

# Giovanni Coppola

## List of Publications by Year in descending order

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Version: 2024-02-01

241  
papers

25,070  
citations

9786

73  
h-index

8866

145  
g-index

261  
all docs

261  
docs citations

261  
times ranked

34103  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic and functional analysis of a Pacific hagfish opioid system. <i>Journal of Neuroscience Research</i> , 2022, 100, 19-34.	2.9	2
2	Epigenetic clock and methylation studies in vervet monkeys. <i>GeroScience</i> , 2022, 44, 699-717.	4.6	18
3	IAPP-induced beta cell stress recapitulates the islet transcriptome in type 2 diabetes. <i>Diabetologia</i> , 2022, 65, 173-187.	6.3	19
4	Cortical and subcortical pathological burden and neuronal loss in an autopsy series of FTLD-TDP-type C. <i>Brain</i> , 2022, 145, 1069-1078.	7.6	12
5	Tau interactome maps synaptic and mitochondrial processes associated with neurodegeneration. <i>Cell</i> , 2022, 185, 712-728.e14.	28.9	114
6	Uninterrupted CAG repeat drives striatum-selective transcriptionopathy and nuclear pathogenesis in human Huntingtin BAC mice. <i>Neuron</i> , 2022, 110, 1173-1192.e7.	8.1	30
7	Genome-wide analysis provides genetic evidence that ACE2 influences COVID-19 risk and yields risk scores associated with severe disease. <i>Nature Genetics</i> , 2022, 54, 382-392.	21.4	97
8	Comprehensive cross-sectional and longitudinal analyses of plasma neurofilament light across FTD spectrum disorders. <i>Cell Reports Medicine</i> , 2022, 3, 100607.	6.5	21
9	Population-scale analysis of common and rare genetic variation associated with hearing loss in adults. <i>Communications Biology</i> , 2022, 5, .	4.4	12
10	Brain volumetric deficits in <i>MAPT</i> mutation carriers: a multisite study. <i>Annals of Clinical and Translational Neurology</i> , 2021, 8, 95-110.	3.7	21
11	Synaptic processes and immune-related pathways implicated in Tourette syndrome. <i>Translational Psychiatry</i> , 2021, 11, 56.	4.8	31
12	Selective axonal translation of the mRNA isoform encoding prenylated Cdc42 supports axon growth. <i>Journal of Cell Science</i> , 2021, 134, .	2.0	16
13	Plasma Neurofilament Light for Prediction of Disease Progression in Familial Frontotemporal Lobar Degeneration. <i>Neurology</i> , 2021, 96, e2296-e2312.	1.1	52
14	Specific and behaviorally consequential astrocyte Gq GPCR signaling attenuation in vivo with $\hat{i}^2$ ARK. <i>Neuron</i> , 2021, 109, 2256-2274.e9.	8.1	47
15	Advancing human genetics research and drug discovery through exome sequencing of the UK Biobank. <i>Nature Genetics</i> , 2021, 53, 942-948.	21.4	234
16	Sequencing of 640,000 exomes identifies <i>GPR75</i> variants associated with protection from obesity. <i>Science</i> , 2021, 373, .	12.6	130
17	GADD45A is a protective modifier of neurogenic skeletal muscle atrophy. <i>JCI Insight</i> , 2021, 6, .	5.0	14
18	HDinHD: A Rich Data Portal for Huntington's Disease Research. <i>Journal of Huntington's Disease</i> , 2021, 10, 405-412.	1.9	11

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19	Molecular and functional properties of cortical astrocytes during peripherally induced neuroinflammation. <i>Cell Reports</i> , 2021, 36, 109508.	6.4	54
20	The glycine arginine-rich domain of the RNA-binding protein nucleolin regulates its subcellular localization. <i>EMBO Journal</i> , 2021, 40, e107158.	7.8	23
21	Topoisomerase I inhibition and peripheral nerve injury induce DNA breaks and ATF3-associated axon regeneration in sensory neurons. <i>Cell Reports</i> , 2021, 36, 109666.	6.4	16
22	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. <i>Molecular Psychiatry</i> , 2021, 26, 7522-7529.	7.9	8
23	Plasma Tau and Neurofilament Light in Frontotemporal Lobar Degeneration and Alzheimer Disease. <i>Neurology</i> , 2021, 96, e671-e683.	1.1	84
24	Exome sequencing and analysis of 454,787 UK Biobank participants. <i>Nature</i> , 2021, 599, 628-634.	27.8	377
25	Transcriptomic profiling of whole blood in 22q11.2 reciprocal copy number variants reveals that cell proportion highly impacts gene expression. <i>Brain, Behavior, &amp; Immunity - Health</i> , 2021, 18, 100386.	2.5	3
26	AD-linked R47H- <i>TREM2</i> mutation induces disease-enhancing microglial states via AKT hyperactivation. <i>Science Translational Medicine</i> , 2021, 13, eabe3947.	12.4	55
27	Assessment of executive function declines in presymptomatic and mildly symptomatic familial frontotemporal dementia: NIH EXAMINER as a potential clinical trial endpoint. <i>Alzheimer's and Dementia</i> , 2020, 16, 11-21.	0.8	32
28	Individualized atrophy scores predict dementia onset in familial frontotemporal lobar degeneration. <i>Alzheimer's and Dementia</i> , 2020, 16, 37-48.	0.8	38
29	Synaptic and Gene Regulatory Mechanisms in Schizophrenia, Autism, and 22q11.2 Copy Number Variant-Mediated Risk for Neuropsychiatric Disorders. <i>Biological Psychiatry</i> , 2020, 87, 150-163.	1.3	46
30	Activation of the HIF1 $\alpha$ /PFKFB3 stress response pathway in beta cells in type 1 diabetes. <i>Diabetologia</i> , 2020, 63, 149-161.	6.3	49
31	The longitudinal evaluation of familial frontotemporal dementia subjects protocol: Framework and methodology. <i>Alzheimer's and Dementia</i> , 2020, 16, 22-36.	0.8	32
32	Association of Cognitive and Behavioral Features Between Adults With Tuberous Sclerosis and Frontotemporal Dementia. <i>JAMA Neurology</i> , 2020, 77, 358.	9.0	14
33	DNA Methylation Analysis Validates Organoids as a Viable Model for Studying Human Intestinal Aging. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2020, 9, 527-541.	4.5	53
34	Clinical and volumetric changes with increasing functional impairment in familial frontotemporal lobar degeneration. <i>Alzheimer's and Dementia</i> , 2020, 16, 49-59.	0.8	27
35	A Ca <sup>2+</sup> -Dependent Switch Activates Axonal Casein Kinase 2 $\beta$ Translation and Drives G3BP1 Granule Disassembly for Axon Regeneration. <i>Current Biology</i> , 2020, 30, 4882-4895.e6.	3.9	22
36	Context-Specific Striatal Astrocyte Molecular Responses Are Phenotypically Exploitable. <i>Neuron</i> , 2020, 108, 1146-1162.e10.	8.1	73

