

Andrea Carmine Belin

List of Publications by Year in descending order

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Version: 2024-02-01

79
papers

4,457
citations

126907

33
h-index

118850

62
g-index

83
all docs

83
docs citations

83
times ranked

9067
citing authors

#	ARTICLE	IF	CITATIONS
1	Mendelian Randomisation Study of Smoking, Alcohol, and Coffee Drinking in Relation to Parkinson's Disease. <i>Journal of Parkinson's Disease</i> , 2022, 12, 267-282.	2.8	21
2	Dairy Intake and Parkinson's Disease: A Mendelian Randomization Study. <i>Movement Disorders</i> , 2022, 37, 857-864.	3.9	15
3	Board Walk " January 2022. <i>Cephalalgia</i> , 2022, 42, 87-87.	3.9	0
4	Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. <i>Nature Genetics</i> , 2022, 54, 152-160.	21.4	135
5	Elucidating the relationship between migraine risk and brain structure using genetic data. <i>Brain</i> , 2022, 145, 3214-3224.	7.6	7
6	The molecular clock gene cryptochrome 1 (<i>CRY1</i>) and its role in cluster headache. <i>Cephalalgia</i> , 2021, 41, 1374-1381.	3.9	8
7	Genome-Wide Association Study Identifies Risk Loci for Cluster Headache. <i>Annals of Neurology</i> , 2021, 90, 193-202.	5.3	31
8	PER Gene Family Polymorphisms in Relation to Cluster Headache and Circadian Rhythm in Sweden. <i>Brain Sciences</i> , 2021, 11, 1108.	2.3	1
9	Analysis of NOS Gene Polymorphisms in Relation to Cluster Headache and Predisposing Factors in Sweden. <i>Brain Sciences</i> , 2021, 11, 34.	2.3	5
10	Genetic Screening of Plasticity Regulating Nogo-Type Signaling Genes in Migraine. <i>Brain Sciences</i> , 2020, 10, 5.	2.3	1
11	Habitual sleep disturbances and migraine: a Mendelian randomization study. <i>Annals of Clinical and Translational Neurology</i> , 2020, 7, 2370-2380.	3.7	18
12	A genome-wide cross-phenotype meta-analysis of the association of blood pressure with migraine. <i>Nature Communications</i> , 2020, 11, 3368.	12.8	49
13	Calcitonin Gene-Related Peptide (CGRP) and Cluster Headache. <i>Brain Sciences</i> , 2020, 10, 30.	2.3	35
14	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinson's disease. <i>Nature Genetics</i> , 2020, 52, 482-493.	21.4	216
15	Cross-trait analyses with migraine reveal widespread pleiotropy and suggest a vascular component to migraine headache. <i>International Journal of Epidemiology</i> , 2020, 49, 1022-1031.	1.9	34
16	Anoctamin 3: A Possible Link between Cluster Headache and Ca ²⁺ Signaling. <i>Brain Sciences</i> , 2019, 9, 184.	2.3	8
17	Low prevalence of known pathogenic mutations in dominant PD genes: A Swedish multicenter study. <i>Parkinsonism and Related Disorders</i> , 2019, 66, 158-165.	2.2	12
18	Involvement of CGRP receptor RAMP1 in cluster headache: A Swedish case-control study. <i>Cephalalgia Reports</i> , 2019, 2, 251581631987988.	0.7	2

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19	Using global team science to identify genetic parkinson's disease worldwide. <i>Annals of Neurology</i> , 2019, 86, 153-157.	5.3	26
20	Experimental Tools to Study the Regulation and Function of the Choroid Plexus. <i>Neuromethods</i> , 2019, , 205-230.	0.3	2
21	Analysis of HCRTR2 Gene Variants and Cluster Headache in Sweden. <i>Headache</i> , 2019, 59, 410-417.	3.9	20
22	Common Variant Burden Contributes to the Familial Aggregation of Migraine in 1,589 Families. <i>Neuron</i> , 2018, 98, 743-753.e4.	8.1	63
23	A genetic CLOCK variant associated with cluster headache causing increased mRNA levels. <i>Cephalalgia</i> , 2018, 38, 496-502.	3.9	43
24	Cluster headache " clinical pattern and a new severity scale in a Swedish cohort. <i>Cephalalgia</i> , 2018, 38, 1286-1295.	3.9	52
25	The choroid plexus harbors a circadian oscillator modulated by estrogens. <i>Chronobiology International</i> , 2018, 35, 270-279.	2.0	28
26	Implications for the migraine SNP rs1835740 in a Swedish cluster headache population. <i>Journal of Headache and Pain</i> , 2018, 19, 100.	6.0	6
27	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	12.6	1,085
28	Molecular genetic overlap between migraine and major depressive disorder. <i>European Journal of Human Genetics</i> , 2018, 26, 1202-1216.	2.8	56
29	S100B polymorphisms are associated with age of onset of Parkinson's disease. <i>BMC Medical Genetics</i> , 2018, 19, 42.	2.1	17
30	Genetic Variations and mRNA Expression of NRF2 in Parkinson's Disease. <i>Parkinson's Disease</i> , 2017, 2017, 1-7.	1.1	12
31	Shared genetic risk between migraine and coronary artery disease: A genome-wide analysis of common variants. <i>PLoS ONE</i> , 2017, 12, e0185663.	2.5	44
32	Screening of genetic variants in ADCYAP1R1, MME and 14q21 in a Swedish cluster headache cohort. <i>Journal of Headache and Pain</i> , 2017, 18, 88.	6.0	20
33	Screening of Two ADH4 Variations in a Swedish Cluster Headache Case-Control Material. <i>Headache</i> , 2016, 56, 835-840.	3.9	19
34	Ryanodine-sensitive intracellular Ca ²⁺ channels are involved in the output from the SCN circadian clock. <i>European Journal of Neuroscience</i> , 2016, 44, 2504-2514.	2.6	14
35	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. <i>Nature Genetics</i> , 2016, 48, 856-866.	21.4	520
36	Strong association between glucocerebrosidase mutations and Parkinson's disease in Sweden. <i>Neurobiology of Aging</i> , 2016, 45, 212.e5-212.e11.	3.1	50

