

Hong Jiang

List of Publications by Year in descending order

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92
papers

2,370
citations

304743
22
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100
all docs

100
docs citations

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times ranked

2852
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of the Largest SCA36 Pedigree in Asia: with Multimodal Neuroimaging Evaluation for the First Time. <i>Cerebellum</i> , 2022, 21, 358-367.	2.5	3
2	Blood Neurofilament Light Chain in Genetic Ataxia: A Meta-Analysis. <i>Movement Disorders</i> , 2022, 37, 171-181.	3.9	8
3	<i>TMEM151A</i> Variants Cause Paroxysmal Kinesigenic Dyskinesia: A Large-Sample Study. <i>Movement Disorders</i> , 2022, 37, 545-552.	3.9	20
4	A Variant in Genes of the NPY System as Modifier Factor of Machado-Joseph Disease in the Chinese Population. <i>Frontiers in Aging Neuroscience</i> , 2022, 14, 822657.	3.4	0
5	Coffin-Siris syndrome in two chinese patients with novel pathogenic variants of ARID1A and SMARCA4. <i>Genes and Genomics</i> , 2022, , 1.	1.4	1
6	Paving the Way Toward Meaningful Trials in Ataxias: An Ataxia Global Initiative Perspective. <i>Movement Disorders</i> , 2022, 37, 1125-1130.	3.9	21
7	Profiling the Genome-Wide Landscape of Short Tandem Repeats by Long-Read Sequencing. <i>Frontiers in Genetics</i> , 2022, 13, .	2.3	4
8	The progression rate of spinocerebellar ataxia type 3 varies with disease stage. <i>Journal of Translational Medicine</i> , 2022, 20, 226.	4.4	5
9	Rare, pathogenic variants in LRP10 are associated with amyotrophic lateral sclerosis in patients from mainland China. <i>Neurobiology of Aging</i> , 2021, 97, 145.e17-145.e22.	3.1	5
10	No genetic evidence for the involvement of GGC repeat expansions of the NOTCH2NLC gene in Chinese patients with multiple system atrophy. <i>Neurobiology of Aging</i> , 2021, 97, 144.e5-144.e7.	3.1	5
11	Prediction of the Age at Onset of Spinocerebellar Ataxia Type 3 with Machine Learning. <i>Movement Disorders</i> , 2021, 36, 216-224.	3.9	11
12	Age is an important independent modifier of SCA3 phenotype severity. <i>Neuroscience Letters</i> , 2021, 741, 135510.	2.1	4
13	Human stem cell models of polyglutamine diseases: Sources for disease models and cell therapy. <i>Experimental Neurology</i> , 2021, 337, 113573.	4.1	5
14	ATP10B variants in Parkinson's disease: a large cohort study in Chinese mainland population. <i>Acta Neuropathologica</i> , 2021, 141, 805-806.	7.7	8
15	Recommendations for the diagnosis and treatment of paroxysmal kinesigenic dyskinesia: an expert consensus in China. <i>Translational Neurodegeneration</i> , 2021, 10, 7.	8.0	19
16	<i>UQCRC1</i> variants in Parkinson's disease: a large cohort study in Chinese mainland population. <i>Brain</i> , 2021, 144, e54-e54.	7.6	5
17	New Model for Estimation of the Age at Onset in Spinocerebellar Ataxia Type 3. <i>Neurology</i> , 2021, 96, e2885-e2895.	1.1	7
18	Generation of spinocerebellar ataxia type 3 patient-derived induced pluripotent stem cell line (CSUXHi005-A) from human urine epithelial cells. <i>Stem Cell Research</i> , 2021, 53, 102289.	0.7	3