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37	Exome sequencing and characterization of 49,960 individuals in the UK Biobank. <i>Nature</i> , 2020, 586, 749-756.	27.8	369
38	DNA methylation study of Huntington's disease and motor progression in patients and in animal models. <i>Nature Communications</i> , 2020, 11, 4529.	12.8	45
39	Lack of Association Between the CCR5-delta32 Polymorphism and Neurodegenerative Disorders. <i>Alzheimer Disease and Associated Disorders</i> , 2020, 34, 244-247.	1.3	11
40	Temporal variant of frontotemporal dementia in C9orf72 repeat expansion carriers: two case studies. <i>Brain Imaging and Behavior</i> , 2020, 14, 336-345.	2.1	3
41	Contribution of common and rare variants to bipolar disorder susceptibility in extended pedigrees from population isolates. <i>Translational Psychiatry</i> , 2020, 10, 74.	4.8	25
42	Immunosuppressive effect and global dysregulation of blood transcriptome in response to psychosocial stress in vervet monkeys ( <i>Chlorocebus sabaeus</i> ). <i>Scientific Reports</i> , 2020, 10, 3459.	3.3	2
43	DYNLRB1 is essential for dynein mediated transport and neuronal survival. <i>Neurobiology of Disease</i> , 2020, 140, 104816.	4.4	15
44	Genetic screening of a large series of North American sporadic and familial frontotemporal dementia cases. <i>Alzheimer's and Dementia</i> , 2020, 16, 118-130.	0.8	43
45	Comparison of sporadic and familial behavioral variant frontotemporal dementia (FTD) in a North American cohort. <i>Alzheimer's and Dementia</i> , 2020, 16, 60-70.	0.8	27
46	Utility of the global CDR <sup>®</sup> plus NACC FTD rating and development of scoring rules: Data from the ARTFL/LEFFTDS Consortium. <i>Alzheimer's and Dementia</i> , 2020, 16, 106-117.	0.8	81
47	Injured adult neurons regress to an embryonic transcriptional growth state. <i>Nature</i> , 2020, 581, 77-82.	27.8	154
48	Revised Self-Monitoring Scale. <i>Neurology</i> , 2020, 94, e2384-e2395.	1.1	23
49	Altered expression of genes regulating inflammation and synaptogenesis during regrowth of afferent neurons to cochlear hair cells. <i>PLoS ONE</i> , 2020, 15, e0238578.	2.5	7
50	Title is missing!. , 2020, 15, e0238578.		0
51	Title is missing!. , 2020, 15, e0238578.		0
52	Title is missing!. , 2020, 15, e0238578.		0
53	Title is missing!. , 2020, 15, e0238578.		0
54	Age- and stress-associated <i>C. elegans</i> granulins impair lysosomal function and induce a compensatory HLH-30/TFEB transcriptional response. <i>PLoS Genetics</i> , 2019, 15, e1008295.	3.5	23

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55	Neurodegenerative Disease Caregiversâ€™ 5-HTTLPR Genotype Moderates the Effect of Patientsâ€™ Empathic Accuracy Deficits on Caregiversâ€™ Well-Being. <i>American Journal of Geriatric Psychiatry</i> , 2019, 27, 1046-1056.	1.2	10
56	Patient-Tailored, Connectivity-Based Forecasts of Spreading Brain Atrophy. <i>Neuron</i> , 2019, 104, 856-868.e5.	8.1	85
57	Astrocyte molecular signatures in Huntingtonâ€™s disease. <i>Science Translational Medicine</i> , 2019, 11, .	12.4	152
58	Time Course of Changes in Peripheral Blood Gene Expression During Medication Treatment for Major Depressive Disorder. <i>Frontiers in Genetics</i> , 2019, 10, 870.	2.3	5
59	Preferential tau aggregation in von Economo neurons and fork cells in frontotemporal lobar degeneration with specific MAPT variants. <i>Acta Neuropathologica Communications</i> , 2019, 7, 159.	5.2	34
60	A Rare Mutation of Î²1-Adrenergic Receptor Affects Sleep/Wake Behaviors. <i>Neuron</i> , 2019, 103, 1044-1055.e7.	8.1	54
61	Longitudinal RNA-Seq analysis of acute and chronic neurogenic skeletal muscle atrophy. <i>Scientific Data</i> , 2019, 6, 179.	5.3	15
62	Tracking white matter degeneration in asymptomatic and symptomatic MAPT mutation carriers. <i>Neurobiology of Aging</i> , 2019, 83, 54-62.	3.1	14
63	Genetic screen in a large series of patients with primary progressive aphasia. <i>Alzheimer's and Dementia</i> , 2019, 15, 553-560.	0.8	30
64	18F-flortaucipir (AV-1451) tau PET in frontotemporal dementia syndromes. <i>Alzheimer's Research and Therapy</i> , 2019, 11, 13.	6.2	121
65	Gyrification abnormalities in presymptomatic <i>c9orf72</i> expansion carriers. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2019, 90, 1005-1010.	1.9	24
66	Partial inhibition of the overactivated Ku80-dependent DNA repair pathway rescues neurodegeneration in <i>C9ORF72</i> -ALS/FTD. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 9628-9633.	7.1	64
67	Hyperactivity with Disrupted Attention by Activation of an Astrocyte Synaptogenic Cue. <i>Cell</i> , 2019, 177, 1280-1292.e20.	28.9	228
68	Dopamine receptor D4 (DRD) polymorphisms with reduced functional potency intensify atrophy in syndrome-specific sites of frontotemporal dementia. <i>NeuroImage: Clinical</i> , 2019, 23, 101822.	2.7	4
69	Genome-wide survey of copy number variants finds MAPT duplications in progressive supranuclear palsy. <i>Movement Disorders</i> , 2019, 34, 1049-1059.	3.9	24
70	Primary familial brain calcification caused by a novel homozygous MYORG mutation in a consanguineous Italian family. <i>Neurogenetics</i> , 2019, 20, 99-102.	1.4	14
71	Interrogating the Genetic Determinants of Touretteâ€™s Syndrome and Other Tic Disorders Through Genome-Wide Association Studies. <i>American Journal of Psychiatry</i> , 2019, 176, 217-227.	7.2	242
72	Thalamo-cortical network hyperconnectivity in preclinical progranulin mutation carriers. <i>NeuroImage: Clinical</i> , 2019, 22, 101751.	2.7	30

