Maria Torres

List of Publications by Year
in descending order

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	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	21.4	281
2	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
3	Development of a Panel of Genome-Wide Ancestry Informative Markers to Study Admixture Throughout the Americas. PLoS Genetics, 2012, 8, e1002554.	3.5	212
4	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
5	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
6	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
7	A European Spectrum of Pharmacogenomic Biomarkers: Implications for Clinical Pharmacogenomics. PLoS ONE, 2016, 11, e0162866.	2.5	96
8	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
9	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
10	Genetic variation in the nuclear factor Î¹B pathway in relation to susceptibility to rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 579-583.	0.9	40
11	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
12	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
13	Analyses of variants located in estrogen metabolism genes (ESR1, ESR2, COMT and APOE) and schizophrenia. Schizophrenia Research, 2008, 100, 308-315.	2.0	23
14	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
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	Residential radon, genetic polymorphisms in DNA damage and repair-related. Lung Cancer, 2019, 135, 10-15.	2.0	21
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	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

#	Article	IF	Citations
19	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
20	High-throughput genotyping assay for the large-scale genetic characterization ofCryptosporidiumparasites from human and bovine samples. Parasitology, 2014, 141, 491-500.	1.5	14
21	Gene Expression Profile in Oral Squamous Cell Carcinoma: A Pilot Study. Journal of Oral and Maxillofacial Surgery, 2005, 63, 786-792.	1.2	13
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	A new approach to long QT syndrome mutation detection by Sequenom MassARRAY^{Â®} system. Electrophoresis, 2010, 31, 1648-1655.	2.4	13
24	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
25	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
26	LIPG endothelial lipase and breast cancer risk by subtypes. Scientific Reports, 2021, 11, 10436.	3.3	2
27	Obesity-related genetic determinants of stroke. Brain Communications, 2021, 3, fcab069.	3.3	1
28	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
29	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
30	Sequenom MassArrayâ„¢ application in the long QT syndrome mutation detection. Forensic Science International: Genetics Supplement Series, 2009, 2, 497-498.	0.3	0
31	Abstract 2781: Genetic susceptibility to breast cancer in a Spanish population. , 2015, , .		0