

Antonio F Pardias

List of Publications by Citations

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

75
papers

2,819
citations

22
h-index

52
g-index

92
ext. papers

4,831
ext. citations

9.9
avg, IF

4.46
L-index

#	Paper	IF	Citations
75	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. <i>Nature Genetics</i> , 2018 , 50, 381-389	36.3	787
74	Genetic identification of brain cell types underlying schizophrenia. <i>Nature Genetics</i> , 2018 , 50, 825-833	36.3	295
73	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018 , 362,	33.3	277
72	Identification of genetic variants associated with Huntington's disease progression: a genome-wide association study. <i>Lancet Neurology</i> , <i>The</i> , 2017 , 16, 701-711	24.1	161
71	Gene expression imputation across multiple brain regions provides insights into schizophrenia risk. <i>Nature Genetics</i> , 2019 , 51, 659-674	36.3	99
70	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021 , 53, 817-829	36.3	83
69	Analysis of Intellectual Disability Copy Number Variants for Association With Schizophrenia. <i>JAMA Psychiatry</i> , 2016 , 73, 963-969	14.5	75
68	High level of mislabeling in Spanish and Greek hake markets suggests the fraudulent introduction of African species. <i>Journal of Agricultural and Food Chemistry</i> , 2011 , 59, 475-80	5.7	72
67	Expression quantitative trait loci in the developing human brain and their enrichment in neuropsychiatric disorders. <i>Genome Biology</i> , 2018 , 19, 194	18.3	69
66	Association Between Schizophrenia-Related Polygenic Liability and the Occurrence and Level of Mood-Incongruent Psychotic Symptoms in Bipolar Disorder. <i>JAMA Psychiatry</i> , 2018 , 75, 28-35	14.5	66
65	Association of Genetic Liability to Psychotic Experiences With Neuropsychotic Disorders and Traits. <i>JAMA Psychiatry</i> , 2019 , 76, 1256-1265	14.5	58
64	Medical consequences of pathogenic CNVs in adults: analysis of the UK Biobank. <i>Journal of Medical Genetics</i> , 2019 , 56, 131-138	5.8	56
63	Genetic influences on eight psychiatric disorders based on family data of 4 408 646 full and half-siblings, and genetic data of 333 748 cases and controls. <i>Psychological Medicine</i> , 2019 , 49, 1166-1173	6.9	52
62	De novo mutations identified by exome sequencing implicate rare missense variants in SLC6A1 in schizophrenia. <i>Nature Neuroscience</i> , 2020 , 23, 179-184	25.5	47
61	Genome-wide common and rare variant analysis provides novel insights into clozapine-associated neutropenia. <i>Molecular Psychiatry</i> , 2017 , 22, 1502-1508	15.1	45
60	Genetic contributors to risk of schizophrenia in the presence of a 22q11.2 deletion. <i>Molecular Psychiatry</i> , 2021 , 26, 4496-4510	15.1	39
59	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2020 , 46, 336-344	1.3	38

