

Vania Pereira

List of Publications by Year in descending order

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38
papers

688
citations

567281

15
h-index

580821

25
g-index

39
all docs

39
docs citations

39
times ranked

733
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic analysis of sixteen autosomal STR loci in three Tunisian populations from Makthar, Nabeul and Sousse. <i>Annals of Human Biology</i> , 2022, , 1-22.	1.0	1
2	Testing the Ion AmpliSeq, HID Y-SNP Research Panel v1 for performance and resolution in admixed South Americans of haplogroup Q. <i>Forensic Science International: Genetics</i> , 2022, 59, 102708.	3.1	3
3	Evaluation of a custom GeneRead, massively parallel sequencing assay with 210 ancestry informative SNPs using the Ion S5, and MiSeq platforms. <i>Forensic Science International: Genetics</i> , 2021, 50, 102411.	3.1	7
4	Ancestry prediction efficiency of the software GenoGeographer using a z-score method and the ancestry informative markers in the Precision ID Ancestry Panel. <i>Forensic Science International: Genetics</i> , 2020, 44, 102154.	3.1	19
5	Evaluation of the Precision of Ancestry Inferences in South American Admixed Populations. <i>Frontiers in Genetics</i> , 2020, 11, 966.	2.3	10
6	Skin pigmentation and genetic variants in an admixed Brazilian population of primarily European ancestry. <i>International Journal of Legal Medicine</i> , 2020, 134, 1569-1579.	2.2	4
7	Paternal and maternal mutations in X-STRs: A GHEP-ISFG collaborative study. <i>Forensic Science International: Genetics</i> , 2020, 46, 102258.	3.1	10
8	Genetic relationships of European, Mediterranean, and SW Asian populations using a panel of 55 AISNPs. <i>European Journal of Human Genetics</i> , 2019, 27, 1885-1893.	2.8	22
9	Development and validation of the EUROFORGEN NAME (North African and Middle Eastern) ancestry panel. <i>Forensic Science International: Genetics</i> , 2019, 42, 260-267.	3.1	46
10	Analysis of 16 autosomal STR loci in Uyghur and Kazakh populations from Xinjiang, China. <i>Forensic Science International: Genetics</i> , 2019, 40, e262-e263.	3.1	3
11	GENETIC PORTRAIT OF THE PUNJABI POPULATION FROM PAKISTAN USING THE PRECISION ID ANCESTRY PANEL. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 87-89.	0.3	1
12	Analysis of mainland Japanese and Okinawan Japanese populations using the precision ID Ancestry Panel. <i>Forensic Science International: Genetics</i> , 2018, 33, 106-109.	3.1	16
13	Sequencing of mitochondrial genomes using the Precision ID mtDNA Whole Genome Panel. <i>Electrophoresis</i> , 2018, 39, 2766-2775.	2.4	25
14	Evaluation of the Precision ID Ancestry Panel for crime case work: A SNP typing assay developed for typing of 165 ancestral informative markers. <i>Forensic Science International: Genetics</i> , 2017, 28, 138-145.	3.1	82
15	Typing of two Middle Eastern populations with the Precision ID Ancestry Panel. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e301-e302.	0.3	9
16	Analysis of 16 autosomal STR loci in Uyghur and Kazakh populations from Xinjiang, China. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e537-e538.	0.3	0
17	Analysis of ancestry informative markers in three main ethnic groups from Ecuador supports a trihybrid origin of Ecuadorians. <i>Forensic Science International: Genetics</i> , 2017, 31, 29-33.	3.1	40
18	Genomic and immunohistochemical characterisation of a lacrimal gland oncocytoma and review of literature. <i>Oncology Letters</i> , 2017, 14, 4176-4182.	1.8	15

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19	NGMSelect [®] and Investigator Argus X-12 analysis in population samples from Albania, Iraq, Lithuania, Slovenia, and Turkey. <i>Forensic Science International: Genetics</i> , 2016, 22, 110-112.	3.1	16
20	Importance of nonsynonymous <i>OCA2</i> variants in human eye color prediction. <i>Molecular Genetics & Genomic Medicine</i> , 2016, 4, 420-430.	1.2	31
21	A study of the peopling of Greenland using next generation sequencing of complete mitochondrial genomes. <i>American Journal of Physical Anthropology</i> , 2016, 161, 698-704.	2.1	15
22	Types of Genomes, Sequences and Genetic Markers (Repeats, SNPs, Indels, Haplotypes). <i>Security Science and Technology</i> , 2016, , 163-191.	0.5	4
23	Peopling of the North Circumpolar Region – Insights from Y Chromosome STR and SNP Typing of Greenlanders. <i>PLoS ONE</i> , 2015, 10, e0116573.	2.5	16
24	The peopling of Greenland: further insights from the analysis of genetic diversity using autosomal and X-chromosomal markers. <i>European Journal of Human Genetics</i> , 2015, 23, 245-251.	2.8	15
25	Mosaic maternal ancestry in the Great Lakes region of East Africa. <i>Human Genetics</i> , 2015, 134, 1013-1027.	3.8	18
26	Next-generation sequencing of multiple individuals per barcoded library by deconvolution of sequenced amplicons using endonuclease fragment analysis. <i>BioTechniques</i> , 2014, 57, 91-4.	1.8	5
27	Refining the genetic portrait of Portuguese Roma through X-chromosomal markers. <i>American Journal of Physical Anthropology</i> , 2012, 148, 389-394.	2.1	9
28	A method for the analysis of 32 X chromosome insertion deletion polymorphisms in a single PCR. <i>International Journal of Legal Medicine</i> , 2012, 126, 97-105.	2.2	45
29	Analysis of 12 X-STRs in Greenlanders, Danes and Somalis using Argus X-12. <i>International Journal of Legal Medicine</i> , 2012, 126, 121-128.	2.2	70
30	Genetic characterization of Somali and Iraqi populations using a set of 33 X-chromosome Indels. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e137-e138.	0.3	4
31	Study of 25 X-chromosome Single Nucleotide Polymorphisms in African and Asian populations. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e139-e140.	0.3	1
32	Study of 25 X-chromosome SNPs in the Portuguese. <i>Forensic Science International: Genetics</i> , 2011, 5, 336-338.	3.1	9
33	A framework for the development of STR genotyping in domestic animal species: Characterization and population study of 12 canine X-chromosome loci. <i>Electrophoresis</i> , 2010, 31, 303-308.	2.4	21
34	Genetic characterization of uniparental lineages in populations from Southwest Iberia with past malaria endemicity. <i>American Journal of Human Biology</i> , 2010, 22, 588-595.	1.6	6
35	A new autosomal STR nineplex for canine identification and parentage testing. <i>Electrophoresis</i> , 2009, 30, 417-423.	2.4	48
36	The Karimojong from Uganda: Genetic characterization using an X-STR decaplex system. <i>Forensic Science International: Genetics</i> , 2009, 3, e127-e128.	3.1	21

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37	Association between STRs from the X chromosome in a sample of Portuguese Gypsies. Forensic Science International: Genetics Supplement Series, 2009, 2, 391-393.	0.3	3
38	Genetic patterns of 10 X chromosome short tandem repeats in an Asian population from Macau. Forensic Science International: Genetics Supplement Series, 2009, 2, 402-404.	0.3	4