

Rub n Fern ndez-Santiago

List of Publications by Year in descending order

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Version: 2024-02-01

62
papers

3,772
citations

186265

28
h-index

149698

56
g-index

63
all docs

63
docs citations

63
times ranked

5775
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	Angiogenin variants in Parkinson disease and amyotrophic lateral sclerosis. <i>Annals of Neurology</i> , 2011, 70, 964-973.	5.3	168
3	Penetrance estimate of <i>LRRK2</i> p.G2019S mutation in individuals of non-Ashkenazi Jewish ancestry. <i>Movement Disorders</i> , 2017, 32, 1432-1438.	3.9	126
4	Identification of blood serum microRNAs associated with idiopathic and <i>LRRK2</i> Parkinson's disease. <i>Journal of Neuroscience Research</i> , 2014, 92, 1071-1077.	2.9	122
5	Aberrant epigenome in <i>iPSC</i> -derived dopaminergic neurons from Parkinson's disease patients. <i>EMBO Molecular Medicine</i> , 2015, 7, 1529-1546.	6.9	117
6	Urinary <i>LRRK2</i> phosphorylation predicts parkinsonian phenotypes in G2019S <i>LRRK2</i> carriers. <i>Neurology</i> , 2016, 86, 994-999.	1.1	114
7	Motor and nonmotor heterogeneity of <i>LRRK2</i> -related and idiopathic Parkinson's disease. <i>Movement Disorders</i> , 2016, 31, 1192-1202.	3.9	102
8	Advances in Parkinson's Disease: 200 Years Later. <i>Frontiers in Neuroanatomy</i> , 2018, 12, 113.	1.7	102
9	Identification of Candidate Parkinson Disease Genes by Integrating Genome-Wide Association Study, Expression, and Epigenetic Data Sets. <i>JAMA Neurology</i> , 2021, 78, 464.	9.0	95
10	Meta-analysis of vascular endothelial growth factor variations in amyotrophic lateral sclerosis: increased susceptibility in male carriers of the -2578AA genotype. <i>Journal of Medical Genetics</i> , 2009, 46, 840-846.	3.2	70
11	Sleep Disorders in Parkinsonian and Nonparkinsonian <i>LRRK2</i> Mutation Carriers. <i>PLoS ONE</i> , 2015, 10, e0132368.	2.5	67
12	Identification of novel Angiogenin (ANG) gene missense variants in German patients with amyotrophic lateral sclerosis. <i>Journal of Neurology</i> , 2009, 256, 1337-1342.	3.6	60
13	Nonsteroidal Anti-inflammatory Use and <i>LRRK2</i> Parkinson's Disease Penetrance. <i>Movement Disorders</i> , 2020, 35, 1755-1764.	3.9	57
14	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
15	MicroRNA alterations in <i>iPSC</i> -derived dopaminergic neurons from Parkinson disease patients. <i>Neurobiology of Aging</i> , 2018, 69, 283-291.	3.1	55
16	HLA and microtubule-associated protein tau H1 haplotype associations in anti-IgLON5 disease. <i>Neurology: Neuroimmunology and NeuroInflammation</i> , 2019, 6, .	6.0	55
17	The <i>MC1R</i> melanoma risk variant p.R160W is associated with Parkinson disease. <i>Annals of Neurology</i> , 2015, 77, 889-894.	5.3	52
18	MicroRNA association with synucleinopathy conversion in rapid eye movement behavior disorder. <i>Annals of Neurology</i> , 2015, 77, 895-901.	5.3	50

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19	The prodromal phase of leucineâ€rich repeat kinase 2â€associated Parkinson disease: Clinical and imaging Studies. <i>Movement Disorders</i> , 2017, 32, 726-738.	3.9	48
20	Mutant superoxide dismutase-1 indistinguishable from wild-type causes ALS. <i>Human Molecular Genetics</i> , 2012, 21, 3568-3574.	2.9	47
21	The Genetic Architecture of Parkinson Disease in Spain: Characterizing Populationâ€Specific Risk, Differential Haplotype Structures, and Providing Etiologic Insight. <i>Movement Disorders</i> , 2019, 34, 1851-1863.	3.9	47
22	CSF microRNA Profiling in Alzheimerâ€™s Disease: a Screening and Validation Study. <i>Molecular Neurobiology</i> , 2017, 54, 6647-6654.	4.0	45
23	Parkin loss of function contributes to RTP801 elevation and neurodegeneration in Parkinsonâ€™s disease. <i>Cell Death and Disease</i> , 2014, 5, e1364-e1364.	6.3	40
24	The Small GTPase RAC1/CED-10 Is Essential in Maintaining Dopaminergic Neuron Function and Survival Against Î±-Synuclein-Induced Toxicity. <i>Molecular Neurobiology</i> , 2018, 55, 7533-7552.	4.0	40
25	Age at Onset in LRRK2-Associated PD is Modified by SNCA Variants. <i>Journal of Molecular Neuroscience</i> , 2012, 48, 245-247.	2.3	34
26	Discovering the 3â€UTR-mediated regulation of alpha-synuclein. <i>Nucleic Acids Research</i> , 2017, 45, 12888-12903.	14.5	32
27	Cerebrospinal fluid levels of coenzyme Q10 are reduced in multiple system atrophy. <i>Parkinsonism and Related Disorders</i> , 2018, 46, 16-23.	2.2	32
28	Absence of <i>LRRK2</i> mutations in a cohort of patients with idiopathic REM sleep behavior disorder. <i>Neurology</i> , 2016, 86, 1072-1073.	1.1	30
29	Possible gender-dependent association of vascular endothelial growth factor (VEGF) gene and ALS. <i>Neurology</i> , 2006, 66, 1929-1931.	1.1	29
30	Investigation of Autosomal Genetic Sex Differences in Parkinson's Disease. <i>Annals of Neurology</i> , 2021, 90, 35-42.	5.3	29
31	Accumulation of mitochondrial 7S DNA in idiopathic and LRRK2 associated Parkinson's disease. <i>EBioMedicine</i> , 2019, 48, 554-567.	6.1	28
32	Cerebrospinal fluid cytokines in multiple system atrophy: A cross-sectional Catalan MSA registry study. <i>Parkinsonism and Related Disorders</i> , 2019, 65, 3-12.	2.2	26
33	The human G93A SOD1 phenotype closely resembles sporadic amyotrophic lateral sclerosis. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2010, 81, 764-767.	1.9	25
34	Î±-Synuclein (<i>SNCA</i>) but not dynamin 3 (<i>DNM3</i>) influences age at onset of leucineâ€rich repeat kinase 2 (<i>LRRK2</i>) Parkinson's disease in Spain. <i>Movement Disorders</i> , 2018, 33, 637-641.	3.9	25
35	Mitochondrial and autophagic alterations in skin fibroblasts from Parkinson disease patients with Parkin mutations. <i>Aging</i> , 2019, 11, 3750-3767.	3.1	25
36	No evidence of association of FLJ10986 and ITPR2 with ALS in a large German cohort. <i>Neurobiology of Aging</i> , 2011, 32, 551.e1-551.e4.	3.1	22