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19	Regional Brain and Spinal Cord Volume Loss in Spinocerebellar Ataxia Type 3. <i>Movement Disorders</i> , 2021, 36, 2273-2281.	3.9	37
20	Anxiety and depression in spinocerebellar ataxia patients during the COVID-19 pandemic in China: A cross-sectional study. <i>Journal of Clinical Neuroscience</i> , 2021, 88, 39-46.	1.5	7
21	Genotype and phenotype distribution of 435 patients with Charcot-Marie-Tooth disease from central south China. <i>European Journal of Neurology</i> , 2021, 28, 3774-3783.	3.3	19
22	Genetic etiology of a Chinese ataxia cohort: Expanding the mutational spectrum of hereditary ataxias. <i>Parkinsonism and Related Disorders</i> , 2021, 89, 120-127.	2.2	5
23	CRISPR/Cas9 mediated gene correction ameliorates abnormal phenotypes in spinocerebellar ataxia type 3 patient-derived induced pluripotent stem cells. <i>Translational Psychiatry</i> , 2021, 11, 479.	4.8	15
24	<i>PSAP</i> variants in Parkinson's disease: a large cohort study in Chinese mainland population. <i>Brain</i> , 2021, 144, e25-e25.	7.6	7
25	Cholecystokinin 1 receptor activation restores normal mTORC1 signaling and is protective to Purkinje cells of SCA mice. <i>Cell Reports</i> , 2021, 37, 109831.	6.4	11
26	Polyglutamine-expanded ataxin3 alter specific gene expressions through changing DNA methylation status in SCA3/MJD. <i>Aging</i> , 2021, 13, 3680-3698.	3.1	4
27	Expansion of GGC repeat in the human-specific NOTCH2NLC gene is associated with essential tremor. <i>Brain</i> , 2020, 143, 222-233.	7.6	139
28	Identification of GGC repeat expansion in the NOTCH2NLC gene in amyotrophic lateral sclerosis. <i>Neurology</i> , 2020, 95, e3394-e3405.	1.1	59
29	Micro-structural white matter abnormalities and cognitive impairment in asymptomatic carotid plaque patients: A DTI study using TBSS analysis. <i>Clinical Neurology and Neurosurgery</i> , 2020, 197, 106096.	1.4	3
30	GCH1 variants contribute to the risk and earlier age-at-onset of Parkinson's disease: a two-cohort case-control study. <i>Translational Neurodegeneration</i> , 2020, 9, 31.	8.0	30
31	MR Imaging of SCA3/MJD. <i>Frontiers in Neuroscience</i> , 2020, 14, 749.	2.8	18
32	Biallelic Intronic AAGGG Expansion of RFC1 is Related to Multiple System Atrophy. <i>Annals of Neurology</i> , 2020, 88, 1132-1143.	5.3	41
33	Association of serum neurofilament light and disease severity in patients with spinocerebellar ataxia type 3. <i>Neurology</i> , 2020, 95, e2977-e2987.	1.1	19
34	Myeloteriosis in an ALPS5 patient with primary immune dysregulation syndrome. <i>CNS Neuroscience and Therapeutics</i> , 2020, 26, 773-775.	3.9	1
35	The role of genetics in Parkinson's disease: a large cohort study in Chinese mainland population. <i>Brain</i> , 2020, 143, 2220-2234.	7.6	97
36	Identification of novel mutations in TSC1 and TSC2 for tuberous sclerosis complex by targeted next-generation sequencing and ACMG guidelines. <i>Child's Nervous System</i> , 2020, 36, 1827-1830.	1.1	0

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37	Profiling of mitochondrial genomes in SCA3/MJD patients from mainland China. <i>Gene</i> , 2020, 738, 144487.	2.2	1
38	Generation of induced pluripotent stem cell line (CSUXHi002-A) from a patient with spinocerebellar ataxia type 1. <i>Stem Cell Research</i> , 2020, 45, 101816.	0.7	6
39	Mutation analysis of CAPN1 in Chinese populations with spastic paraplegia and related neurodegenerative diseases. <i>Journal of the Neurological Sciences</i> , 2020, 411, 116691.	0.6	4
40	Central motor conduction time in spinocerebellar ataxia: a meta-analysis. <i>Aging</i> , 2020, 12, 25718-25729.	3.1	6
41	RNA Expression Profile and Potential Biomarkers in Patients With Spinocerebellar Ataxia Type 3 From Mainland China. <i>Frontiers in Genetics</i> , 2019, 10, 566.	2.3	4
42	Alterations of the Gut Microbiota in Multiple System Atrophy Patients. <i>Frontiers in Neuroscience</i> , 2019, 13, 1102.	2.8	42
43	Gene-Related Cerebellar Neurodegeneration in SCA3/MJD: A Case-Controlled Imaging-Genetic Study. <i>Frontiers in Neurology</i> , 2019, 10, 1025.	2.4	21
44	Clinical findings of autosomal-dominant striatal degeneration and PDE8B mutation screening in parkinsonism and related disorders. <i>Parkinsonism and Related Disorders</i> , 2019, 69, 94-98.	2.2	5
45	Generation of an induced pluripotent stem cell line (XHCSUi001-A) from urine cells of a patient with spinocerebellar ataxia type 3. <i>Stem Cell Research</i> , 2019, 40, 101555.	0.7	6
46	Genetic and clinical analyses of spinocerebellar ataxia type 8 in mainland China. <i>Journal of Neurology</i> , 2019, 266, 2979-2986.	3.6	7
47	Identification of a potential exosomal biomarker in spinocerebellar ataxia Type 3/Machado-Joseph disease. <i>Epigenomics</i> , 2019, 11, 1037-1056.	2.1	23
48	Expansion of Human-Specific GGC Repeat in Neuronal Intranuclear Inclusion Disease-Related Disorders. <i>American Journal of Human Genetics</i> , 2019, 105, 166-176.	6.2	212
49	(CAG) _n loci as genetic modifiers of age at onset in patients with spinocerebellar ataxia type 1 from mainland China. <i>European Journal of Neurology</i> , 2019, 26, 1130-1136.	3.3	11
50	Long-read sequencing identified intronic repeat expansions in SAMD12 from Chinese pedigrees affected with familial cortical myoclonic tremor with epilepsy. <i>Journal of Medical Genetics</i> , 2019, 56, 265-270.	3.2	82
51	Polymorphisms in DNA methylation-related genes are linked to the phenotype of Machado-Joseph disease. <i>Neurobiology of Aging</i> , 2019, 75, 225.e1-225.e8.	3.1	5
52	A novel mutation in MYORG causes primary familial brain calcification with central neuropathic pain. <i>Clinical Genetics</i> , 2019, 95, 433-435.	2.0	17
53	Analysis of (CAG) _n expansion in ATXN1, ATXN2 and ATXN3 in Chinese patients with multiple system atrophy. <i>Scientific Reports</i> , 2018, 8, 3889.	3.3	11
54	Updated frequency analysis of spinocerebellar ataxia in China. <i>Brain</i> , 2018, 141, e22-e22.	7.6	33

