

Christina Amory

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

24
papers

433
citations

12
h-index

20
g-index

28
ext. papers

550
ext. citations

5.4
avg, IF

3.55
L-index

#	Paper	IF	Citations
24	Mitochondrial DNA Footprints from Western Eurasia in Modern Mongolia.. <i>Frontiers in Genetics</i> , 2021 , 12, 819337	4.5	0
23	Evidence for multi-copy Mega-NUMTs in the human genome. <i>Nucleic Acids Research</i> , 2021 , 49, 1517-1531	10.1	15
22	Evaluation of DNA Extraction Methods Developed for Forensic and Ancient DNA Applications Using Bone Samples of Different Age. <i>Genes</i> , 2021 , 12,	4.2	7
21	Genetic and phylogeographic evidence for Jewish Holocaust victims at the Sobibor death camp. <i>Genome Biology</i> , 2021 , 22, 200	18.3	1
20	The lot-to-lot variability in the mitochondrial genome of controls. <i>Forensic Science International: Genetics</i> , 2020 , 47, 102298	4.3	4
19	The mitogenome portrait of Umbria in Central Italy as depicted by contemporary inhabitants and pre-Roman remains. <i>Scientific Reports</i> , 2020 , 10, 10700	4.9	4
18	Distinguishing mitochondrial DNA and NUMT sequences amplified with the precision ID mtDNA whole genome panel. <i>Mitochondrion</i> , 2020 , 55, 122-133	4.9	13
17	Developmental Validation of a MPS Workflow with a PCR-Based Short Amplicon Whole Mitochondrial Genome Panel. <i>Genes</i> , 2020 , 11,	4.2	11
16	House of the dead-exceptional burials of the Avar period (seventh century AD) in Podersdorf am See (Burgenland/A). <i>Archaeological and Anthropological Sciences</i> , 2020 , 12, 1	1.8	
15	Mitochondrial DNA analysis of a Viking age mass grave in Sweden. <i>Forensic Science International: Genetics</i> , 2019 , 42, 268-274	4.3	8
14	SD quants-Sensitive detection tetraplex-system for nuclear and mitochondrial DNA quantification and degradation inference. <i>Forensic Science International: Genetics</i> , 2019 , 42, 39-44	4.3	15
13	Evaluation of mitogenome sequence concordance, heteroplasmy detection, and haplogrouping in a worldwide lineage study using the Precision ID mtDNA Whole Genome Panel. <i>Forensic Science International: Genetics</i> , 2019 , 42, 244-251	4.3	22
12	The maternal inheritance of Alto Parani revealed by full mitogenome sequences. <i>Forensic Science International: Genetics</i> , 2019 , 39, 66-72	4.3	9
11	Evaluation of the precision ID whole MtDNA genome panel for forensic analyses. <i>Forensic Science International: Genetics</i> , 2018 , 35, 21-25	4.3	49
10	Working towards implementation of whole genome mitochondrial DNA sequencing into routine casework. <i>Forensic Science International: Genetics Supplement Series</i> , 2017 , 6, e388-e389	0.5	12
9	Lack of gene-language correlation due to reciprocal female but directional male admixture in Austronesians and non-Austronesians of East Timor. <i>European Journal of Human Genetics</i> , 2017 , 25, 246-252	5.3	7
8	Optimized mtDNA Control Region Primer Extension Capture Analysis for Forensically Relevant Samples and Highly Compromised mtDNA of Different Age and Origin. <i>Genes</i> , 2017 , 8,	4.2	34

7	Human settlement history between Sunda and Sahul: a focus on East Timor (Timor-Leste) and the Pleistocenic mtDNA diversity. <i>BMC Genomics</i> , 2015 , 16, 70	4-5	28
6	Full mtGenome reference data: development and characterization of 588 forensic-quality haplotypes representing three U.S. populations. <i>Forensic Science International: Genetics</i> , 2015 , 14, 141-55	4-3	59
5	Evidence for frequent and tissue-specific sequence heteroplasmy in human mitochondrial DNA. <i>Mitochondrion</i> , 2015 , 20, 82-94	4-9	66
4	Helena, the hidden beauty: Resolving the most common West Eurasian mtDNA control region haplotype by massively parallel sequencing an Italian population sample. <i>Forensic Science International: Genetics</i> , 2015 , 15, 21-6	4-3	20
3	Development of forensic-quality full mtGenome haplotypes: success rates with low template specimens. <i>Forensic Science International: Genetics</i> , 2014 , 10, 73-79	4-3	15
2	Reprint of: Evaluation of next generation mtGenome sequencing using the Ion Torrent Personal Genome Machine (PGM). <i>Forensic Science International: Genetics</i> , 2013 , 7, 632-639	4-3	32
1	Social Relations, Deprivation and Violence at Schleinbach, Lower Austria. Insights from an Interdisciplinary Analysis of the Early Bronze Age Human Remains. <i>Archaeologica Austriaca</i> , Band 104/2020, 13-52		2