58	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , 2022 ,	50.4	35
57	Socio-cultural factors in dental diseases in the Medieval and early Modern Age of northern Spain. <i>HOMO- Journal of Comparative Human Biology</i> , 2012 , 63, 21-42	0.5	29
56	A transcriptome-wide association study implicates specific pre- and post-synaptic abnormalities in schizophrenia. <i>Human Molecular Genetics</i> , 2020 , 29, 159-167	5.6	27
55	Novel Insight Into the Etiology of Autism Spectrum Disorder Gained by Integrating Expression Data With Genome-wide Association Statistics. <i>Biological Psychiatry</i> , 2019 , 86, 265-273	7.9	26
54	Recent Demographic History Inferred by High-Resolution Analysis of Linkage Disequilibrium. <i>Molecular Biology and Evolution</i> , 2020 , 37, 3642-3653	8.3	25
53	Pharmacogenomic Variants and Drug Interactions Identified Through the Genetic Analysis of Clozapine Metabolism. <i>American Journal of Psychiatry</i> , 2019 , 176, 477-486	11.9	21
52	Targeted Sequencing of 10,198 Samples Confirms Abnormalities in Neuronal Activity and Implicates Voltage-Gated Sodium Channels in Schizophrenia Pathogenesis. <i>Biological Psychiatry</i> , 2019 , 85, 554-562	7.9	21
51	Common schizophrenia alleles are enriched in mutation-intolerant genes and maintained by background selection		20
50	Genome-wide association studies in schizophrenia: Recent advances, challenges and future perspective. <i>Schizophrenia Research</i> , 2020 , 217, 4-12	3.6	20
49	Conditional GWAS analysis to identify disorder-specific SNPs for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2070-2081	15.1	19
48	Clinical indicators of treatment-resistant psychosis. <i>British Journal of Psychiatry</i> , 2020 , 216, 259-266	5.4	19
47	A genome-wide association study in individuals of African ancestry reveals the importance of the Duffy-null genotype in the assessment of clozapine-related neutropenia. <i>Molecular Psychiatry</i> , 2019 , 24, 328-337	15.1	18
46	Role of upwelling on larval dispersal and productivity of gooseneck barnacle populations in the Cantabrian Sea: management implications. <i>PLoS ONE</i> , 2013 , 8, e78482	3.7	13
45	Evidence of trepanations in a medieval population (13th-14th century) of northern Spain (Gormaz, Soria). <i>Anthropological Science</i> , 2011 , 119, 247-257	1.3	12
44	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology		11
43	Climate change and oceanic barriers: genetic differentiation in <i>Pomatomus saltatrix</i> (Pisces: Pomatomidae) in the North Atlantic Ocean and the Mediterranean Sea. <i>Journal of Fish Biology</i> , 2010 , 77, 1993-8	1.9	9
42	The effect of a genetic variant at the schizophrenia associated AS3MT/BORCS7 locus on striatal dopamine function: A PET imaging study. <i>Psychiatry Research - Neuroimaging</i> , 2019 , 291, 34-41	2.9	8
41	Paleoclimate Shaped Bluefish Structure in the Northern Hemisphere. <i>Fisheries</i> , 2014 , 39, 578-586	1.1	8

40	Assessing the genetic influence of ancient sociopolitical structure: micro-differentiation patterns in the population of Asturias (Northern Spain). <i>PLoS ONE</i> , 2012 , 7, e50206	3.7	8
39	Satisfaction with higher education of Spanish graduates in the pre-Bologna era: a mirror of employment conditions?. <i>Journal of Further and Higher Education</i> , 2012 , 36, 519-534	1.5	7
38	Introducing human population biology through an easy laboratory exercise on mitochondrial DNA. <i>Biochemistry and Molecular Biology Education</i> , 2010 , 38, 110-5	1.3	7
37	Genetic identification Of brain cell types underlying schizophrenia		7
36	Genetic association of FMRP targets with psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2977-2990	5.1	7
35	Gene expression imputation across multiple brain regions reveals schizophrenia risk throughout development		6
34	Conditional GWAS analysis identifies putative disorder-specific SNPs for psychiatric disorders		6
33	Convergent Evidence That ZNF804A Is a Regulator of Pre-messenger RNA Processing and Gene Expression. <i>Schizophrenia Bulletin</i> , 2019 , 45, 1267-1278	1.3	6
32	The Relationship Between Common Variant Schizophrenia Liability and Number of Offspring in the UK Biobank. <i>American Journal of Psychiatry</i> , 2019 , 176, 661-666	11.9	6
31	The Relationship Between Common Variant Schizophrenia Liability and Number of Offspring in the UK Biobank: Response to Lawn et al. <i>American Journal of Psychiatry</i> , 2019 , 176, 574-575	11.9	5
30	Polygenic risk for schizophrenia and subcortical brain anatomy in the UK Biobank cohort. <i>Translational Psychiatry</i> , 2020 , 10, 309	8.6	5
29	Clozapine Metabolism is Associated With Absolute Neutrophil Count in Individuals With Treatment-Resistant Schizophrenia. <i>Frontiers in Pharmacology</i> , 2021 , 12, 658734	5.6	5
28	The effect of the DISC1 Ser704Cys polymorphism on striatal dopamine synthesis capacity: an [18F]-DOPA PET study. <i>Human Molecular Genetics</i> , 2018 , 27, 3498-3506	5.6	5
27	Rare Copy Number Variants Are Associated With Poorer Cognition in Schizophrenia. <i>Biological Psychiatry</i> , 2021 , 90, 28-34	7.9	5
26	Over the sands and far away: interpreting an Iberian mitochondrial lineage with ancient Western African origins. <i>American Journal of Human Biology</i> , 2014 , 26, 777-83	2.7	4
25	Treponemal disease in the Old World? Integrated palaeopathological assessment of a 9th-11th century skeleton from north-central Spain. <i>Anthropological Science</i> , 2017 , 125, 101-114	1.3	4
24	Mitochondrial diversity patterns and the Magdalenian resettlement of Europe: new insights from the edge of the Franco-Cantabrian refuge. <i>Journal of Human Genetics</i> , 2012 , 57, 717-26	4.3	4
23	A Transcriptome Wide Association Study implicates specific pre- and post-synaptic abnormalities in Schizophrenia		