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73	Regeneration Enhances Metastasis: A Novel Role for Neurovascular Signaling in Promoting Melanoma Brain Metastasis. <i>Frontiers in Neuroscience</i> , 2019, 13, 297.	2.8	14
74	miRNA expression profiles and molecular networks in resting and LPS-activated BV-2 microglia—Effect of cannabinoids. <i>PLoS ONE</i> , 2019, 14, e0212039.	2.5	41
75	White Matter Stroke Induces a Unique Oligo-Astrocyte Niche That Inhibits Recovery. <i>Journal of Neuroscience</i> , 2019, 39, 9343-9359.	3.6	29
76	ForestQC: Quality control on genetic variants from next-generation sequencing data using random forest. <i>PLoS Computational Biology</i> , 2019, 15, e1007556.	3.2	17
77	Frequency of the TREM2 R47H Variant in Various Neurodegenerative Disorders. <i>Alzheimer Disease and Associated Disorders</i> , 2019, 33, 327-330.	1.3	6
78	Gene-environment regulatory circuits of right ventricular pathology in tetralogy of fallot. <i>Journal of Molecular Medicine</i> , 2019, 97, 1711-1722.	3.9	11
79	Nonlinear Z-score modeling for improved detection of cognitive abnormality. <i>Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring</i> , 2019, 11, 797-808.	2.4	12
80	Transcriptional profiling of isogenic Friedreich ataxia neurons and effect of an HDAC inhibitor on disease signatures. <i>Journal of Biological Chemistry</i> , 2019, 294, 1846-1859.	3.4	28
81	Frontotemporal dementia spectrum: first genetic screen in a Greek cohort. <i>Neurobiology of Aging</i> , 2019, 75, 224.e1-224.e8.	3.1	16
82	Rare variants in the neuronal ceroid lipofuscinosis gene MFSD8 are candidate risk factors for frontotemporal dementia. <i>Acta Neuropathologica</i> , 2019, 137, 71-88.	7.7	29
83	Neurons selectively targeted in frontotemporal dementia reveal early stage TDP-43 pathobiology. <i>Acta Neuropathologica</i> , 2019, 137, 27-46.	7.7	87
84	Elevated TREM2 Gene Dosage Reprograms Microglia Responsivity and Ameliorates Pathological Phenotypes in Alzheimer's Disease Models. <i>Neuron</i> , 2018, 97, 1032-1048.e5.	8.1	246
85	Neurodegenerative disease biomarkers A $\beta$ <sub>40</sub> , A $\beta$ <sub>42</sub> , tau, and p-tau <sub>181</sub> in the vervet monkey cerebrospinal fluid: Relation to normal aging, genetic influences, and cerebral amyloid angiopathy. <i>Brain and Behavior</i> , 2018, 8, e00903.	2.2	45
86	Poly(GP), neurofilament and grey matter deficits in <i>C9orf72</i> expansion carriers. <i>Annals of Clinical and Translational Neurology</i> , 2018, 5, 583-597.	3.7	48
87	Benign hereditary chorea and deletions outside NKX2-1: What's the role of MBIP?. <i>European Journal of Medical Genetics</i> , 2018, 61, 581-584.	1.3	9
88	Expanding the global prevalence of spinocerebellar ataxia type 42. <i>Neurology: Genetics</i> , 2018, 4, e232.	1.9	14
89	Impairment of memory generalization in preclinical autosomal dominant Alzheimer's disease mutation carriers. <i>Neurobiology of Aging</i> , 2018, 65, 149-157.	3.1	7
90	Mechanistic Differences in Neuropathic Pain Modalities Revealed by Correlating Behavior with Global Expression Profiling. <i>Cell Reports</i> , 2018, 22, 1301-1312.	6.4	142