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37	Fused in Sarcoma (FUS) gene mutations are not a frequent cause of essential tremor in Europeans. <i>Neurobiology of Aging</i> , 2013, 34, 2441.e9-2441.e11.	3.1	22
38	Exhaustion of mitochondrial and autophagic reserve may contribute to the development of LRRK2 G2019S -Parkinson's disease. <i>Journal of Translational Medicine</i> , 2018, 16, 160.	4.4	22
39	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
40	Peripheral insulin and amylin levels in Parkinson's disease. <i>Parkinsonism and Related Disorders</i> , 2020, 79, 91-96.	2.2	20
41	MTOR Pathway-Based Discovery of Genetic Susceptibility to L-DOPA-Induced Dyskinesia in Parkinson's Disease Patients. <i>Molecular Neurobiology</i> , 2019, 56, 2092-2100.	4.0	17
42	Whole-genome DNA hyper-methylation in iPSC-derived dopaminergic neurons from Parkinson's disease patients. <i>Clinical Epigenetics</i> , 2019, 11, 108.	4.1	16
43	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
44	A Novel p.Glu298Lys Mutation in the ACMSD Gene in Sporadic Parkinson's Disease. <i>Journal of Parkinson's Disease</i> , 2017, 7, 459-463.	2.8	15
45	MicroRNA Deregulation in Blood Serum Identifies Multiple System Atrophy Altered Pathways. <i>Movement Disorders</i> , 2020, 35, 1873-1879.	3.9	15
46	CCAAT/enhancer binding protein β is a transcriptional repressor of α -synuclein. <i>Cell Death and Differentiation</i> , 2020, 27, 509-524.	11.2	14
47	Epigenetic Research of Neurodegenerative Disorders Using Patient iPSC-Based Models. <i>Stem Cells International</i> , 2016, 2016, 1-16.	2.5	13
48	Transcriptional alterations in skin fibroblasts from Parkinson's disease patients with parkin mutations. <i>Neurobiology of Aging</i> , 2018, 65, 206-216.	3.1	13
49	Lack of interaction of SNCA and MAPT genotypes in Parkinson's disease. <i>European Journal of Neurology</i> , 2011, 18, e32-e32.	3.3	12
50	MAPT association with REM sleep behavior disorder. <i>Neurology: Genetics</i> , 2017, 3, e131.	1.9	10
51	Differential PhosphoSignatures in Blood Cells Identify LRRK2 G2019S Carriers in Parkinson's Disease. <i>Movement Disorders</i> , 2022, 37, 1004-1015.	3.9	9
52	What have we learned from genome-wide association studies (GWAS) in Parkinson's disease?. <i>Ageing Research Reviews</i> , 2022, 79, 101648.	10.9	9
53	Disrupted Mitochondrial and Metabolic Plasticity Underlie Comorbidity between Age-Related and Degenerative Disorders as Parkinson Disease and Type 2 Diabetes Mellitus. <i>Antioxidants</i> , 2020, 9, 1063.	5.1	8
54	Altered expression of the immunoregulatory ligand-receptor pair CD200-CD200R1 in the brain of Parkinson's disease patients. <i>Npj Parkinson's Disease</i> , 2022, 8, 27.	5.3	8

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55	Transcriptomic differences in MSA clinical variants. <i>Scientific Reports</i> , 2020, 10, 10310.	3.3	7
56	A Common Variant in the MC1R Gene (p.V92M) is associated with Alzheimer's Disease Risk. <i>Journal of Alzheimer's Disease</i> , 2017, 56, 1065-1074.	2.6	5
57	Reply. <i>Annals of Neurology</i> , 2016, 79, 161-163.	5.3	3
58	Parkinson's disease as a systemic pathology. <i>Aging</i> , 2019, 11, 1081-1082.	3.1	3
59	Smoking is associated with age at disease onset in Parkinson's disease. <i>Parkinsonism and Related Disorders</i> , 2022, 97, 79-83.	2.2	2
60	Reply. <i>Annals of Neurology</i> , 2015, 78, 153-154.	5.3	1
61	Transcriptome analysis in LRRK2 and idiopathic Parkinson's disease at different glucose levels. <i>Npj Parkinson's Disease</i> , 2021, 7, 109.	5.3	1
62	Reply. <i>Annals of Neurology</i> , 2016, 79, 868-868.	5.3	0