Maria Torres

List of Publications by Year
in descending order

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Version: 2024-02-01

31 papers	1,665 citations	430874 18 h-index	454955 30 g-index
33 all docs	33 docs citations	33 times ranked	4137 citing authors

#	Article	IF	Citations
1	Obesity-related genetic determinants of stroke. Brain Communications, 2021, 3, fcab069.	3.3	1
2	LIPG endothelial lipase and breast cancer risk by subtypes. Scientific Reports, 2021, 11, 10436.	3.3	2
3	Ancestry analysis using autosomal SNPs in northern South America, reveals interpretation differences between an AIM panel and an identification panel. Forensic Science International, 2021, 326, 110934.	2.2	1
4	Residential radon, genetic polymorphisms in DNA damage and repair-related. Lung Cancer, 2019, 135, 10-15.	2.0	21
5	A new panel of SNPs to assess thyroid carcinoma risk: a pilot study in a Brazilian admixture population. BMC Medical Genetics, 2017, 18, 140.	2.1	3
6	A European Spectrum of Pharmacogenomic Biomarkers: Implications for Clinical Pharmacogenomics. PLoS ONE, 2016, 11, e0162866.	2.5	96
7	Development of a methylation marker set for forensic age estimation using analysis of public methylation data and the Agena Bioscience EpiTYPER system. Forensic Science International: Genetics, 2016, 24, 65-74.	3.1	127
8	Investigation of geneâ€environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. International Journal of Cancer, 2015, 136, E685-96.	5.1	34
9	Sequenom MassARRAY approach in the arrhythmogenic right ventricular cardiomyopathy post-mortem setting: clinical and forensic implications. International Journal of Legal Medicine, 2015, 129, 1-10.	2.2	18
10	Abstract 2781: Genetic susceptibility to breast cancer in a Spanish population. , 2015, , .		0
11	Cuba: Exploring the History of Admixture and the Genetic Basis of Pigmentation Using Autosomal and Uniparental Markers. PLoS Genetics, 2014, 10, e1004488.	3.5	57
12	Building a forensic ancestry panel from the ground up: The EUROFORGEN Global AIM-SNP set. Forensic Science International: Genetics, 2014, 11, 13-25.	3.1	116
13	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	21.4	281
14	High-throughput genotyping assay for the large-scale genetic characterization ofCryptosporidiumparasites from human and bovine samples. Parasitology, 2014, 141, 491-500.	1.5	14
15	Association of thromboxane A1 synthase (TBXAS1) gene polymorphism with acute urticaria induced by nonsteroidal anti-inflammatory drugs. Journal of Allergy and Clinical Immunology, 2013, 132, 989-991.	2.9	21
16	Development of a Panel of Genome-Wide Ancestry Informative Markers to Study Admixture Throughout the Americas. PLoS Genetics, 2012, 8, e1002554.	3.5	212
17	New technologies in the genetic approach to sudden cardiac death in the young. Forensic Science International, 2010, 203, 15-24.	2.2	19
18	A new approach to long QT syndrome mutation detection by Sequenom MassARRAY^{Â®}system. Electrophoresis, 2010, 31, 1648-1655.	2.4	13

#	Article	IF	Citations
19	Common variants at 2q37.3, 8q24.21, 15q21.3 and 16q24.1 influence chronic lymphocytic leukemia risk. Nature Genetics, 2010, 42, 132-136.	21.4	223
20	Identification of a novel mutation in the human PDE6A gene in autosomal recessive retinitis pigmentosa: homology with the nmf28/nmf28 mice model. Clinical Genetics, 2010, 78, 495-498.	2.0	21
21	Genetic variation in the nuclear factor ÎºB pathway in relation to susceptibility to rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 579-583.	0.9	40
22	Involvement of hypertrophic cardiomyopathy genes in sudden infant death syndrome (SIDS). Forensic Science International: Genetics Supplement Series, 2009, 2, 495-496.	0.3	13
23	Sequenom MassArrayâ„¢ application in the long QT syndrome mutation detection. Forensic Science International: Genetics Supplement Series, 2009, 2, 497-498.	0.3	0
24	Association of schizophrenia with DTNBP1 but not with DAO, DAOA, NRG1 and RGS4 nor their genetic interaction. Journal of Psychiatric Research, 2008, 42, 278-288.	3.1	80
25	Analyses of variants located in estrogen metabolism genes (ESR1, ESR2, COMT and APOE) and schizophrenia. Schizophrenia Research, 2008, 100, 308-315.	2.0	23
26	Gene Expression Profile in Oral Squamous Cell Carcinoma: A Pilot Study. Journal of Oral and Maxillofacial Surgery, 2005, 63, 786-792.	1.2	13
27	Relative efficiency of the linkage disequilibrium mapping approach in detecting candidate genes for schizophrenia in different European populations. Genomics, 2005, 86, 280-286.	2.9	9
28	Analysis ofBRCA1andBRCA2genes in Spanish breast/ovarian cancer patients: A high proportion of mutations unique to Spain and evidence of founder effects. Human Mutation, 2003, 22, 301-312.	2.5	154
29	A Novel Loss-of-Function Mutation (N48K) in the PTEN Gene in a Spanish Patient with Cowden Disease. Journal of Investigative Dermatology, 2003, 121, 1356-1359.	0.7	15
30	Analysis of BRCA1 and BRCA2 in breast and breast/ovarian cancer families shows population substructure in the Iberian peninsula. Annals of Human Genetics, 2002, 66, 29-36.	0.8	30
31	BRCA1 and BRCA2 mutations in breast and breast/ovarian cancer families from Galicia (NW Spain). European Journal of Cancer, 2001, 37, S178.	2.8	0