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55	Association of <i>TNF-Î±</i> rs1799964 and <i>IL-1Î²</i> rs16944 polymorphisms with multiple system atrophy in Chinese Han population. <i>International Journal of Neuroscience</i> , 2018, 128, 761-764.	1.6	16
56	Familial paroxysmal kinesigenic dyskinesia is associated with mutations in the KCNA1 gene. <i>Human Molecular Genetics</i> , 2018, 27, 625-637.	2.9	43
57	Identifying SYNE1 Ataxia With Novel Mutations in a Chinese Population. <i>Frontiers in Neurology</i> , 2018, 9, 1111.	2.4	14
58	Roles of Post-translational Modifications in Spinocerebellar Ataxias. <i>Frontiers in Cellular Neuroscience</i> , 2018, 12, 290.	3.7	13
59	Genetic modifiers of age-at-onset in polyglutamine diseases. <i>Ageing Research Reviews</i> , 2018, 48, 99-108.	10.9	20
60	Noncoding RNAs and Base Modifications: Epigenomic Players Implicated in Neurological Disorders and Tumorigenesis. <i>International Journal of Genomics</i> , 2018, 2018, 1-2.	1.6	0
61	Coding mutations in NUS1 contribute to Parkinson's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 11567-11572.	7.1	78
62	Cerebellar lncRNA Expression Profile Analysis of SCA3/MJD Mice. <i>International Journal of Genomics</i> , 2018, 2018, 1-6.	1.6	5
63	Investigation on modulation of DNA repair pathways in Chinese MJD patients. <i>Neurobiology of Aging</i> , 2018, 71, 267.e5-267.e6.	3.1	5
64	Birt-Hogg-DubÃ© syndrome in two Chinese families with mutations in the FLCN gene. <i>BMC Medical Genetics</i> , 2018, 19, 14.	2.1	13
65	Is the High Frequency of Machado-Joseph Disease in China Due to New Mutational Origins?. <i>Frontiers in Genetics</i> , 2018, 9, 740.	2.3	17
66	The Effects of Dracocephalum heterophyllum Benth Flavonoid on Hypertrophic Cardiomyocytes Induced by Angiotensin II in Rats. <i>Medical Science Monitor</i> , 2018, 24, 6322-6330.	1.1	5
67	Alteration of methylation status in the ATXN3 gene promoter region is linked to the SCA3/MJD. <i>Neurobiology of Aging</i> , 2017, 53, 192.e5-192.e10.	3.1	18
68	Ubiquitin-related network underlain by (CAG) _n loci modulate age at onset in Machado-Joseph disease. <i>Brain</i> , 2017, 140, e25-e25.	7.6	10
69	Novel mutations in ADSL for Adenylosuccinate Lyase Deficiency identified by the combination of Trio-WES and constantly updated guidelines. <i>Scientific Reports</i> , 2017, 7, 1625.	3.3	12
70	Rare GCH1 heterozygous variants contributing to Parkinson's disease. <i>Brain</i> , 2017, 140, e41-e41.	7.6	21
71	Genetic and clinical analysis of spinocerebellar ataxia type 36 in Mainland China. <i>Clinical Genetics</i> , 2016, 90, 141-148.	2.0	13
72	(CAG) _n loci as genetic modifiers of age-at-onset in patients with Machado-Joseph disease from mainland China. <i>Brain</i> , 2016, 139, e41-e41.	7.6	37

