

Christopher E Shaw

List of Publications by Year in descending order

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339
papers

50,410
citations

1704

104
h-index

1857

209
g-index

354
all docs

354
docs citations

354
times ranked

36280
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , 2009, 41, 1088-1093.	21.4	2,697
2	Mutations in FUS, an RNA Processing Protein, Cause Familial Amyotrophic Lateral Sclerosis Type 6. <i>Science</i> , 2009, 323, 1208-1211.	12.6	2,295
3	TDP-43 Mutations in Familial and Sporadic Amyotrophic Lateral Sclerosis. <i>Science</i> , 2008, 319, 1668-1672.	12.6	2,268
4	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates APOE, tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019, 51, 414-430.	21.4	1,962
5	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , 2011, 43, 429-435.	21.4	1,708
6	Mutations in prion-like domains in hnRNPA2B1 and hnRNPA1 cause multisystem proteinopathy and ALS. <i>Nature</i> , 2013, 495, 467-473.	27.8	1,249
7	Characterizing the RNA targets and position-dependent splicing regulation by TDP-43. <i>Nature Neuroscience</i> , 2011, 14, 452-458.	14.8	956
8	Effects of non-invasive ventilation on survival and quality of life in patients with amyotrophic lateral sclerosis: a randomised controlled trial. <i>Lancet Neurology</i> , 2006, 5, 140-147.	10.2	922
9	Amyotrophic lateral sclerosis. <i>Nature Reviews Disease Primers</i> , 2017, 3, 17071.	30.5	885
10	Pathological TDP-43 distinguishes sporadic amyotrophic lateral sclerosis from amyotrophic lateral sclerosis with SOD1 mutations. <i>Annals of Neurology</i> , 2007, 61, 427-434.	5.3	840
11	Exome sequencing in amyotrophic lateral sclerosis identifies risk genes and pathways. <i>Science</i> , 2015, 347, 1436-1441.	12.6	823
12	VEGF is a modifier of amyotrophic lateral sclerosis in mice and humans and protects motoneurons against ischemic death. <i>Nature Genetics</i> , 2003, 34, 383-394.	21.4	794
13	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	21.4	783
14	Evidence of widespread cerebral microglial activation in amyotrophic lateral sclerosis: an [11C](R)-PK11195 positron emission tomography study. <i>Neurobiology of Disease</i> , 2004, 15, 601-609.	4.4	630
15	Axonal Transport of TDP-43 mRNA Granules Is Impaired by ALS-Causing Mutations. <i>Neuron</i> , 2014, 81, 536-543.	8.1	521
16	Genome-wide Analyses Identify KIF5A as a Novel ALS Gene. <i>Neuron</i> , 2018, 97, 1268-1283.e6.	8.1	517
17	Variants of the elongator protein 3 (ELP3) gene are associated with motor neuron degeneration. <i>Human Molecular Genetics</i> , 2009, 18, 472-481.	2.9	512
18	Oxidative stress in ALS: Key role in motor neuron injury and therapeutic target. <i>Free Radical Biology and Medicine</i> , 2010, 48, 629-641.	2.9	512