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91	The Longitudinal Trajectory of Default Mode Network Connectivity in Healthy Older Adults Varies As a Function of Age and Is Associated with Changes in Episodic Memory and Processing Speed. <i>Journal of Neuroscience</i> , 2018, 38, 2809-2817.	3.6	161
92	A molecular cascade modulates MAP1B and confers resistance to mTOR inhibition in human glioblastoma. <i>Neuro-Oncology</i> , 2018, 20, 764-775.	1.2	22
93	Progranulin levels in blood in Alzheimer's disease and mild cognitive impairment. <i>Annals of Clinical and Translational Neurology</i> , 2018, 5, 616-629.	3.7	23
94	Combined Pathologies in FTLT-DTP Types A and C. <i>Journal of Neuropathology and Experimental Neurology</i> , 2018, 77, 405-412.	1.7	8
95	P1â€433: GRAY MATTER DEFICITS IN SYMPTOMATIC AND PRESYMPTOMATIC <i>MAPT</i> MUTATION CARRIERS. <i>Alzheimer's and Dementia</i> , 2018, 14, P475.	0.8	0
96	O2â€14â€06: DIFFERENCES BETWEEN SPORADIC AND FAMILIAL BEHAVIORAL VARIANT FTD IN ADVANCING RESEARCH AND TREATMENT FOR FTD (ARTFL) CLINICAL RESEARCH CONSORTIUM. <i>Alzheimer's and Dementia</i> , 2018, 14, P658.	0.8	0
97	P1â€281: NONLINEAR Nâ€SCORE ESTIMATION FOR ESTABLISHING COGNITIVE NORMS FROM THE NATIONAL ALZHEIMER'S COORDINATING CENTER (NACC) DATASET. <i>Alzheimer's and Dementia</i> , 2018, 14, P390.	0.8	1
98	CSIG-22. RECONCILING TUMOR HETEROGENEITY IN GLIOBLASTOMA USING A PATHWAY-BASED APPROACH. <i>Neuro-Oncology</i> , 2018, 20, vi47-vi47.	1.2	0
99	Frequency of frontotemporal dementia gene variants in &em&gt;C9ORF72&/em&gt;, &em&gt;MAPT&/em&gt;, and &em&gt;GRN&/em&gt; in academic versus commercial laboratory cohorts. <i>Advances in Genomics and Genetics</i> , 2018, Volume 8, 23-33.	0.8	7
100	O1â€08â€01: THE NIHâ€EXAMINER IS SENSITIVE TO COGNITIVE CHANGES IN ASYMPTOMATIC AND MILDLY SYMPTOMATIC FAMILIAL FRONTOTEMPORAL DEMENTIA. <i>Alzheimer's and Dementia</i> , 2018, 14, P235.	0.8	0
101	O2â€14â€01: CHARACTERISTICS AND PROGRESS OF 320 SUBJECTS IN THE LONGITUDINAL EVALUATION OF FAMILIAL FRONTOTEMPORAL DEMENTIA SUBJECTS (LEFFTDs) PROTOCOL. <i>Alzheimer's and Dementia</i> , 2018, 14, P656.	0.8	0
102	P1â€419: USING A BRAIN NETWORK APPROACH TO PREDICT GENETIC MUTATION IN INDIVIDUAL PATIENTS WITH FAMILIAL FRONTOTEMPORAL DEMENTIA. <i>Alzheimer's and Dementia</i> , 2018, 14, P465.	0.8	0
103	Nâ€acetylcysteine targets 5 lipoxygenaseâ€derived, toxic lipids and can synergize with prostaglandin E<sub>2</sub> to inhibit ferroptosis and improve outcomes following hemorrhagic stroke in mice. <i>Annals of Neurology</i> , 2018, 84, 854-872.	5.3	195
104	Differential effects of partial and complete loss of TREM2 on microglial injury response and tauopathy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10172-10177.	7.1	118
105	De Novo Sequence and Copy Number Variants Are Strongly Associated with Tourette Disorder and Implicate Cell Polarity in Pathogenesis. <i>Cell Reports</i> , 2018, 24, 3441-3454.e12.	6.4	91
106	Mixed TDP-43 proteinopathy and tauopathy in frontotemporal lobar degeneration: nine case series. <i>Journal of Neurology</i> , 2018, 265, 2960-2971.	3.6	17
107	Required growth facilitators propel axon regeneration across complete spinal cord injury. <i>Nature</i> , 2018, 561, 396-400.	27.8	341
108	Reducing Astrocyte Calcium Signaling InÂVivo Alters Striatal Microcircuits and Causes Repetitive Behavior. <i>Neuron</i> , 2018, 99, 1170-1187.e9.	8.1	234

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109	Adult rat myelin enhances axonal outgrowth from neural stem cells. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	28
110	Peripheral blood gene expression reveals an inflammatory transcriptomic signature in Friedreichâ€™s ataxia patients. <i>Human Molecular Genetics</i> , 2018, 27, 2965-2977.	2.9	36
111	Primary brain calcification: an international study reporting novel variants and associated phenotypes. <i>European Journal of Human Genetics</i> , 2018, 26, 1462-1477.	2.8	48
112	Generation and post-injury integration of human spinal cord neural stem cells. <i>Nature Methods</i> , 2018, 15, 723-731.	19.0	132
113	hnRNPs Interacting with mRNA Localization Motifs Define Axonal RNA Regulons. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 2091-2106.	3.8	32
114	Metabolic characterization of isocitrate dehydrogenase (IDH) mutant and IDH wildtype gliomaspheres uncovers cell type-specific vulnerabilities. <i>Cancer &amp; Metabolism</i> , 2018, 6, 4.	5.0	55
115	In the setting of $\beta$ -cell stress, the pancreatic duct gland transcriptome shows characteristics of an activated regenerative response. <i>American Journal of Physiology - Renal Physiology</i> , 2018, 315, G848-G854.	3.4	4
116	Joint genome-wide association study of progressive supranuclear palsy identifies novel susceptibility loci and genetic correlation to neurodegenerative diseases. <i>Molecular Neurodegeneration</i> , 2018, 13, 41.	10.8	77
117	Bioinformatics and genomic databases. <i>Handbook of Clinical Neurology</i> / Edited By P J Vinken and G W Bruyn, 2018, 147, 75-92.	1.8	15
118	CREB controls cortical circuit plasticity and functional recovery after stroke. <i>Nature Communications</i> , 2018, 9, 2250.	12.8	96
119	MicroRNA signatures of endogenous Huntingtin CAG repeat expansion in mice. <i>PLoS ONE</i> , 2018, 13, e0190550.	2.5	39
120	Translatome Regulation in Neuronal Injury and Axon Regrowth. <i>ENeuro</i> , 2018, 5, ENEURO.0276-17.2018.	1.9	26
121	Genome-wide association study identifies <i>MAPT</i> locus influencing human plasma tau levels. <i>Neurology</i> , 2017, 88, 669-676.	1.1	33
122	Precipitous Deterioration of Motor Function, Cognition, and Behavior. <i>JAMA Neurology</i> , 2017, 74, 591.	9.0	0
123	De Novo Coding Variants Are Strongly Associated with Tourette Disorder. <i>Neuron</i> , 2017, 94, 486-499.e9.	8.1	155
124	Sox11 Expression Promotes Regeneration of Some Retinal Ganglion Cell Types but Kills Others. <i>Neuron</i> , 2017, 94, 1112-1120.e4.	8.1	151
125	Rare Copy Number Variants in NRXN1 and CNTN6 Increase Risk for Tourette Syndrome. <i>Neuron</i> , 2017, 94, 1101-1111.e7.	8.1	137
126	Network degeneration and dysfunction in presymptomatic C9ORF72 expansion carriers. <i>NeuroImage: Clinical</i> , 2017, 14, 286-297.	2.7	129



