Maria Martinez

List of Publications by Year
in descending order

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Version: 2024-02-01

96 papers	7,805 citations	94433 37 h-index	51608 86 g-index
99 all docs	99 docs citations	99 times ranked	11933 citing authors

#	Article	IF	Citations
1	Dark Matter Searches Using NaI(Tl) at the Canfranc Underground Laboratory: Past, Present and Future. Universe, 2022, 8, 75.	2.5	1
2	Transient nozzle flow analysis and near field characterization of gasoline direct fuel injector using Large Eddy Simulation. International Journal of Multiphase Flow, 2022, 148, 103920.	3.4	10
3	Computational study of the Premixed Charge Compression Ignition combustion in a Rapid Compression Expansion Machine: Impact of multiple injection strategy on mixing, ignition and combustion processes. Fuel, 2022, 318, 123388.	6.4	2
4	Validation of a three-phase Eulerian CFD model to account for cavitation and spray atomization phenomena. Journal of the Brazilian Society of Mechanical Sciences and Engineering, 2021, 43, 1.	1.6	3
5	Dead in life. Lives pierced by death. Death Studies, 2020, 44, 677-680.	2.7	2
6	Living dead: Suspended lives during/after gender violence. Death Studies, 2020, 44, 718-726.	2.7	15
7	Inverted Exception. Ideas for Thinking about the New Disappearances through Two Case Studies. Journal of Latin American Cultural Studies, 2020, 29, 581-604.	0.2	3
8	Naming what has no name: thinking about migrant â€œdisappearanceâ€•at the Mexico-US border (Arizona). Athenea Digital, 2020, 20, 2692.	0.2	2
9	The exceeded disappearance. Introduction. Athenea Digital, 2020, 20, 2874.	0.2	1
10	Result on the Neutrinoless Double Beta Decay Search of 82Se with the CUPID-0 Experiment. Universe, 2019, 5, 2.	2.5	0
11	Phonon traps reduce the quasiparticle density in superconducting circuits. Applied Physics Letters, 2019, 115, .	3.3	34
12	Results from the Cuore Experiment â€. Universe, 2019, 5, 10.	2.5	5
13	IntroducciÃ³n. La desapariciÃ³n forzada de personas: circulaciÃ³n transnacional y usos sociales de una categorÃa de los derechos humanos (Introduction. The forced disappearance of persons:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 2019, 9, 145-154.	0.4	1
14	MÃ¡s allÃ¡ de los tÃ³picos en el estudio de la desapariciÃ³n. Hacia el desaparecido social (Beyond the) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 9, 169-182.	0.4	3
15	DesapariciÃ³n. La vida en sus lÃmites (Disappearance. Life in the limits). Onati Socio-Legal Series, 2019, 9, 222-236.	0.4	4
16	Una (breve y no muy sistemÃ¡tica) aproximaciÃ³n a la nociÃ³n de agencia desde la vulnerabilidad. Papeles Del CEIC: International Journal on Collective Identity Research, 2019, 2019, .	0.2	4
17	La â€œtÃ©cnicaâ€•de la desapariciÃ³n (The â€œtechniqueâ€•of disappearance). Onati Socio-Legal Series, 2019, 9, 252-266.	0.4	0
18	From the Subjected Subject to the Vulnerable Subject: An Unfinished Discussion in Contemporary Spanish Feminisms. Signs, 2018, 43, 327-351.	0.9	4

