Catarina Xavier

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/8866935/catarina-xavier-publications-by-year.pdf

Version: 2024-04-11

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

32	523	14	22
papers	citations	h-index	g-index
33 ext. papers	761 ext. citations	4.6 avg, IF	3.94 L-index

#	Paper	IF	Citations
32	Evaluation of the VISAGE basic tool for appearance and ancestry inference using ForenSeq ^[] chemistry on the MiSeq FGx ^[] system <i>Forensic Science International: Genetics</i> , 2022 , 58, 102675	4.3	O
31	The Ancestry of Eastern Paraguay: A Typical South American Profile with a Unique Pattern of Admixture. <i>Genes</i> , 2021 , 12,	4.2	1
30	Development of the VISAGE enhanced tool and statistical models for epigenetic age estimation in blood, buccal cells and bones. <i>Aging</i> , 2021 , 13, 6459-6484	5.6	11
29	Evidence for multi-copy Mega-NUMTs in the human genome. <i>Nucleic Acids Research</i> , 2021 , 49, 1517-153	1 20.1	15
28	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
27	Development and Evaluation of the Ancestry Informative Marker Panel of the VISAGE Basic Tool. <i>Genes</i> , 2021 , 12,	4.2	2
26	Epigenetic age prediction in semen - marker selection and model development. <i>Aging</i> , 2021 , 13, 19145-	1 9 .664	1
25	Impact of excessive alcohol abuse on age prediction using the VISAGE enhanced tool for epigenetic age estimation in blood. <i>International Journal of Legal Medicine</i> , 2021 , 135, 2209-2219	3.1	1
24	Broadening the Applicability of a Custom Multi-Platform Panel of Microhaplotypes: Bio-Geographical Ancestry Inference and Expanded Reference Data. <i>Frontiers in Genetics</i> , 2020 , 11, 581	04:₹	8
23	Development and optimization of the VISAGE basic prototype tool for forensic age estimation. <i>Forensic Science International: Genetics</i> , 2020 , 48, 102322	4.3	13
22	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
21	Evaluation of the VISAGE Basic Tool for Appearance and Ancestry Prediction Using PowerSeq Chemistry on the MiSeq FGx System. <i>Genes</i> , 2020 , 11,	4.2	15
20	Development and validation of the VISAGE AmpliSeq basic tool to predict appearance and ancestry from DNA. <i>Forensic Science International: Genetics</i> , 2020 , 48, 102336	4.3	22
19	Body fluid identification and assignment to donors using a targeted mRNA massively parallel sequencing approach - results of a second EUROFORGEN / EDNAP collaborative exercise. <i>Forensic Science International: Genetics</i> , 2020 , 45, 102208	4.3	9
18	Building a custom large-scale panel of novel microhaplotypes for forensic identification using MiSeq and Ion S5 massively parallel sequencing systems. <i>Forensic Science International: Genetics</i> , 2020 , 45, 102213	4.3	27
17	Forensic evaluation of the Asia Pacific ancestry-informative MAPlex assay. <i>Forensic Science International: Genetics</i> , 2020 , 48, 102344	4.3	8
16	DNA Testing Reveals the Putative Identity of JB55, a 19th Century Vampire Buried in Griswold, Connecticut. <i>Genes</i> , 2019 , 10,	4.2	4

LIST OF PUBLICATIONS

15	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	
14	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	
13	Mitochondrial DNA control region variation in Lebanon, Jordan, and Bahrain. <i>Forensic Science International: Genetics</i> , 2019 , 42, 99-102	4.3	4	
12	HIrisPlex-S system for eye, hair, and skin color prediction from DNA: Massively parallel sequencing solutions for two common forensically used platforms. <i>Forensic Science International: Genetics</i> , 2019 , 43, 102152	4.3	24	
11	The maternal inheritance of the Ashaninka native group from Peru. <i>Forensic Science International: Genetics Supplement Series</i> , 2019 , 7, 135-137	0.5	O	
10	The maternal inheritance of Alto Paranirevealed by full mitogenome sequences. <i>Forensic Science International: Genetics</i> , 2019 , 39, 66-72	4.3	9	
9	Body fluid identification using a targeted mRNA massively parallel sequencing approach - results of a EUROFORGEN/EDNAP collaborative exercise. <i>Forensic Science International: Genetics</i> , 2018 , 34, 105-17	1 \$.3	42	
8	Resolving the matrilineal relationship of seven Late Bronze Age individuals from Stillfried, Austria. <i>Forensic Science International: Genetics</i> , 2018 , 36, 148-151	4.3	5	
7	Evaluation of the Illumina ForenSeqIDNA Signature Prep Kit - MPS forensic application for the MiSeq FGxIbenchtop sequencer. <i>Forensic Science International: Genetics</i> , 2017 , 28, 188-194	4.3	61	
6	Paraguay: Unveiling migration patterns with ancestry genetic markers. <i>Forensic Science International: Genetics Supplement Series</i> , 2017 , 6, e226-e228	0.5	3	
5	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	•
4	High-quality mtDNA control region sequences from 680 individuals sampled across the Netherlands to establish a national forensic mtDNA reference database. <i>Forensic Science International: Genetics</i> , 2016 , 21, 158-67	4.3	15	
3	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, 141-5	5 4·3	59	
2	Admixture and genetic diversity distribution patterns of non-recombining lineages of Native American ancestry in Colombian populations. <i>PLoS ONE</i> , 2015 , 10, e0120155	3.7	16	
1	Massively parallel sequencing of complete mitochondrial genomes from hair shaft samples. <i>Forensic Science International: Genetics</i> , 2015 , 15, 8-15	4.3	66	