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127	Ancient hybridization and strong adaptation to viruses across African vervet monkey populations. <i>Nature Genetics</i> , 2017, 49, 1705-1713.	21.4	107
128	Genetic variation and gene expression across multiple tissues and developmental stages in a nonhuman primate. <i>Nature Genetics</i> , 2017, 49, 1714-1721.	21.4	57
129	Self-Organized Cerebral Organoids with Human-Specific Features Predict Effective Drugs to Combat Zika Virus Infection. <i>Cell Reports</i> , 2017, 21, 517-532.	6.4	305
130	Clinicopathological correlations in behavioural variant frontotemporal dementia. <i>Brain</i> , 2017, 140, 3329-3345.	7.6	226
131	Linking tuberous sclerosis complex, excessive mTOR signaling, and age-related neurodegeneration: a new association between TSC1 mutation and frontotemporal dementia. <i>Acta Neuropathologica</i> , 2017, 134, 813-816.	7.7	11
132	Timing of Smarcb1 and Nf2 inactivation determines schwannoma versus rhabdoid tumor development. <i>Nature Communications</i> , 2017, 8, 300.	12.8	70
133	Neural Circuit-Specialized Astrocytes: Transcriptomic, Proteomic, Morphological, and Functional Evidence. <i>Neuron</i> , 2017, 95, 531-549.e9.	8.1	556
134	Enhanced Neuronal Regeneration in the CAST/Ei Mouse Strain Is Linked to Expression of Differentiation Markers after Injury. <i>Cell Reports</i> , 2017, 20, 1136-1147.	6.4	26
135	Activity-Dependent Regulation of Alternative Cleavage and Polyadenylation During Hippocampal Long-Term Potentiation. <i>Scientific Reports</i> , 2017, 7, 17377.	3.3	38
136	A152T tau allele causes neurodegeneration that can be ameliorated in a zebrafish model by autophagy induction. <i>Brain</i> , 2017, 140, 1128-1146.	7.6	84
137	Brain calcifications and <i>PCDH12</i> variants. <i>Neurology: Genetics</i> , 2017, 3, e166.	1.9	15
138	Mapping Gene Expression in Excitatory Neurons during Hippocampal Late-Phase Long-Term Potentiation. <i>Frontiers in Molecular Neuroscience</i> , 2017, 10, 39.	2.9	49
139	Wnt11 regulates cardiac chamber development and disease during perinatal maturation. <i>JCI Insight</i> , 2017, 2, .	5.0	21
140	Fine-mapping of the human leukocyte antigen locus as a risk factor for Alzheimer disease: A case-control study. <i>PLoS Medicine</i> , 2017, 14, e1002272.	8.4	67
141	Huntington's disease accelerates epigenetic aging of human brain and disrupts DNA methylation levels. <i>Aging</i> , 2016, 8, 1485-1512.	3.1	192
142	Forward Genetic Screen in <i>Caenorhabditis elegans</i> Suggests F57A10.2 and <i>acp-4</i> As Suppressors of C9ORF72 Related Phenotypes. <i>Frontiers in Molecular Neuroscience</i> , 2016, 9, 113.	2.9	16
143	Characterization of Expression Quantitative Trait Loci in Pedigrees from Colombia and Costa Rica Ascertained for Bipolar Disorder. <i>PLoS Genetics</i> , 2016, 12, e1006046.	3.5	4
144	Identification of an Efficient Gene Expression Panel for Glioblastoma Classification. <i>PLoS ONE</i> , 2016, 11, e0164649.	2.5	12