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37	No Association Between rs7077361 in ITGA8 and Parkinson's Disease in Sweden. <i>The Open Neurology Journal</i> , 2016, 10, 25-29.	0.4	2
38	ReMo-SNPs: a new software tool for identification of polymorphisms in regions and motifs genome-wide. <i>Genetical Research</i> , 2015, 97, e8.	0.9	1
39	A replication study of GWAS findings in migraine identifies association in a Swedish case-control sample. <i>BMC Medical Genetics</i> , 2014, 15, 38.	2.1	30
40	The HLA-DRA variation rs3129882 is not associated with Parkinson's disease in Sweden. <i>Parkinsonism and Related Disorders</i> , 2013, 19, 701-702.	2.2	8
41	Association of a protective paraoxonase 1 (PON1) polymorphism in Parkinson's disease. <i>Neuroscience Letters</i> , 2012, 522, 30-35.	2.1	27
42	Adh1 and Adh1/4 knockout mice as possible rodent models for presymptomatic Parkinson's disease. <i>Behavioural Brain Research</i> , 2012, 227, 252-257.	2.2	13
43	MAG11 Copy Number Variation in Bipolar Affective Disorder and Schizophrenia. <i>Biological Psychiatry</i> , 2012, 71, 922-930.	1.3	41
44	Genetic Screening of the Mitochondrial Rho GTPases MIRO1 and MIRO2 in Parkinson's Disease. <i>The Open Neurology Journal</i> , 2012, 6, 1-5.	0.4	10
45	Modeling Parkinson's disease genetics: Altered function of the dopamine system in Adh4 knockout mice. <i>Behavioural Brain Research</i> , 2011, 217, 439-445.	2.2	12
46	Genetic studies of the protein kinase AKT1 in Parkinson's disease. <i>Neuroscience Letters</i> , 2011, 501, 41-44.	2.1	9
47	Altered enzymatic activity and allele frequency of OMI/HTRA2 in Alzheimer's disease. <i>FASEB Journal</i> , 2011, 25, 1345-1352.	0.5	25
48	DJ-1 Mutations are Rare in a Swedish Parkinson Cohort. <i>The Open Neurology Journal</i> , 2011, 5, 8-11.	0.4	3
49	Possible Involvement of a Mitochondrial Translation Initiation Factor 3 Variant Causing Decreased mRNA Levels in Parkinson's Disease. <i>Parkinson's Disease</i> , 2010, 2010, 1-5.	1.1	14
50	Variations of the CAG trinucleotide repeat in DNA polymerase gamma (POLG1) is associated with Parkinson's disease in Sweden. <i>Neuroscience Letters</i> , 2010, 485, 117-120.	2.1	32
51	PITX3 polymorphism is associated with early onset Parkinson's disease. <i>Neurobiology of Aging</i> , 2010, 31, 114-117.	3.1	65
52	Do polymorphisms in transcription factors LMX1A and LMX1B influence the risk for Parkinson's disease?. <i>Journal of Neural Transmission</i> , 2009, 116, 333-338.	2.8	39
53	Cytochrome P450 2E1 gene polymorphisms/haplotypes and Parkinson's disease in a Swedish population. <i>Journal of Neural Transmission</i> , 2009, 116, 567-573.	2.8	26
54	Association of a polymorphism in the ABCB1 gene with Parkinson's disease. <i>Parkinsonism and Related Disorders</i> , 2009, 15, 422-424.	2.2	38

