <i>Forensic Science International: Genetics Supplement Series</i> , 2009, 2, 278-280.	0.3	12
15	RepeatAround: A software tool for finding and visualizing repeats in circular genomes and its application to a human mtDNA database. <i>Mitochondrion</i> , 2006, 6, 218-224.	3.4	10
16	Portuguese mitochondrial DNA genetic diversity—An update and a phylogenetic revision. <i>Forensic Science International: Genetics</i> , 2015, 15, 27-32.	3.1	10
17	Mitochondrial DNA pseudogenes in the nuclear genome as possible sources of contamination. <i>International Congress Series</i> , 2006, 1288, 697-699.	0.2	7
18	Identification of mouse inbred strains through mitochondrial DNA single-nucleotide extension. <i>Electrophoresis</i> , 2008, 29, 4795-4802.	2.4	7

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19	Analysis of uni and bi-parental markers in mixture samples: Lessons from the 22nd GHEP-ISFG Intercomparison Exercise. <i>Forensic Science International: Genetics</i> , 2016, 25, 63-72.	3.1	7
20	Mitochondrial DNA-control region sequence variation in the NE Portuguese Jewish community. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e51-e52.	0.3	5
21	mtDNAoffice: A software to assign human mtDNA macro haplogroups through automated analysis of the protein coding region. <i>Mitochondrion</i> , 2012, 12, 666-668.	3.4	4
22	Evaluation of heteroplasmy detection in the Ion Torrent PGM. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e13-e15.	0.3	4
23	mtDNA single macrodeletions associated with myopathies: Absence of haplogroup-related increased risk. <i>Journal of Inherited Metabolic Disease</i> , 2005, 28, 769-778.	3.6	3
24	A new algorithm for mtDNA sequence clustering. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e315-e316.	0.3	3
25	The Islamization of Iberian Peninsula: A demographic shift or a cultural change? Search for an answer using extant and ancient DNA from MÃ©rtola (Southeast Portugal). <i>International Congress Series</i> , 2006, 1288, 828-830.	0.2	2
26	Sampling efficiency for Amerindian female lineages. <i>International Congress Series</i> , 2006, 1288, 322-324.	0.2	0
27	A SNaPshot multiplex typing strategy for mtDNA identification of mouse inbred strains. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 596-597.	0.3	0