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145	Cardiac Dysfunction in the BACHD Mouse Model of Huntington's Disease. PLoS ONE, 2016, 11, e0147269.	2.5	30
146	A Novel Protocol for Directed Differentiation of C9orf72-Associated Human Induced Pluripotent Stem Cells Into Contractile Skeletal Myotubes. Stem Cells Translational Medicine, 2016, 5, 1461-1472.	3.3	38
147	Genetic Prion Disease Caused by PRNP Q160X Mutation Presenting with an Orbitofrontal Syndrome, Cyclic Diarrhea, and Peripheral Neuropathy. Journal of Alzheimer's Disease, 2016, 55, 249-258.	2.6	13
148	New Transgenic Mouse Lines for Selectively Targeting Astrocytes and Studying Calcium Signals in Astrocyte Processes In Situ and In Vivo. Neuron, 2016, 92, 1181-1195.	8.1	283
149	Astrocyte scar formation aids central nervous system axon regeneration. Nature, 2016, 532, 195-200.	27.8	1,390
150	Inhibition of Nucleotide Synthesis Targets Brain Tumor Stem Cells in a Subset of Glioblastoma. Molecular Cancer Therapeutics, 2016, 15, 1271-1278.	4.1	13
151	Progranulin Deficiency Promotes Circuit-Specific Synaptic Pruning by Microglia via Complement Activation. Cell, 2016, 165, 921-935.	28.9	558
152	Large-scale assessment of the gliomasphere model system. Neuro-Oncology, 2016, 18, 1367-1378.	1.2	82
153	Decoding the Long Noncoding RNA During Cardiac Maturation. Circulation: Cardiovascular Genetics, 2016, 9, 395-407.	5.1	39
154	Pathways and gene networks mediating the regulatory effects of cannabidiol, a nonpsychoactive cannabinoid, in autoimmune T cells. Journal of Neuroinflammation, 2016, 13, 136.	7.2	59
155	Widespread white matter and conduction defects in PSEN1-related spastic paraparesis. Neurobiology of Aging, 2016, 47, 201-209.	3.1	17
156	Incidence and impact of subclinical epileptiform activity in Alzheimer's disease. Annals of Neurology, 2016, 80, 858-870.	5.3	373
157	Human iPSC-Derived Neuronal Model of Tau-A152T Frontotemporal Dementia Reveals Tau-Mediated Mechanisms of Neuronal Vulnerability. Stem Cell Reports, 2016, 7, 325-340.	4.8	92
158	Design, Characterization, and Use of a Novel Amyloid $\beta$ -Protein Control for Assembly, Neurotoxicity, and Gene Expression Studies. Biochemistry, 2016, 55, 5049-5060.	2.5	5
159	Increased prevalence of autoimmune disease within C9 and FTD/MND cohorts. Neurology: Neuroimmunology and Neuroinflammation, 2016, 3, e301.	6.0	78
160	Distinct Subtypes of Behavioral Variant Frontotemporal Dementia Based on Patterns of Network Degeneration. JAMA Neurology, 2016, 73, 1078.	9.0	115
161	Therapeutic targeting of oxygen-sensing prolyl hydroxylases abrogates ATF4-dependent neuronal death and improves outcomes after brain hemorrhage in several rodent models. Science Translational Medicine, 2016, 8, 328ra29.	12.4	106
162	Timing and significance of pathological features in C9orf72 expansion-associated frontotemporal dementia. Brain, 2016, 139, 3202-3216.	7.6	136

#	ARTICLE	IF	CITATIONS
163	Rare TREM2 variants associated with Alzheimer's disease display reduced cell surface expression. <i>Acta Neuropathologica Communications</i> , 2016, 4, 98.	5.2	46
164	Suberoylanilide hydroxamic acid increases progranulin production in iPSC-derived cortical neurons of frontotemporal dementia patients. <i>Neurobiology of Aging</i> , 2016, 42, 35-40.	3.1	18
165	Co-expression networks in generation of induced pluripotent stem cells. <i>Biology Open</i> , 2016, 5, 300-310.	1.2	3
166	Transcriptomics and the mechanisms of antidepressant efficacy. <i>European Neuropsychopharmacology</i> , 2016, 26, 105-112.	0.7	19
167	Differential regulation of type III secretion and virulence genes in <i>Bordetella pertussis</i> and <i>Bordetella bronchiseptica</i> by a secreted anti-If factor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 2341-2348.	7.1	54
168	Neuropathology of Autosomal Dominant Alzheimer Disease in the National Alzheimer Coordinating Center Database. <i>Journal of Neuropathology and Experimental Neurology</i> , 2016, 75, 284-290.	1.7	71
169	A Systems-Level Analysis of the Peripheral Nerve Intrinsic Axonal Growth Program. <i>Neuron</i> , 2016, 89, 956-970.	8.1	314
170	Integrated genomics and proteomics define huntingtin CAG length-dependent networks in mice. <i>Nature Neuroscience</i> , 2016, 19, 623-633.	14.8	342
171	Primary familial brain calcification in a Norwegian family, caused by a novel SLC20A2 gene mutation. <i>Journal of Neurology</i> , 2016, 263, 594-596.	3.6	2
172	C9orf72 repeat expansions that cause frontotemporal dementia are detectable among patients with psychosis. <i>Psychiatry Research</i> , 2016, 235, 200-202.	3.3	22
173	Common variants in ABCA7 and MS4A6A are associated with cortical and hippocampal atrophy. <i>Neurobiology of Aging</i> , 2016, 39, 82-89.	3.1	40
174	Early-onset Alzheimer's disease versus frontotemporal dementia: resolution with genetic diagnoses?. <i>Neurocase</i> , 2016, 22, 161-167.	0.6	3
175	Amyloid in dementia associated with familial FTL: not an innocent bystander. <i>Neurocase</i> , 2016, 22, 76-83.	0.6	12
176	The 5-HTTLPR variant in the serotonin transporter gene modifies degeneration of brain regions important for emotion in behavioral variant frontotemporal dementia. <i>NeuroImage: Clinical</i> , 2015, 9, 283-290.	2.7	7
177	No Evidence to Suggest that the Use of Acetylcholinesterase Inhibitors Confounds the Results of Two Blood-Based Biomarker Studies in Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2015, 47, 741-750.	2.6	2
178	Genomic Analysis Reveals Disruption of Striatal Neuronal Development and Therapeutic Targets in Human Huntington's Disease Neural Stem Cells. <i>Stem Cell Reports</i> , 2015, 5, 1023-1038.	4.8	117
179	Brain calcification process and phenotypes according to age and sex: Lessons from SLC20A2, PDGFB, and PDGFRB mutation carriers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 586-594.	1.7	74
180	Apolipoprotein $\mu$ 4 Is Associated with Lower Brain Volume in Cognitively Normal Chinese but Not White Older Adults. <i>PLoS ONE</i> , 2015, 10, e0118338.	2.5	12

