

Antonio F Pardias

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/883140/antonio-f-pardinas-publications-by-year.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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	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
74	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
73	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , 2022 ,	50.4	35
72	Pharmacogenomics: A road ahead for precision medicine in psychiatry. <i>Neuron</i> , 2021 ,	13.9	1
71	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
70	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
69	Conditional GWAS analysis to identify disorder-specific SNPs for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2070-2081	15.1	19
68	Genetic association of FMRP targets with psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2977-2990	15.1	7
67	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
66	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
65	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
64	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
63	Rare Copy Number Variants Are Associated With Poorer Cognition in Schizophrenia. <i>Biological Psychiatry</i> , 2021 , 90, 28-34	7.9	5
62	Associations Between Schizophrenia Polygenic Liability, Symptom Dimensions, and Cognitive Ability in Schizophrenia. <i>JAMA Psychiatry</i> , 2021 , 78, 1143-1151	14.5	4
61	Developmental Profile of Psychiatric Risk Associated With Voltage-Gated Cation Channel Activity. <i>Biological Psychiatry</i> , 2021 , 90, 399-408	7.9	3
60	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021 , 16, e0248254	3.7	2
59	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2020 , 46, 336-344	1.3	38

58	Recent Demographic History Inferred by High-Resolution Analysis of Linkage Disequilibrium. <i>Molecular Biology and Evolution</i> , 2020 , 37, 3642-3653	8.3	25
57	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
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	Genomic treatment response prediction in schizophrenia 2020 , 413-422		0
54	Genome-wide association studies in schizophrenia: Recent advances, challenges and future perspective. <i>Schizophrenia Research</i> , 2020 , 217, 4-12	3.6	20
53	The Duffy-null genotype and risk of infection. <i>Human Molecular Genetics</i> , 2020 , 29, 3341-3349	5.6	1
52	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
51	Polygenic risk for schizophrenia and subcortical brain anatomy in the UK Biobank cohort. <i>Translational Psychiatry</i> , 2020 , 10, 309	8.6	5
50	Clinical indicators of treatment-resistant psychosis. <i>British Journal of Psychiatry</i> , 2020 , 216, 259-266	5.4	19
49	T66. PREDICTING TREATMENT RESISTANT SCHIZOPHRENIA AT FIRST-EPISODE OF PSYCHOSIS. <i>Schizophrenia Bulletin</i> , 2019 , 45, S229-S230	1.3	1
48	Association of Genetic Liability to Psychotic Experiences With Neuropsychotic Disorders and Traits. <i>JAMA Psychiatry</i> , 2019 , 76, 1256-1265	14.5	58
47	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
46	Gene expression imputation across multiple brain regions provides insights into schizophrenia risk. <i>Nature Genetics</i> , 2019 , 51, 659-674	36.3	99
45	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
44	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
43	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
42	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
41	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

40	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
39	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
38	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
37	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
36	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
35	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
34	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
33	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
32	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018 , 362,	33.3	277
31	Genetic identification of brain cell types underlying schizophrenia. <i>Nature Genetics</i> , 2018 , 50, 825-833	36.3	295
30	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
29	Identification of genetic variants associated with Huntington's disease progression: a genome-wide association study. <i>Lancet Neurology</i> , 2017 , 16, 701-711	24.1	161
28	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
27	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
26	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
25	Analysis of Intellectual Disability Copy Number Variants for Association With Schizophrenia. <i>JAMA Psychiatry</i> , 2016 , 73, 963-969	14.5	75
24	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
23	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	

22	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	
21	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
20	Paleoclimate Shaped Bluefish Structure in the Northern Hemisphere. <i>Fisheries</i> , 2014 , 39, 578-586	1.1	8
19	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
18	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
17	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
16	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
15	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
14	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
13	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
12	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	
11	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
10	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
9	DLG2 knockout reveals neurogenic transcriptional programs underlying neuropsychiatric disorders and cognition		1
8	Common schizophrenia alleles are enriched in mutation-intolerant genes and maintained by background selection		20
7	Genetic identification Of brain cell types underlying schizophrenia		7
6	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology		11
5	Gene expression imputation across multiple brain regions reveals schizophrenia risk throughout development	6	

- 4 Association between schizophrenia and both loss of function and missense mutations in paralog conserved sites of voltage-gated sodium channels 2
- 3 A Transcriptome Wide Association Study implicates specific pre- and post-synaptic abnormalities in Schizophrenia
- 2 Conditional GWAS analysis identifies putative disorder-specific SNPs for psychiatric disorders 6
- 1 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