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19	CALDER: The Second-Generation Light Detectors. IEEE Transactions on Applied Superconductivity, 2018, 28, 1-3.	1.7	1
20	Relational iterations and emotional activations: towards a radicalization of processuality of collective identities. Athenea Digital, 2018, 18, 293.	0.2	1
21	Desapariciones. Usos locales, circulaciones globales. Gabriel Gatti (Ed.). BogotÃ¡: Siglo del Hombre Editores - Universidad de los Andes, 2017. 288 pp.. Universitas HumanÃstica, 2018, 86, .	0.3	4
22	Phonon-Mediated KIDs as Light Detectors for Rare Event Search: The CALDER Project. IEEE Transactions on Applied Superconductivity, 2017, 27, 1-5.	1.7	1
23	When school meets the Other: intercultural policies in the Basque inclusive school. Patterns of Prejudice, 2017, 51, 89-110.	0.6	2
24	Transethnic meta-analysis identifies GSDMA and PRDM1 as susceptibility genes to systemic sclerosis. Annals of the Rheumatic Diseases, 2017, 76, 1150-1158.	0.9	77
25	Ce que la loi fait aux victimes. Construction de la figure de la Â«Âfemme-victimeÂÂ» par la loi contre la violence de genre en Espagne. Pensee Plurielle, 2017, nÂ° 45, 123-138.	0.2	0
26	La vÃctima simulada. Identidades forzadas en la violencia de gÃ©nero. Politica Y Sociedad, 2016, 53, 879-896.	0.2	7
27	Neutron Spectrometry With Scintillating Bolometers of LiF and Sapphire. IEEE Transactions on Nuclear Science, 2016, 63, 1967-1975.	2.0	3
28	Further support for the association of GNPAT variant rs11558492 with severe iron overload in hemochromatosis. Hepatology, 2016, 63, 2054-2055.	7.3	14
29	Fragmented victims: Women victims of gender-based violence in the face of expert discourses and practices in Spain. Women's Studies International Forum, 2016, 59, 39-47.	1.1	1
30	WES/WGS Reporting of Mutations from Cardiovascular â€œActionableâ€•Genes in Clinical Practice: A Key Role for UMD Knowledgebases in the Era of Big Databases. Human Mutation, 2016, 37, 1308-1317.	2.5	5
31	Les victimes peuvent-elles parler et agirÂ? Deux paradoxes Ã lâ€™Ã¨re des citoyens-victimes. Pensee Plurielle, 2016, nÂ° 43, 155-167.	0.2	4
32	NeuroX, a fast and efficient genotyping platform for investigation of neurodegenerative diseases. Neurobiology of Aging, 2015, 36, 1605.e7-1605.e12.	3.1	96
33	Diagnosis of Parkinson's disease on the basis of clinical and genetic classification: a population-based modelling study. Lancet Neurology, The, 2015, 14, 1002-1009.	10.2	179
34	RARE EVENT SEARCHES AT CANFRANC: ANAIS EXPERIMENT. , 2015, , .		0
35	Response of parylene-coated NaI(Tl) scintillators at low temperature. EPJ Web of Conferences, 2014, 65, 02001.	0.3	3
36	Genetic comorbidities in Parkinson's disease. Human Molecular Genetics, 2014, 23, 831-841.	2.9	57

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37	Large-scale meta-analysis of genome-wide association data identifies six new risk loci for Parkinson's disease. Nature Genetics, 2014, 46, 989-993.	21.4	1,685
38	Single-marker and multi-marker mixed models for polygenic score analysis in family-based data. BMC Proceedings, 2014, 8, S63.	1.6	1
39	Analysis of Genome-Wide Association Studies of Alzheimer Disease and of Parkinson Disease to Determine If These 2 Diseases Share a Common Genetic Risk. JAMA Neurology, 2013, 70, 1268-76.	9.0	51
40	A pathway-based analysis provides additional support for an immune-related genetic susceptibility to Parkinson's disease. Human Molecular Genetics, 2013, 22, 1039-1049.	2.9	122
41	The Val158Met COMT polymorphism is a modifier of the age at onset in Parkinson's disease with a sexual dimorphism. Journal of Neurology, Neurosurgery and Psychiatry, 2013, 84, 666-673.	1.9	43
42	Using genome-wide complex trait analysis to quantify 'missing heritability' in Parkinson's disease. Human Molecular Genetics, 2012, 21, 4996-5009.	2.9	176
43	Comprehensive Research Synopsis and Systematic Meta-Analyses in Parkinson's Disease Genetics: The PDGene Database. PLoS Genetics, 2012, 8, e1002548.	3.5	495
44	Evidence for caveolin-1 as a new susceptibility gene regulating tissue fibrosis in systemic sclerosis. Annals of the Rheumatic Diseases, 2012, 71, 1034-1041.	0.9	33
45	Use of support vector machines for disease risk prediction in genome-wide association studies: Concerns and opportunities. Human Mutation, 2012, 33, 1708-1718.	2.5	42
46	Cooperative Genome-Wide Analysis Shows Increased Homozygosity in Early Onset Parkinson's Disease. PLoS ONE, 2012, 7, e28787.	2.5	21
47	Bivariate association analysis in selected samples: application to a GWAS of two bone mineral density phenotypes in males with high or low BMD. European Journal of Human Genetics, 2011, 19, 710-716.	2.8	17
48	Imputation of sequence variants for identification of genetic risks for Parkinson's disease: a meta-analysis of genome-wide association studies. Lancet, The, 2011, 377, 641-649.	13.7	845
49	Comparative study of statistical methods for detecting association with rare variants in exome-resequencing data. BMC Proceedings, 2011, 5, S33.	1.6	5
50	Genome-wide association study confirms BST1 and suggests a locus on 12q24 as the risk loci for Parkinson's disease in the European population. Human Molecular Genetics, 2011, 20, 615-627.	2.9	155
51	Genome-Wide Scan Identifies TNIP1, PSORS1C1, and RHOB as Novel Risk Loci for Systemic Sclerosis. PLoS Genetics, 2011, 7, e1002091.	3.5	205
52	The p.Asp216His TOR1A allele effect is not found in the French population. Movement Disorders, 2009, 24, 919-921.	3.9	15
53	No Significant Association of 14 Candidate Genes With Schizophrenia in a Large European Ancestry Sample: Implications for Psychiatric Genetics. American Journal of Psychiatry, 2008, 165, 497-506.	7.2	323
54	Genome-Wide Linkage Screen of Bone Mineral Density (BMD) in European Pedigrees Ascertained through a Male Relative with Low BMD Values: Evidence for Quantitative Trait Loci on 17q21â€“23, 11q12â€“13, 13q12â€“14, and 22q11. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 3755-3762.	3.6	44

