

Oswaldo Lorenzo-Betancor

List of Publications by Year in descending order

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Version: 2024-02-01

53
papers

2,388
citations

218677

26
h-index

214800

47
g-index

55
all docs

55
docs citations

55
times ranked

6421
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of Essential Tremor With Novel Risk Loci. <i>JAMA Neurology</i> , 2022, 79, 185.	9.0	17
2	Novel compound heterozygous FBXO7 mutations in a family with early onset Parkinson's disease. <i>Parkinsonism and Related Disorders</i> , 2020, 80, 142-147.	2.2	8
3	X-Linked Lymphoproliferative Syndrome Presenting as Adult-Onset Multi-Infarct Dementia. <i>Journal of Neuropathology and Experimental Neurology</i> , 2019, 78, 460-466.	1.7	6
4	The distribution and risk effect of GBA variants in a large cohort of PD patients from Colombia and Peru. <i>Parkinsonism and Related Disorders</i> , 2019, 63, 204-208.	2.2	31
5	Target-enriched sequencing of chromosome 17q21.31 in sporadic tauopathies reveals no candidate variants. <i>Neurobiology of Aging</i> , 2018, 66, 177.e7-177.e10.	3.1	1
6	Multiple system atrophy and apolipoprotein E. <i>Movement Disorders</i> , 2018, 33, 647-650.	3.9	15
7	<i>PCNT</i> point mutations and familial intracranial aneurysms. <i>Neurology</i> , 2018, 91, e2170-e2181.	1.1	22
8	Association study between multiple system atrophy and TREM2 p.R47H. <i>Neurology: Genetics</i> , 2018, 4, e257.	1.9	9
9	Age- and disease-dependent increase of the mitophagy marker phospho-ubiquitin in normal aging and Lewy body disease. <i>Autophagy</i> , 2018, 14, 1404-1418.	9.1	87
10	Pooled-DNA target sequencing of Parkinson genes reveals novel phenotypic associations in Spanish population. <i>Neurobiology of Aging</i> , 2018, 70, 325.e1-325.e5.	3.1	6
11	MAPT haplotype diversity in multiple system atrophy. <i>Parkinsonism and Related Disorders</i> , 2016, 30, 40-45.	2.2	23
12	RAB39B gene mutations are not a common cause of Parkinson's disease or dementia with Lewy bodies. <i>Neurobiology of Aging</i> , 2016, 45, 107-108.	3.1	21
13	A family study of DRD3 rs6280, SLC1A2 rs3794087 and MAPT rs1052553 variants in essential tremor. <i>Neurological Research</i> , 2016, 38, 880-887.	1.3	8
14	TREM2 p.R47H substitution is not associated with dementia with Lewy bodies. <i>Neurology: Genetics</i> , 2016, 2, e85.	1.9	16
15	LRRK2 variation and dementia with Lewy bodies. <i>Parkinsonism and Related Disorders</i> , 2016, 31, 98-103.	2.2	30
16	Genome-wide association study in essential tremor identifies three new loci. <i>Brain</i> , 2016, 139, 3163-3169.	7.6	78
17	Epigenetic regulation in Parkinson's disease. <i>Acta Neuropathologica</i> , 2016, 132, 515-530.	7.7	84
18	Rare variants in <i>MC1R/TUBB3</i> exon 1 are not associated with Parkinson's disease. <i>Annals of Neurology</i> , 2016, 79, 331-331.	5.3	10