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55	Expression of multi-drug resistance 1 mRNA in human and rodent tissues: reduced levels in Parkinson patients. <i>Cell and Tissue Research</i> , 2008, 334, 179-185.	2.9	41
56	Parkinson's disease: A genetic perspective. <i>FEBS Journal</i> , 2008, 275, 1377-1383.	4.7	97
57	Developmental regulation of leucine-rich repeat kinase 1 and 2 expression in the brain and other rodent and human organs: Implications for Parkinson's disease. <i>Neuroscience</i> , 2008, 152, 429-436.	2.3	96
58	Investigation of genes related to familial forms of Parkinson's disease – With focus on the Parkin gene. <i>Parkinsonism and Related Disorders</i> , 2008, 14, 520-522.	2.2	7
59	Cerebellar α -synuclein levels are decreased in Parkinson's disease and do not correlate with <i>SNCA</i> polymorphisms associated with disease in a Swedish material. <i>FASEB Journal</i> , 2008, 22, 3509-3514.	0.5	41
60	S18Y, UCH-L1 and Parkinson's Disease. <i>European Neurological Review</i> , 2008, 3, 41.	0.5	1
61	S18Y in ubiquitin carboxy-terminal hydrolase L1 (UCH-L1) associated with decreased risk of Parkinson's disease in Sweden. <i>Parkinsonism and Related Disorders</i> , 2007, 13, 295-298.	2.2	46
62	DJ-1 and UCH-L1 gene activity patterns in the brains of controls, Parkinson and schizophrenia patients and in rodents. <i>Physiology and Behavior</i> , 2007, 92, 46-53.	2.1	21
63	Association study of two genetic variants in mitochondrial transcription factor A (TFAM) in Alzheimer's and Parkinson's disease. <i>Neuroscience Letters</i> , 2007, 420, 257-262.	2.1	41
64	Cyclooxygenase-2 polymorphisms in Parkinson's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007, 144B, 367-369.	1.7	10
65	High and complementary expression patterns of alcohol and aldehyde dehydrogenases in the gastrointestinal tract. <i>FEBS Journal</i> , 2007, 274, 1212-1223.	4.7	30
66	Lack of replication of thirteen single-nucleotide polymorphisms implicated in Parkinson's disease: a large-scale international study. <i>Lancet Neurology</i> , The, 2006, 5, 917-923.	10.2	83
67	Leucine-rich repeat kinase 2 (LRRK2) mutations in a Swedish Parkinson cohort and a healthy nonagenarian. <i>Movement Disorders</i> , 2006, 21, 1731-1734.	3.9	47
68	LRRK2 expression linked to dopamine-innervated areas. <i>Annals of Neurology</i> , 2006, 59, 714-719.	5.3	166
69	Interaction of polymorphisms in the genes encoding interleukin-6 and estrogen receptor beta on the susceptibility to Parkinson's disease. , 2005, 133B, 88-92.		68
70	Investigation of genes coding for inflammatory components in Parkinson's disease. <i>Movement Disorders</i> , 2005, 20, 569-573.	3.9	46
71	Tissue- and species-specific expression patterns of class I, III, and IV Adh and Aldh1 mRNAs in rodent embryos. <i>Cell and Tissue Research</i> , 2005, 322, 227-236.	2.9	38
72	A Rare Truncating Mutation in ADH1C (G78Stop) Shows Significant Association With Parkinson Disease in a Large International Sample. <i>Archives of Neurology</i> , 2005, 62, 74.	4.5	57

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73	Association between the estrogen receptor beta gene and age of onset of Parkinson's disease. Psychoneuroendocrinology, 2004, 29, 993-998.	2.7	49
74	Lack of association between the BDNF Val66Met polymorphism and Parkinson's disease in a Swedish population. Annals of Neurology, 2003, 53, 823-823.	5.3	44
75	Distribution of class I, III and IV alcohol dehydrogenase mRNAs in the adult rat, mouse and human brain. FEBS Journal, 2003, 270, 1316-1326.	0.2	78
76	Decreased ethanol preference and wheel running in Nurr1-deficient mice. European Journal of Neuroscience, 2003, 17, 2418-2424.	2.6	25
77	ALDH1 mRNA: presence in human dopamine neurons and decreases in substantia nigra in Parkinson's disease and in the ventral tegmental area in schizophrenia. Neurobiology of Disease, 2003, 14, 637-647.	4.4	148
78	Two NOTCH4 polymorphisms and their relation to schizophrenia susceptibility and different personality traits. Psychiatric Genetics, 2003, 13, 23-28.	1.1	16
79	Further evidence for an association of the Paraoxonase 1 (PON1) Met-54 allele with Parkinson's disease. Movement Disorders, 2002, 17, 764-766.	3.9	49