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19	Molecular pathways of motor neuron injury in amyotrophic lateral sclerosis. <i>Nature Reviews Neurology</i> , 2011, 7, 616-630.	10.1	512
20	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016, 48, 1043-1048.	21.4	494
21	Common variants at 7p21 are associated with frontotemporal lobar degeneration with TDP-43 inclusions. <i>Nature Genetics</i> , 2010, 42, 234-239.	21.4	479
22	ERâ€™mitochondria associations are regulated by the VAPBâ€™PTPIP51 interaction and are disrupted by ALS/FTD-associated TDP-43. <i>Nature Communications</i> , 2014, 5, 3996.	12.8	463
23	Senataxin, the ortholog of a yeast RNA helicase, is mutant in ataxia-ocular apraxia 2. <i>Nature Genetics</i> , 2004, 36, 225-227.	21.4	454
24	Controversies and priorities in amyotrophic lateral sclerosis. <i>Lancet Neurology</i> , The, 2013, 12, 310-322.	10.2	454
25	Gain of Toxicity from ALS/FTD-Linked Repeat Expansions in C9ORF72 Is Alleviated by Antisense Oligonucleotides Targeting GGGGCC-Containing RNAs. <i>Neuron</i> , 2016, 90, 535-550.	8.1	437
26	p62 positive, TDP-43 negative, neuronal cytoplasmic and intranuclear inclusions in the cerebellum and hippocampus define the pathology of C9orf72-linked FTL and MND/ALS. <i>Acta Neuropathologica</i> , 2011, 122, 691-702.	7.7	432
27	VAPB interacts with the mitochondrial protein PTPIP51 to regulate calcium homeostasis. <i>Human Molecular Genetics</i> , 2012, 21, 1299-1311.	2.9	423
28	Hexanucleotide Repeats in ALS/FTD Form Length-Dependent RNA Foci, Sequester RNA Binding Proteins, and Are Neurotoxic. <i>Cell Reports</i> , 2013, 5, 1178-1186.	6.4	419
29	Novel Mutations in TARDBP (TDP-43) in Patients with Familial Amyotrophic Lateral Sclerosis. <i>PLoS Genetics</i> , 2008, 4, e1000193.	3.5	393
30	ALS-linked TDP-43 mutations produce aberrant RNA splicing and adult-onset motor neuron disease without aggregation or loss of nuclear TDP-43. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E736-45.	7.1	370
31	Autophagy induction enhances TDP43 turnover and survival in neuronal ALS models. <i>Nature Chemical Biology</i> , 2014, 10, 677-685.	8.0	368
32	Familial amyotrophic lateral sclerosis-linked SOD1 mutants perturb fast axonal transport to reduce axonal mitochondria content. <i>Human Molecular Genetics</i> , 2007, 16, 2720-2728.	2.9	365
33	Familial amyotrophic lateral sclerosis with frontotemporal dementia is linked to a locus on chromosome 9p13.2â€™21.3. <i>Brain</i> , 2006, 129, 868-876.	7.6	363
34	The role of mitochondria in amyotrophic lateral sclerosis. <i>Neuroscience Letters</i> , 2019, 710, 132933.	2.1	356
35	Genetic Evidence Implicates the Immune System and Cholesterol Metabolism in the Aetiology of Alzheimer's Disease. <i>PLoS ONE</i> , 2010, 5, e13950.	2.5	347
36	Genome-wide association study identifies 19p13.3 (UNC13A) and 9p21.2 as susceptibility loci for sporadic amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2009, 41, 1083-1087.	21.4	344