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55	<i>DTNBP1 (Dystrobrevin Binding Protein 1)</i> and Schizophrenia: Association Evidence in the 3â€² End of the Gene. Human Heredity, 2007, 64, 97-106.	0.8	35
56	LRRK2emph Exon 41 Mutations in Sporadic Parkinson Disease in Europeans. Archives of Neurology, 2007, 64, 425.	4.5	51
57	Gene expression profiling of Hfe-/- liver and duodenum in mouse strains with differing susceptibilities to iron loading: identification of transcriptional regulatory targets of Hfe and potential hemochromatosis modifiers. Genome Biology, 2007, 8, R221.	9.6	16
58	Summary of contributions to GAW15 Group 16: Processing/normalization of expression traits. Genetic Epidemiology, 2007, 31, S132-S138.	1.3	2
59	Genomewide Linkage Scan of 409 European-Ancestry and African American Families with Schizophrenia: Suggestive Evidence of Linkage at 8p23.3-p21.2 and 11p13.1-q14.1 in the Combined Sample. American Journal of Human Genetics, 2006, 78, 315-333.	6.2	141
60	Chromosome 5 and Parkinson disease. European Journal of Human Genetics, 2006, 14, 1106-1110.	2.8	4
61	Neuregulin 1 (NRG1) and schizophrenia: analysis of a US family sample and the evidence in the balance. Psychological Medicine, 2005, 35, 1599-1610.	4.5	46
62	Apolipoprotein E4 is probably responsible for the chromosome 19 linkage peak for Parkinson's disease. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 136B, 72-74.	1.7	27
63	A non-DM1, non-DM2 multisystem myotonic disorder with frontotemporal dementia: phenotype and suggestive mapping of the DM3 locus to chromosome 15q21-24. Brain, 2004, 127, 1979-1992.	7.6	38
64	Polymorphisms in the Trace Amine Receptor 4 (TRAR4) Gene on Chromosome 6q23.2 Are Associated with Susceptibility to Schizophrenia. American Journal of Human Genetics, 2004, 75, 624-638.	6.2	101
65	Analysis of ten candidate genes in autism by association and linkage. American Journal of Medical Genetics Part A, 2002, 114, 125-128.	2.4	20
66	Association between tumor necrosis factor receptor II and familial, but not sporadic, rheumatoid arthritis: Evidence for genetic heterogeneity. Arthritis and Rheumatism, 2002, 46, 2039-2044.	6.7	62
67	Evidence of a Cohort Effect for Age at Onset of Schizophrenia. American Journal of Psychiatry, 2001, 158, 489-492.	7.2	29
68	Different familial transmission patterns in bipolar I disorder with onset before and after age 25. American Journal of Medical Genetics Part A, 2001, 105, 765-773.	2.4	74
69	A Third Major Locus for Autosomal Dominant Hypercholesterolemia Maps to 1p34.1-p32. American Journal of Human Genetics, 1999, 64, 1378-1387.	6.2	154
70	Early-Onset Autosomal Dominant Alzheimer Disease: Prevalence, Genetic Heterogeneity, and Mutation Spectrum. American Journal of Human Genetics, 1999, 65, 664-670.	6.2	696
71	The Ile93Met mutation in the ubiquitin carboxy-terminal-hydrolase-L1 gene is not observed in European cases with familial Parkinson's disease. Neuroscience Letters, 1999, 270, 1-4.	2.1	75
72	Apolipoprotein E Ïµ4 Allele and Familial Aggregation of Alzheimer Disease. Archives of Neurology, 1998, 55, 810.	4.5	65