#	ARTICLE	IF	CITATIONS
181	Transcriptome Profiling of Peripheral Blood in 22q11.2 Deletion Syndrome Reveals Functional Pathways Related to Psychosis and Autism Spectrum Disorder. <i>PLoS ONE</i> , 2015, 10, e0132542.	2.5	40
182	Robust Axonal Regeneration Occurs in the Injured CAST/Ei Mouse CNS. <i>Neuron</i> , 2015, 86, 1215-1227.	8.1	87
183	Clinicopathological Study of Patients With <i>C9ORF72</i> -Associated Frontotemporal Dementia Presenting With Delusions. <i>Journal of Geriatric Psychiatry and Neurology</i> , 2015, 28, 99-107.	2.3	41
184	N17 Modifies Mutant Huntingtin Nuclear Pathogenesis and Severity of Disease in HD BAC Transgenic Mice. <i>Neuron</i> , 2015, 85, 726-741.	8.1	66
185	SIRT1 Deficiency in Microglia Contributes to Cognitive Decline in Aging and Neurodegeneration via Epigenetic Regulation of IL-1 $\beta$ . <i>Journal of Neuroscience</i> , 2015, 35, 807-818.	3.6	212
186	A Multiancestral Genome-Wide Exome Array Study of Alzheimer Disease, Frontotemporal Dementia, and Progressive Supranuclear Palsy. <i>JAMA Neurology</i> , 2015, 72, 414.	9.0	37
187	Update and Mutational Analysis of <i>SLC20A2</i> : A Major Cause of Primary Familial Brain Calcification. <i>Human Mutation</i> , 2015, 36, 489-495.	2.5	80
188	A new model to study neurodegeneration in ataxia oculomotor apraxia type 2. <i>Human Molecular Genetics</i> , 2015, 24, 5759-5774.	2.9	34
189	The choroid plexus transcriptome reveals changes in type I and II interferon responses in a mouse model of Alzheimer's disease. <i>Brain, Behavior, and Immunity</i> , 2015, 49, 280-292.	4.1	60
190	Tet3 regulates synaptic transmission and homeostatic plasticity via DNA oxidation and repair. <i>Nature Neuroscience</i> , 2015, 18, 836-843.	14.8	164
191	Novel candidate blood-based transcriptional biomarkers of Machado-Joseph disease. <i>Movement Disorders</i> , 2015, 30, 968-975.	3.9	28
192	Cannabidiol, a non-psychoactive cannabinoid, leads to EGR2-dependent energy in activated encephalitogenic T cells. <i>Journal of Neuroinflammation</i> , 2015, 12, 52.	7.2	37
193	Mutations in XPR1 cause primary familial brain calcification associated with altered phosphate export. <i>Nature Genetics</i> , 2015, 47, 579-581.	21.4	237
194	GDF10 is a signal for axonal sprouting and functional recovery after stroke. <i>Nature Neuroscience</i> , 2015, 18, 1737-1745.	14.8	144
195	First Japanese family with primary familial brain calcification due to a mutation in the <i>PDGFB</i> gene: An exome analysis study. <i>Psychiatry and Clinical Neurosciences</i> , 2015, 69, 77-83.	1.8	22
196	Decision tree analysis of genetic risk for clinically heterogeneous Alzheimer's disease. <i>BMC Neurology</i> , 2015, 15, 47.	1.8	25
197	Loss of functional connectivity is greater outside the default mode network in nonfamilial early-onset Alzheimer's disease variants. <i>Neurobiology of Aging</i> , 2015, 36, 2678-2686.	3.1	72
198	Whole-genome sequencing suggests a chemokine gene cluster that modifies age at onset in familial Alzheimer's disease. <i>Molecular Psychiatry</i> , 2015, 20, 1294-1300.	7.9	55

#	ARTICLE	IF	CITATIONS
199	Divergent CSF A $\beta$ alterations in two common tauopathies: Alzheimer's disease and progressive supranuclear palsy. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2015, 86, 244-250.	1.9	101
200	Increased CYFIP1 dosage alters cellular and dendritic morphology and dysregulates mTOR. <i>Molecular Psychiatry</i> , 2015, 20, 1069-1078.	7.9	98
201	Hydroxamic Acid-Based Histone Deacetylase (HDAC) Inhibitors Can Mediate Neuroprotection Independent of HDAC Inhibition. <i>Journal of Neuroscience</i> , 2014, 34, 14328-14337.	3.6	25
202	Novel Roles for Osteopontin and Clusterin in Peripheral Motor and Sensory Axon Regeneration. <i>Journal of Neuroscience</i> , 2014, 34, 1689-1700.	3.6	61
203	An Epigenetic Signature in Peripheral Blood Associated with the Haplotype on 17q21.31, a Risk Factor for Neurodegenerative Tauopathy. <i>PLoS Genetics</i> , 2014, 10, e1004211.	3.5	65
204	Altered network connectivity in frontotemporal dementia with C9orf72 hexanucleotide repeat expansion. <i>Brain</i> , 2014, 137, 3047-3060.	7.6	140
205	Cerebrospinal fluid neurofilament concentration reflects disease severity in frontotemporal degeneration. <i>Annals of Neurology</i> , 2014, 75, 116-126.	5.3	213
206	A Novel PSEN1 Mutation (I238M) associated with Early-Onset Alzheimer's Disease in an African-American Woman. <i>Journal of Alzheimer's Disease</i> , 2014, 40, 271-275.	2.6	14
207	Mutation of senataxin alters disease-specific transcriptional networks in patients with ataxia with oculomotor apraxia type 2. <i>Human Molecular Genetics</i> , 2014, 23, 4758-4769.	2.9	43
208	Life Extension Factor Klotho Enhances Cognition. <i>Cell Reports</i> , 2014, 7, 1065-1076.	6.4	243
209	Chronic administration of cholesterol oximes in mice increases transcription of cytoprotective genes and improves transcriptome alterations induced by alpha-synuclein overexpression in nigrostriatal dopaminergic neurons. <i>Neurobiology of Disease</i> , 2014, 69, 263-275.	4.4	28
210	Abstract A10: Suppressor of cytokine signaling (SOCS)-3 and the C-X-C chemokines CXCL1 and CXCL2 promote tumor aggressiveness and radiation resistance in pancreatic cancer. , 2014, , .		0
211	Mutations in the gene encoding PDGF-B cause brain calcifications in humans and mice. <i>Nature Genetics</i> , 2013, 45, 1077-1082.	21.4	273
212	Mutations in SLC20A2 are a major cause of familial idiopathic basal ganglia calcification. <i>Neurogenetics</i> , 2013, 14, 11-22.	1.4	131
213	Regional brain volume differences in symptomatic and presymptomatic carriers of familial Alzheimer's disease mutations. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2013, 84, 154-162.	1.9	47
214	<i>C9ORF72</i> repeat expansions in cases with previously identified pathogenic mutations. <i>Neurology</i> , 2013, 81, 1332-1341.	1.1	84
215	The effect of the serotonin transporter polymorphism (5-HTTLPR) on empathic and self-conscious emotional reactivity.. <i>Emotion</i> , 2013, 13, 25-35.	1.8	65
216	Progranulin Mutations as Risk Factors for Alzheimer Disease. <i>JAMA Neurology</i> , 2013, 70, 774.	9.0	114