22	Associations Between Schizophrenia Polygenic Liability, Symptom Dimensions, and Cognitive Ability in Schizophrenia. <i>JAMA Psychiatry</i> , 2021 , 78, 1143-1151	14.5	4
21	Transcriptional programs regulating neuronal differentiation are disrupted in DLG2 knockout human embryonic stem cells and enriched for schizophrenia and related disorders risk variants.. <i>Nature Communications</i> , 2022 , 13, 27	17.4	3
20	HLA-DQB1 6672G>C (rs113332494) is associated with clozapine-induced neutropenia and agranulocytosis in individuals of European ancestry. <i>Translational Psychiatry</i> , 2021 , 11, 214	8.6	3
19	Developmental Profile of Psychiatric Risk Associated With Voltage-Gated Cation Channel Activity. <i>Biological Psychiatry</i> , 2021 , 90, 399-408	7.9	3
18	Medical diagnostic methods applied to a medieval female with vitamin D deficiency from the north of Spain. <i>International Journal of Paleopathology</i> , 2018 , 22, 109-120	1.5	2
17	Phenotyping the ancient world: The physical appearance and ancestry of very degraded samples from a chalcolithic human remains. <i>Forensic Science International: Genetics Supplement Series</i> , 2017 , 6, e484-e486	0.5	2
16	Association between schizophrenia and both loss of function and missense mutations in paralog conserved sites of voltage-gated sodium channels		2
15	Analyses of rare and common alleles in parent-proband trios implicate rare missense variants in SLC6A1 in schizophrenia and confirm the involvement of loss of function intolerant and neurodevelopmental disorder genes		2
14	The quest for the soldier's rest: combining anthropological and archaeochemical approaches to study social and occupational diversity in the medieval graveyard of San Andr� de Arroyo (Palencia, Spain). <i>Anthropological Science</i> , 2016 , 124, 169-184	1.3	2
13	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021 , 16, e0248254	3.7	2
12	T66. PREDICTING TREATMENT RESISTANT SCHIZOPHRENIA AT FIRST-EPISEDE OF PSYCHOSIS. <i>Schizophrenia Bulletin</i> , 2019 , 45, S229-S230	1.3	1
11	DLG2 knockout reveals neurogenic transcriptional programs underlying neuropsychiatric disorders and cognition		1
10	Pharmacogenomics: A road ahead for precision medicine in psychiatry. <i>Neuron</i> , 2021 ,	13.9	1
9	The Duffy-null genotype and risk of infection. <i>Human Molecular Genetics</i> , 2020 , 29, 3341-3349	5.6	1
8	Impact of schizophrenia genetic liability on the association between schizophrenia and physical illness: data-linkage study. <i>BJPsych Open</i> , 2020 , 6, e139	5	1
7	Genome-wide association analyses of symptom severity among clozapine-treated patients with schizophrenia spectrum disorders.. <i>Translational Psychiatry</i> , 2022 , 12, 145	8.6	1
6	Post-partum psychosis and its association with bipolar disorder in the UK: a case-control study using polygenic risk scores. <i>Lancet Psychiatry</i> , 2021 , 8, 1045-1052	23.3	0
5	Genomic treatment response prediction in schizophrenia 2020 , 413-422		0

4	Assessing individual and population variability in degenerative joint disease prevalence using generalized linear mixed models. <i>American Journal of Physical Anthropology</i> , 2021 , 175, 611-625	2.5	0
3	Evaluation of large-scale genetic structure in complex demographic and historical scenarios: the mitochondrial DNA and Y-chromosome pools of the Iberian Atlantic façade. <i>American Journal of Physical Anthropology</i> , 2014 , 153, 617-26	2.5	
2	Type I Monteggia fracture-dislocation in a monk from a 17th-18th century necropolis of Valladolid (Spain). <i>Anthropological Science</i> , 2011 , 119, 39-47	1.3	
1	Summaries of plenary, symposia, and oral sessions at the XXII World Congress of Psychiatric Genetics, Copenhagen, Denmark, 12-16 October 2014. <i>Psychiatric Genetics</i> , 2016 , 26, 1-47	2.9	