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37	Prognosis for patients with amyotrophic lateral sclerosis: development and validation of a personalised prediction model. <i>Lancet Neurology</i> , The, 2018, 17, 423-433.	10.2	342
38	The C9orf72 protein interacts with Rab1a and the <scp>ULK</scp>1 complex to regulate initiation of autophagy. <i>EMBO Journal</i> , 2016, 35, 1656-1676.	7.8	327
39	Oxidative damage to protein in sporadic motor neuron disease spinal cord. <i>Annals of Neurology</i> , 1995, 38, 691-695.	5.3	312
40	Mutant induced pluripotent stem cell lines recapitulate aspects of TDP-43 proteinopathies and reveal cell-specific vulnerability. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5803-5808.	7.1	308
41	Exome-wide Rare Variant Analysis Identifies TUBA4A Mutations Associated with Familial ALS. <i>Neuron</i> , 2014, 84, 324-331.	8.1	308
42	Analysis of amyotrophic lateral sclerosis as a multistep process: a population-based modelling study. <i>Lancet Neurology</i> , The, 2014, 13, 1108-1113.	10.2	302
43	Astrocyte pathology and the absence of non-cell autonomy in an induced pluripotent stem cell model of TDP-43 proteinopathy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 4697-4702.	7.1	301
44	A proposed staging system for amyotrophic lateral sclerosis. <i>Brain</i> , 2012, 135, 847-852.	7.6	296
45	Direct conversion of patient fibroblasts demonstrates non-cell autonomous toxicity of astrocytes to motor neurons in familial and sporadic ALS. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 829-832.	7.1	296
46	Clinico-pathological features in amyotrophic lateral sclerosis with expansions in C9ORF72. <i>Brain</i> , 2012, 135, 751-764.	7.6	293
47	Detection of long repeat expansions from PCR-free whole-genome sequence data. <i>Genome Research</i> , 2017, 27, 1895-1903.	5.5	277
48	Sequestration of multiple RNA recognition motif-containing proteins by C9orf72 repeat expansions. <i>Brain</i> , 2014, 137, 2040-2051.	7.6	253
49	TDP-43 Proteinopathy and ALS: Insights into Disease Mechanisms and Therapeutic Targets. <i>Neurotherapeutics</i> , 2015, 12, 352-363.	4.4	246
50	Human iPSC-derived motoneurons harbouring TARDBP or C9ORF72 ALS mutations are dysfunctional despite maintaining viability. <i>Nature Communications</i> , 2015, 6, 5999.	12.8	241
51	Widespread binding of FUS along nascent RNA regulates alternative splicing in the brain. <i>Scientific Reports</i> , 2012, 2, 603.	3.3	231
52	Familial amyotrophic lateral sclerosis is associated with a mutation in D-amino acid oxidase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 7556-7561.	7.1	229
53	FUS-SMN Protein Interactions Link the Motor Neuron Diseases ALS and SMA. <i>Cell Reports</i> , 2012, 2, 799-806.	6.4	229
54	Overexpression of human wild-type FUS causes progressive motor neuron degeneration in an age- and dose-dependent fashion. <i>Acta Neuropathologica</i> , 2013, 125, 273-288.	7.7	225

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55	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. <i>Nature Genetics</i> , 2021, 53, 1636-1648.	21.4	223
56	CSF and Plasma Amino Acid Levels in Motor Neuron Disease: Elevation of CSF Glutamate in a Subset of Patients. <i>Experimental Neurology</i> , 1995, 4, 209-216.	1.7	221
57	NEK1 variants confer susceptibility to amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016, 48, 1037-1042.	21.4	218
58	Differential roles of the ubiquitin proteasome system (UPS) and autophagy in the clearance of soluble and aggregated TDP-43 species. <i>Journal of Cell Science</i> , 2014, 127, 1263-78.	2.0	216
59	Chromosome 9p21 in sporadic amyotrophic lateral sclerosis in the UK and seven other countries: a genome-wide association study. <i>Lancet Neurology</i> , The, 2010, 9, 986-994.	10.2	205
60	Mutations in CHMP2B in Lower Motor Neuron Predominant Amyotrophic Lateral Sclerosis (ALS). <i>PLoS ONE</i> , 2010, 5, e9872.	2.5	204
61	The C9ORF72 expansion mutation is a common cause of ALS+FTD in Europe and has a single founder. <i>European Journal of Human Genetics</i> , 2013, 21, 102-108.	2.8	201
62	ALS/FTD-associated FUS activates GSK3 β to disrupt the VAPB-PTPIP51 interaction and ER-mitochondria associations. <i>EMBO Reports</i> , 2016, 17, 1326-1342.	4.5	201
63	Analysis of alternative splicing associated with aging and neurodegeneration in the human brain. <i>Genome Research</i> , 2011, 21, 1572-1582.	5.5	199
64	ALS mutant FUS disrupts nuclear localization and sequesters wild-type FUS within cytoplasmic stress granules. <i>Human Molecular Genetics</i> , 2013, 22, 2676-2688.	2.9	199
65	Neurofilament heavy chain side arm phosphorylation regulates axonal transport of neurofilaments. <i>Journal of Cell Biology</i> , 2003, 161, 489-495.	5.2	185
66	ALS/FTD-Linked Mutation in FUS Suppresses Intra-axonal Protein Synthesis and Drives Disease Without Nuclear Loss-of-Function of FUS. <i>Neuron</i> , 2018, 100, 816-830.e7.	8.1	185
67	Charcot-Marie-Tooth disease neurofilament mutations disrupt neurofilament assembly and axonal transport. <i>Human Molecular Genetics</i> , 2002, 11, 2837-2844.	2.9	183
68	Gene expression profiling in human neurodegenerative disease. <i>Nature Reviews Neurology</i> , 2012, 8, 518-530.	10.1	183
69	Amyotrophic lateral sclerosis in an urban setting. <i>Journal of Neurology</i> , 2006, 253, 1642-1643.	3.6	181
70	Reduced expression of the Kinesin-Associated Protein 3 (KIFAP3) gene increases survival in sporadic amyotrophic lateral sclerosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 9004-9009.	7.1	177
71	CCNF mutations in amyotrophic lateral sclerosis and frontotemporal dementia. <i>Nature Communications</i> , 2016, 7, 11253.	12.8	174
72	ALS-specific cognitive and behavior changes associated with advancing disease stage in ALS. <i>Neurology</i> , 2018, 91, e1370-e1380.	1.1	170

