

Sara BandrÃ©s Ciga

List of Publications by Year in descending order

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Version: 2024-02-01

78
papers

4,295
citations

236925

25
h-index

149698

56
g-index

115
all docs

115
docs citations

115
times ranked

5244
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of novel risk loci, causal insights, and heritable risk for Parkinson's disease: a meta-analysis of genome-wide association studies. <i>Lancet Neurology</i> , The, 2019, 18, 1091-1102.	10.2	1,414
2	Parkinson's disease age at onset genome-wide association study: Defining heritability, genetic loci, and ß-synuclein mechanisms. <i>Movement Disorders</i> , 2019, 34, 866-875.	3.9	258
3	Genetics of Parkinson's disease: An introspection of its journey towards precision medicine. <i>Neurobiology of Disease</i> , 2020, 137, 104782.	4.4	241
4	Genome sequencing analysis identifies new loci associated with Lewy body dementia and provides insights into its genetic architecture. <i>Nature Genetics</i> , 2021, 53, 294-303.	21.4	198
5	Genetic risk factors in Parkinson's disease. <i>Cell and Tissue Research</i> , 2018, 373, 9-20.	2.9	159
6	Genetic modifiers of risk and age at onset in GBA associated Parkinson's disease and Lewy body dementia. <i>Brain</i> , 2020, 143, 234-248.	7.6	149
7	Shared polygenic risk and causal inferences in amyotrophic lateral sclerosis. <i>Annals of Neurology</i> , 2019, 85, 470-481.	5.3	118
8	A Missense Mutation in KCTD17 Causes Autosomal Dominant Myoclonus-Dystonia. <i>American Journal of Human Genetics</i> , 2015, 96, 938-947.	6.2	109
9	Type 2 Diabetes as a Determinant of Parkinson's Disease Risk and Progression. <i>Movement Disorders</i> , 2021, 36, 1420-1429.	3.9	108
10	The role of monogenic genes in idiopathic Parkinson's disease. <i>Neurobiology of Disease</i> , 2019, 124, 230-239.	4.4	97
11	Mitochondria function associated genes contribute to Parkinson's Disease risk and later age at onset. <i>Npj Parkinson's Disease</i> , 2019, 5, 8.	5.3	95
12	The Parkinson's Disease ^{Genome-Wide} Association Study Locus Browser. <i>Movement Disorders</i> , 2020, 35, 2056-2067.	3.9	68
13	Large-scale pathway specific polygenic risk and transcriptomic community network analysis identifies novel functional pathways in Parkinson disease. <i>Acta Neuropathologica</i> , 2020, 140, 341-358.	7.7	68
14	The endocytic membrane trafficking pathway plays a major role in the risk of Parkinson's disease. <i>Movement Disorders</i> , 2019, 34, 460-468.	3.9	66
15	Parkinson's disease determinants, prediction and gene-environment interactions in the UK Biobank. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2020, 91, 1046-1054.	1.9	59
16	Genetic analysis of amyotrophic lateral sclerosis identifies contributing pathways and cell types. <i>Science Advances</i> , 2021, 7, .	10.3	59
17	Penetrance of Parkinson's Disease in <i>LRRK2</i> p.G2019S Carriers Is Modified by a Polygenic Risk Score. <i>Movement Disorders</i> , 2020, 35, 774-780.	3.9	57
18	The Parkinson's Disease Mendelian Randomization Research Portal. <i>Movement Disorders</i> , 2019, 34, 1864-1872.	3.9	50