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73	Major psychoses and a potential vulnerability locus on 6p24-p22 in the Eastern QuÃ©bec population. Schizophrenia Research, 1997, 24, 52.	2.0	1
74	Childhood-early adolescence onset and adult onset schizophrenia: Heterogeneity at the dopamine D3 receptor gene. Schizophrenia Research, 1997, 24, 92.	2.0	0
75	Suggestive Evidence for a Schizophrenia Susceptibility Locus on Chromosome 6q and a Confirmation in an Independent Series of Pedigrees. Genomics, 1997, 43, 1-8.	2.9	229
76	Serotonin transporter gene polymorphism and schizophrenia: An association study. Biological Psychiatry, 1997, 42, 634-636.	1.3	29
77	Childhood/early adolescence-onsetand adult-onset schizophrenia. British Journal of Psychiatry, 1997, 170, 27-30.	2.8	16
78	A novel presenilin 1 mutation resulting in familial AlzheimerÊ¼s disease with an onset age of 29 years. NeuroReport, 1996, 7, 1582-1584.	1.2	75
79	Long-Term Stability of Diagnosis and Symptom Dimensions in a Systematic Sample of Patients with Onset of Schizophrenia in Childhood and Early Adolescence. I: Nosology, Sex and Age of Onset. British Journal of Psychiatry, 1996, 169, 361-370.	2.8	68
80	Long-Term Stability of Diagnosis and Symptom Dimensions in a Systematic Sample of Patients with Onset of Schizophrenia in Childhood and Early Adolescence. II: Positive/Negative Distinction and Childhood Predictors of Adult Outcome. British Journal of Psychiatry, 1996, 169, 371-378.	2.8	77
81	Allelic association at the D14S43 locus in early onset Alzheimer's disease. American Journal of Medical Genetics Part A, 1995, 60, 91-93.	2.4	4
82	Evidence for apolipoprotein E Îµ4 association in early-onset Alzheimer's patients with late-onset relatives. American Journal of Medical Genetics Part A, 1995, 60, 550-553.	2.4	28
83	Mutations of the presenilin I gene in families with early-onset Alzheimer's disease. Human Molecular Genetics, 1995, 4, 2373-2377.	2.9	268
84	Further evidence for anticipation in schizophrenia. Psychiatry Research, 1995, 59, 25-33.	3.3	48
85	No evidence for linkage or association between the dopamine transporter gene and schizophrenia in a French population. Psychiatry Research, 1995, 59, 1-6.	3.3	27
86	Follow-up of a report of a potential linkage for schizophrenia on chromosome 22q12-q13.1: Part 2. American Journal of Medical Genetics Part A, 1994, 54, 44-50.	2.4	145
87	Genetic study of dopamine D1, D2, and D4 receptors in schizophrenia. Psychiatry Research, 1994, 51, 215-230.	3.3	53
88	Pseudoautosomal region in schizophrenia: Linkage analysis of seven loci by sib-pair and lod-score methods. Psychiatry Research, 1994, 52, 135-147.	3.3	13
89	Linkage detection by the Affected-Pedigree-Member method: What is really tested?. Genetic Epidemiology, 1993, 10, 389-394.	1.3	44
90	Is a single mutation at the same locus responsible for all affected cases in a large Alzheimer pedigree (FAD4)?. Genetic Epidemiology, 1993, 10, 431-435.	1.3	0

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91	Genetic linkage mapping of the m4 human muscarinic receptor (CHRM4). Genomics, 1992, 13, 239-240.	2.9	7
92	Genetic mapping of the Gs-Î± subunit gene (GNAS1) to the distal long arm of chromosome 20 using a polymorphism detected by denaturing gradient gel electrophoresis. Genomics, 1991, 9, 782-783.	2.9	74
93	Genetic mapping of common diseases: the challenges of manic-depressive illness and schizophrenia. Trends in Genetics, 1990, 6, 282-286.	6.7	24
94	Power of the linkage test for a heterogeneous disorder due to two independent inherited causes: A simulation study. Genetic Epidemiology, 1990, 7, 219-230.	1.3	31
95	Performance of linkage analysis under misclassification error when the genetic model is unknown. Genetic Epidemiology, 1989, 6, 253-258.	1.3	40
96	Genetic analysis of human breast cancer: Implications for family study designs. Genetic Epidemiology, 1988, 5, 225-233.	1.3	14