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19	<i>MAPT</i> haplotype H1G is associated with increased risk of dementia with Lewy bodies. <i>Alzheimer's and Dementia</i> , 2016, 12, 1297-1304.	0.8	32
20	Pathophysiological relevance of PINK1-dependent ubiquitin phosphorylation. <i>EMBO Reports</i> , 2015, 16, 1114-1130.	4.5	147
21	Automated Neuromelanin Imaging as a Diagnostic Biomarker for Parkinson's Disease. <i>Movement Disorders</i> , 2015, 30, 945-952.	3.9	138
22	DNAJC13 p.Asn855Ser mutation screening in Parkinson's disease and pathologically confirmed Lewy body disease patients. <i>European Journal of Neurology</i> , 2015, 22, 1323-1325.	3.3	21
23	Heme Oxygenase 1 and 2 Common Genetic Variants and Risk for Essential Tremor. <i>Medicine (United Tj ETQq1 1 0,784314 rgBT /Ove</i>	1.0	20
24	Mitochondrial targeting sequence variants of the <i>CHCHD2</i> gene are a risk for Lewy body disorders. <i>Neurology</i> , 2015, 85, 2016-2025.	1.1	51
25	Rare variants in β -Amyloid precursor protein (APP) and Parkinson's disease. <i>European Journal of Human Genetics</i> , 2015, 23, 1328-1333.	2.8	50
26	Missense mutations in <i>TENM4</i> , a regulator of axon guidance and central myelination, cause essential tremor. <i>Human Molecular Genetics</i> , 2015, 24, 5677-5686.	2.9	134
27	TREM2 R47H variant and risk of essential tremor: A cross-sectional international multicenter study. <i>Parkinsonism and Related Disorders</i> , 2015, 21, 306-309.	2.2	28
28	Role for the microtubule-associated protein tau variant p.A152T in risk of α -synucleinopathies. <i>Neurology</i> , 2015, 85, 1680-1686.	1.1	31
29	Exonic Re-Sequencing of the Chromosome 2q24.3 Parkinson's Disease Locus. <i>PLoS ONE</i> , 2015, 10, e0128586.	2.5	0
30	Analysis of Nuclear Export Sequence Regions of FUS-Related RNA-Binding Proteins in Essential Tremor. <i>PLoS ONE</i> , 2014, 9, e111989.	2.5	10
31	An association study between Heme oxygenase-1 genetic variants and Parkinson's disease. <i>Frontiers in Cellular Neuroscience</i> , 2014, 8, 298.	3.7	39
32	Analysis of COQ2 gene in multiple system atrophy. <i>Molecular Neurodegeneration</i> , 2014, 9, 44.	10.8	40
33	Rare coding variants in the phospholipase D3 gene confer risk for Alzheimer's disease. <i>Nature</i> , 2014, 505, 550-554.	27.8	425
34	LINGO1 rs9652490 and rs11856808 polymorphisms are not associated with risk for multiple sclerosis. <i>BMC Neurology</i> , 2013, 13, 34.	1.8	10
35	Mutational Screening of PARKIN Identified a 3' UTR Variant (rs62637702) Associated with Parkinson's Disease. <i>Journal of Molecular Neuroscience</i> , 2013, 50, 264-269.	2.3	11
36	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

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37	Update on genetics of essential tremor. <i>Acta Neurologica Scandinavica</i> , 2013, 128, 359-371.	2.1	41
38	No association of the SLC1A2 rs3794087 allele with risk for essential tremor in the Spanish population. <i>Pharmacogenetics and Genomics</i> , 2013, 23, 587-590.	1.5	22
39	Dopaminergic Neuronal Imaging in Genetic Parkinson's Disease: Insights into Pathogenesis. <i>PLoS ONE</i> , 2013, 8, e69190.	2.5	55
40	Common variation in the <i>LRRK2</i> gene is a risk factor for Parkinson's disease. <i>Movement Disorders</i> , 2012, 27, 1823-1826.	3.9	14
41	LINGO1 rs9652490 and rs11856808 are not associated with the risk of Parkinson's disease: Results of a meta-analysis. <i>Parkinsonism and Related Disorders</i> , 2012, 18, 657-659.	2.2	7
42	LINGO1 and risk for essential tremor: Results of a meta-analysis of rs9652490 and rs11856808. <i>Journal of the Neurological Sciences</i> , 2012, 317, 52-57.	0.6	39
43	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
44	H1-MAPT and the Risk for Familial Essential Tremor. <i>PLoS ONE</i> , 2012, 7, e41581.	2.5	17
45	A Search for SNCA 3' UTR Variants Identified SNP rs356165 as a Determinant of Disease Risk and Onset Age in Parkinson's Disease. <i>Journal of Molecular Neuroscience</i> , 2012, 47, 425-430.	2.3	49
46	<i>LRRK2</i> haplotype-sharing analysis in Parkinson's disease reveals a novel p.S1761R mutation. <i>Movement Disorders</i> , 2012, 27, 146-150.	3.9	19
47	Gamma-aminobutyric acid GABRA4, GABRE, and GABRQ receptor polymorphisms and risk for essential tremor. <i>Pharmacogenetics and Genomics</i> , 2011, 21, 436-439.	1.5	28
48	A polymorphism located at an ATG transcription start site of the heme oxygenase-2 gene is associated with classical Parkinson's disease. <i>Pharmacogenetics and Genomics</i> , 2011, 21, 565-571.	1.5	18
49	Lack of association of LINGO1 rs9652490 and rs11856808 SNPs with familial essential tremor. <i>European Journal of Neurology</i> , 2011, 18, 1085-1089.	3.3	30
50	Gamma-aminobutyric acid (GABA) receptor rho (GABRR) polymorphisms and risk for essential tremor. <i>Journal of Neurology</i> , 2011, 258, 203-211.	3.6	28
51	<i>LINGO1</i> gene analysis in Parkinson's disease phenotypes. <i>Movement Disorders</i> , 2011, 26, 722-727.	3.9	17
52	Replication of <i>MAPT</i> and <i>SNCA</i> , but not <i>PARK16</i> , as susceptibility genes for Parkinson's disease. <i>Movement Disorders</i> , 2011, 26, 819-823.	3.9	64
53	PINK1-linked parkinsonism is associated with Lewy body pathology. <i>Brain</i> , 2010, 133, 1128-1142.	7.6	223