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19	The Genetic Architecture of Parkinson Disease in Spain: Characterizing Population-Specific Risk, Differential Haplotype Structures, and Providing Etiologic Insight. <i>Movement Disorders</i> , 2019, 34, 1851-1863.	3.9	47
20	Finding genetically-supported drug targets for Parkinson's disease using Mendelian randomization of the druggable genome. <i>Nature Communications</i> , 2021, 12, 7342.	12.8	44
21	Lower Lymphocyte Count is Associated With Increased Risk of Parkinson's Disease. <i>Annals of Neurology</i> , 2021, 89, 803-812.	5.3	38
22	Proteotoxicity and Neurodegenerative Diseases. <i>International Journal of Molecular Sciences</i> , 2020, 21, 5646.	4.1	37
23	Genome-wide assessment of Parkinson's disease in a Southern Spanish population. <i>Neurobiology of Aging</i> , 2016, 45, 213.e3-213.e9.	3.1	35
24	Heritability Enrichment Implicates Microglia in Parkinson's Disease Pathogenesis. <i>Annals of Neurology</i> , 2021, 89, 942-951.	5.3	35
25	Comprehensive assessment of PINK1 variants in Parkinson's disease. <i>Neurobiology of Aging</i> , 2020, 91, 168.e1-168.e5.	3.1	32
26	Investigation of Autosomal Genetic Sex Differences in Parkinson's Disease. <i>Annals of Neurology</i> , 2021, 90, 35-42.	5.3	29
27	The <i>CACNA1B</i> R1389H variant is not associated with myoclonus-dystonia in a large European multicentric cohort. <i>Human Molecular Genetics</i> , 2015, 24, 5326-5329.	2.9	28
28	Heterozygous <i>PRKN</i> mutations are common but do not increase the risk of Parkinson's disease. <i>Brain</i> , 2022, 145, 2077-2091.	7.6	26
29	Analysis of the genetic variability in Parkinson's disease from Southern Spain. <i>Neurobiology of Aging</i> , 2016, 37, 210.e1-210.e5.	3.1	23
30	SNCA and mTOR Pathway Single Nucleotide Polymorphisms Interact to Modulate the Age at Onset of Parkinson's Disease. <i>Movement Disorders</i> , 2019, 34, 1333-1344.	3.9	21
31	Mendelian Randomization: A Journey From Obscurity to Center Stage With a Few Potholes Along the Way. <i>JAMA Neurology</i> , 2020, 77, 7.	9.0	21
32	An Examination of the Mechanisms Involved in Secondary Clinical Failure to Adalimumab or Etanercept in Inflammatory Arthropathies. <i>Journal of Clinical Rheumatology</i> , 2015, 21, 115-119.	0.9	19
33	ARSA variants in α -synucleinopathies. <i>Brain</i> , 2019, 142, e70-e70.	7.6	17
34	Tumor Necrosis Factor Inhibition and Parkinson Disease. <i>Neurology</i> , 2021, 96, e1672-e1679.	1.1	17
35	Advances in Proteomic and Metabolomic Profiling of Neurodegenerative Diseases. <i>Frontiers in Neurology</i> , 2021, 12, 792227.	2.4	17
36	Genetic Risk Profiling in Parkinson's Disease and Utilizing Genetics to Gain Insight into Disease-Related Biological Pathways. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7332.	4.1	16

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37	Analysis of DNMT3 and VAMP4 as genetic modifiers of LRRK2 Parkinson's disease. <i>Neurobiology of Aging</i> , 2021, 97, 148.e17-148.e24.	3.1	16
38	Unhealthy Behaviours and Risk of Parkinson's Disease: A Mendelian Randomisation Study. <i>Journal of Parkinson's Disease</i> , 2021, 11, 1981-1993.	2.8	16
39	LRP10 in α -synucleinopathies. <i>Lancet Neurology</i> , The, 2018, 17, 1032.	10.2	15
40	ATP10B and the risk for Parkinson's disease. <i>Acta Neuropathologica</i> , 2020, 140, 401-402.	7.7	14
41	Genetic Stratification of Age-Dependent Parkinson's Disease Risk by Polygenic Hazard Score. <i>Movement Disorders</i> , 2022, 37, 62-69.	3.9	13
42	Alpha-synuclein triggers T _H cell response. Is Parkinson's disease an autoimmune disorder?. <i>Movement Disorders</i> , 2017, 32, 1327-1327.	3.9	12
43	Lack of evidence for genetic association of saposins A, B, C and D with Parkinson's disease. <i>Brain</i> , 2020, 143, e72-e72.	7.6	11
44	Evaluating Lipid-Lowering Drug Targets for Parkinson's Disease Prevention with Mendelian Randomization. <i>Annals of Neurology</i> , 2020, 88, 1043-1047.	5.3	11
45	Exome-wide rare variant analysis in familial essential tremor. <i>Parkinsonism and Related Disorders</i> , 2021, 82, 109-116.	2.2	11
46	The Parkinson's Disease <sc>DNA</sc> Variant Browser. <i>Movement Disorders</i> , 2021, 36, 1250-1258.	3.9	11
47	LRRK2 p.M1646T is associated with glucocerebrosidase activity and with Parkinson's disease. <i>Neurobiology of Aging</i> , 2021, 103, 142.e1-142.e5.	3.1	11
48	Genetic and Transcriptomic Biomarkers in Neurodegenerative Diseases: Current Situation and the Road Ahead. <i>Cells</i> , 2021, 10, 1030.	4.1	11
49	<sc> α -Synuclein</sc> Deposition in Sympathetic Nerve Fibers in Genetic Forms of Parkinson's Disease. <i>Movement Disorders</i> , 2021, 36, 2346-2357.	3.9	11
50	Structural genomic variations and Parkinson's disease. <i>Minerva Medica</i> , 2017, 108, 438-447.	0.9	11
51	Genome-wide estimates of heritability and genetic correlations in essential tremor. <i>Parkinsonism and Related Disorders</i> , 2019, 64, 262-267.	2.2	10
52	No genetic evidence for involvement of alcohol dehydrogenase genes in risk for Parkinson's disease. <i>Neurobiology of Aging</i> , 2020, 87, 140.e19-140.e22.	3.1	10
53	Polygenic Resilience Modulates the Penetrance of Parkinson Disease Genetic Risk Factors. <i>Annals of Neurology</i> , 2022, 92, 270-278.	5.3	10
54	An integrated genomic approach to dissect the genetic landscape regulating the cell-to-cell transfer of α -synuclein. <i>Cell Reports</i> , 2021, 35, 109189.	6.4	8