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73	<i>ATXN2</i> polymorphism modulates age at onset in Machado-Joseph disease. <i>Brain</i> , 2016, 139, aww176.	7.6	16
74	Identification of a de novo DYNC1H1 mutation via WES according to published guidelines. <i>Scientific Reports</i> , 2016, 6, 20423.	3.3	20
75	miRNA profiling in autism spectrum disorder in China. <i>Genomics Data</i> , 2015, 6, 108-109.	1.3	6
76	Primary erythromelalgia: a review. <i>Orphanet Journal of Rare Diseases</i> , 2015, 10, 127.	2.7	90
77	Two Novel SNPs in ATXN3 3' UTR May Decrease Age at Onset of SCA3/MJD in Chinese Patients. <i>PLoS ONE</i> , 2015, 10, e0117488.	2.5	19
78	Analysis of the GGGGCC Repeat Expansions of the C9orf72 Gene in SCA3/MJD Patients from China. <i>PLoS ONE</i> , 2015, 10, e0130336.	2.5	11
79	Polygenic determinants of Parkinson's disease in a Chinese population. <i>Neurobiology of Aging</i> , 2015, 36, 1765.e1-1765.e6.	3.1	73
80	Investigation of Gene Regulatory Networks Associated with Autism Spectrum Disorder Based on miRNA Expression in China. <i>PLoS ONE</i> , 2015, 10, e0129052.	2.5	50
81	High Serum GFAP Levels in SCA3/MJD May Not Correlate with Disease Progression. <i>Cerebellum</i> , 2015, 14, 677-681.	2.5	9
82	Targeted Next-Generation Sequencing Revealed Novel Mutations in Chinese Ataxia Telangiectasia Patients: A Precision Medicine Perspective. <i>PLoS ONE</i> , 2015, 10, e0139738.	2.5	8
83	miR-25 alleviates polyQ-mediated cytotoxicity by silencing <i>ATXN3</i> . <i>FEBS Letters</i> , 2014, 588, 4791-4798.	2.8	37
84	Huntingtin gene CAG repeat numbers in Chinese patients with Huntington's disease and controls. <i>European Journal of Neurology</i> , 2014, 21, 637-642.	3.3	20
85	The APOE ϵ 2 allele may decrease the age at onset in patients with spinocerebellar ataxia type 3 or Machado-Joseph disease from the Chinese Han population. <i>Neurobiology of Aging</i> , 2014, 35, 2179.e15-2179.e18.	3.1	25
86	Using next-generation sequencing as a genetic diagnostic tool in rare autosomal recessive neurologic Mendelian disorders. <i>Neurobiology of Aging</i> , 2013, 34, 2442.e11-2442.e17.	3.1	19
87	Spinocerebellar ataxia type 27 (SCA27) is an uncommon cause of dominant ataxia among Chinese Han population. <i>Neuroscience Letters</i> , 2012, 520, 16-19.	2.1	12
88	Identification of PRRT2 as the causative gene of paroxysmal kinesigenic dyskinesias. <i>Brain</i> , 2011, 134, 3493-3501.	7.6	263
89	<i>PLA2G6</i> gene mutation in autosomal recessive early-onset parkinsonism in a Chinese cohort. <i>Neurology</i> , 2011, 77, 75-81.	1.1	87
90	Analysis of SCA2 and SCA3/MJD repeats in Parkinson's disease in mainland China: Genetic, clinical, and positron emission tomography findings. <i>Movement Disorders</i> , 2009, 24, 2007-2011.	3.9	43

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91	Mutation analysis of <i>Parkin</i> , <i>PINK1</i> , <i>DJ-1</i> and <i>ATP13A2</i> genes in Chinese patients with autosomal recessive early-onset Parkinsonism. <i>Movement Disorders</i> , 2008, 23, 2074-2079.	3.9	61
92	Mutation analysis of hereditary multiple exostoses in the Chinese. <i>Human Genetics</i> , 1999, 105, 45-50.	3.8	16