#	ARTICLE	IF	CITATIONS
217	Microarray and Pathway Analysis Reveal Distinct Mechanisms Underlying Cannabinoid-Mediated Modulation of LPS-Induced Activation of BV-2 Microglial Cells. <i>PLoS ONE</i> , 2013, 8, e61462.	2.5	94
218	Evidence for a role of the rare p.A152T variant in MAPT in increasing the risk for FTD-spectrum and Alzheimer's diseases. <i>Human Molecular Genetics</i> , 2012, 21, 3500-3512.	2.9	198
219	Frontotemporal dementia due to <i>C9ORF72</i> mutations. <i>Neurology</i> , 2012, 79, 1002-1011.	1.1	183
220	Mitochondrial Dysfunction and Immune Activation are Detectable in Early Alzheimer's Disease Blood. <i>Journal of Alzheimer's Disease</i> , 2012, 30, 685-710.	2.6	141
221	Inflammatory Mediators Alter the Astrocyte Transcriptome and Calcium Signaling Elicited by Multiple G-Protein-Coupled Receptors. <i>Journal of Neuroscience</i> , 2012, 32, 14489-14510.	3.6	178
222	Subcellular Knockout of Importin $\beta 1$ Perturbs Axonal Retrograde Signaling. <i>Neuron</i> , 2012, 75, 294-305.	8.1	180
223	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012, 44, 552-561.	21.4	594
224	Network Organization of the Huntingtin Proteomic Interactome in Mammalian Brain. <i>Neuron</i> , 2012, 75, 41-57.	8.1	262
225	Designing, Performing, and Interpreting a Microarray-Based Gene Expression Study. <i>Methods in Molecular Biology</i> , 2011, 793, 417-439.	0.9	22
226	Expanded GGGGCC Hexanucleotide Repeat in Noncoding Region of C9ORF72 Causes Chromosome 9p-Linked FTD and ALS. <i>Neuron</i> , 2011, 72, 245-256.	8.1	4,176
227	A gene expression phenotype in lymphocytes from friedreich ataxia patients. <i>Annals of Neurology</i> , 2011, 70, 790-804.	5.3	58
228	Suberoylanilide Hydroxamic Acid (Vorinostat) Up-regulates Progranulin Transcription. <i>Journal of Biological Chemistry</i> , 2011, 286, 16101-16108.	3.4	138
229	Signaling to Transcription Networks in the Neuronal Retrograde Injury Response. <i>Science Signaling</i> , 2010, 3, ra53.	3.6	159
230	An age-related sprouting transcriptome provides molecular control of axonal sprouting after stroke. <i>Nature Neuroscience</i> , 2010, 13, 1496-1504.	14.8	291
231	Inosine Alters Gene Expression and Axonal Projections in Neurons Contralateral to a Cortical Infarct and Improves Skilled Use of the Impaired Limb. <i>Journal of Neuroscience</i> , 2009, 29, 8187-8197.	3.6	93
232	Functional genomic analysis of frataxin deficiency reveals tissue-specific alterations and identifies the PPAR $\beta$ pathway as a therapeutic target in Friedreich's ataxia. <i>Human Molecular Genetics</i> , 2009, 18, 2452-2461.	2.9	109
233	Functional and Evolutionary Insights into Human Brain Development through Global Transcriptome Analysis. <i>Neuron</i> , 2009, 62, 494-509.	8.1	555
234	Tauopathy with paired helical filaments in an aged chimpanzee. <i>Journal of Comparative Neurology</i> , 2008, 509, 259-270.	1.6	129

#	ARTICLE	IF	CITATIONS
235	Gene expression study on peripheral blood identifies progranulin mutations. <i>Annals of Neurology</i> , 2008, 64, 92-96.	5.3	91
236	ATF4 is an oxidative stress-inducible, prodeath transcription factor in neurons in vitro and in vivo. <i>Journal of Experimental Medicine</i> , 2008, 205, 1227-1242.	8.5	219
237	HDAC Inhibitors Correct Frataxin Deficiency in a Friedreich Ataxia Mouse Model. <i>PLoS ONE</i> , 2008, 3, e1958.	2.5	193
238	Phenotypic variability associated with progranulin haploinsufficiency in patients with the common 1477C>T (Arg493X) mutation: an international initiative. <i>Lancet Neurology</i> , The, 2007, 6, 857-868.	10.2	199
239	Microarrays and the microscope: balancing throughput with resolution. <i>Journal of Physiology</i> , 2006, 575, 353-359.	2.9	11
240	Gene expression profiling in frataxin deficient mice: Microarray evidence for significant expression changes without detectable neurodegeneration. <i>Neurobiology of Disease</i> , 2006, 22, 302-311.	4.4	50
241	Patient-Tailored, Connectivity-Based Forecasts of Spreading Brain Atrophy. <i>SSRN Electronic Journal</i> , 0, , .	0.4	1