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55	Predicting progression in patients with Parkinson's disease. <i>Lancet Neurology</i> , The, 2017, 16, 860-862.	10.2	7
56	Lack of evidence for association of UQCRC1 with Parkinson's disease in Europeans. <i>Neurobiology of Aging</i> , 2021, 101, 297.e1-297.e4.	3.1	7
57	Mapping the Diverse and Inclusive Future of Parkinson's Disease Genetics and Its Widespread Impact. <i>Genes</i> , 2021, 12, 1681.	2.4	7
58	Advancing Personalized Medicine in Common Forms of Parkinson's Disease through Genetics: Current Therapeutics and the Future of Individualized Management. <i>Journal of Personalized Medicine</i> , 2021, 11, 169.	2.5	6
59	Immunogenetic Determinants of Parkinson's Disease Etiology. <i>Journal of Parkinson's Disease</i> , 2022, 12, S13-S27.	2.8	6
60	<i>MIDN</i> locus structural variants and Parkinson's Disease risk. <i>Annals of Clinical and Translational Neurology</i> , 2020, 7, 602-603.	3.7	5
61	A population scale analysis of rare SNCA variation in the UK Biobank. <i>Neurobiology of Disease</i> , 2021, 148, 105182.	4.4	5
62	Insights on Genetic and Environmental Factors in Parkinson's Disease from a Regional Swedish Case-Control Cohort. <i>Journal of Parkinson's Disease</i> , 2022, 12, 153-171.	2.8	5
63	Assessment of Genetic Association Between Parkinson Disease and Bipolar Disorder. <i>JAMA Neurology</i> , 2020, 77, 1034.	9.0	4
64	Assessment of LIN28A variants in Parkinson's disease in large European cohorts. <i>Neurobiology of Aging</i> , 2021, 100, 118.e1-118.e3.	3.1	4
65	The role of RHOT1 and RHOT2 genetic variation on Parkinson disease risk and onset. <i>Neurobiology of Aging</i> , 2021, 97, 144.e1-144.e3.	3.1	3
66	Replication assessment of NUS1 variants in Parkinson's disease. <i>Neurobiology of Aging</i> , 2021, 101, 300.e1-300.e3.	3.1	3
67	No Evidence for a Causal Relationship Between Cancers and Parkinson's Disease. <i>Journal of Parkinson's Disease</i> , 2021, 11, 801-809.	2.8	3
68	Lack of Causal Effects or Genetic Correlation between Restless Legs Syndrome and Parkinson's Disease. <i>Movement Disorders</i> , 2021, 36, 1967-1972.	3.9	3
69	The Genetic Landscape of Parkinsonism-Related Dystonias and Atypical Parkinsonism-Related Syndromes. <i>International Journal of Molecular Sciences</i> , 2021, 22, 8100.	4.1	3
70	Black and African American Connections to Parkinson's Disease Study: Addressing Missing Diversity in Parkinson's Disease Genetics. <i>Movement Disorders</i> , 2022, 37, 1559-1561.	3.9	3
71	The appendix and the risk of Parkinson's disease: Appended notes on correlation and causation. <i>Movement Disorders</i> , 2019, 34, 199-199.	3.9	2
72	Unraveling the genetic complexity of Alzheimer disease with Mendelian Randomization. <i>Neurology: Genetics</i> , 2019, 5, e313.	1.9	1

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73	Analysis of PTRHD1 common and rare variants in European patients with Parkinson's disease. <i>Neurobiology of Aging</i> , 2021, 107, 178-180.	3.1	1
74	Common and rare variants in HFE are not associated with Parkinson's disease in Europeans. <i>Neurobiology of Aging</i> , 2021, 107, 174-177.	3.1	1
75	Assessment of ANG variants in Parkinson's disease. <i>Neurobiology of Aging</i> , 2021, 104, 111.e1-111.e4.	3.1	1
76	Identifying Genetic Markers Associated with the Progression of Cognitive Decline in Parkinson's Disease: A Call Out for Replication. <i>Movement Disorders</i> , 2021, 36, 2506-2507.	3.9	1
77	RIC3 variants are not associated with Parkinson's disease in large European, Latin American, or East Asian cohorts. <i>Neurobiology of Aging</i> , 2022, 109, 264-268.	3.1	0
78	17q21.31 Sub-Haplotypes Underlying H1-Associated Risk for Parkinson's Disease and Progressive Supranuclear Palsy Converge on Altered Glial Regulation. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0