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73	Nuclear import impairment causes cytoplasmic trans-activation response DNA-binding protein accumulation and is associated with frontotemporal lobar degeneration. <i>Brain</i> , 2010, 133, 1763-1771.	7.6	165
74	Novel insertion in the KSP region of the neurofilament heavy gene in amyotrophic lateral sclerosis (ALS). <i>NeuroReport</i> , 1998, 9, 3967-3970.	1.2	157
75	The role of <i>TREM2</i> R47H as a risk factor for Alzheimer's disease, frontotemporal lobar degeneration, amyotrophic lateral sclerosis, and Parkinson's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 1407-1416.	0.8	152
76	C9ORF72 repeat expansion causes vulnerability of motor neurons to Ca ²⁺ -permeable AMPA receptor-mediated excitotoxicity. <i>Nature Communications</i> , 2018, 9, 347.	12.8	151
77	The widening spectrum of C9ORF72-related disease; genotype/phenotype correlations and potential modifiers of clinical phenotype. <i>Acta Neuropathologica</i> , 2014, 127, 333-345.	7.7	150
78	Glutamate Slows Axonal Transport of Neurofilaments in Transfected Neurons. <i>Journal of Cell Biology</i> , 2000, 150, 165-176.	5.2	149
79	Antisense RNA foci in the motor neurons of C9ORF72-ALS patients are associated with TDP-43 proteinopathy. <i>Acta Neuropathologica</i> , 2015, 130, 63-75.	7.7	149
80	Mutational analysis reveals the <i>FUS</i> homolog <i>TAF15</i> as a candidate gene for familial amyotrophic lateral sclerosis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 285-290.	1.7	148
81	Oligodendrocytes contribute to motor neuron death in ALS via SOD1-dependent mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6496-E6505.	7.1	139
82	Direct evidence for axonal transport defects in a novel mouse model of mutant spastin-induced hereditary spastic paraplegia (HSP) and human HSP patients. <i>Journal of Neurochemistry</i> , 2009, 110, 34-44.	3.9	135
83	Mechanisms, models and biomarkers in amyotrophic lateral sclerosis. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2013, 14, 19-32.	1.7	135
84	Molecular pathology and genetic advances in amyotrophic lateral sclerosis: an emerging molecular pathway and the significance of glial pathology. <i>Acta Neuropathologica</i> , 2011, 122, 657-671.	7.7	134
85	Unravelling the enigma of selective vulnerability in neurodegeneration: motor neurons resistant to degeneration in ALS show distinct gene expression characteristics and decreased susceptibility to excitotoxicity. <i>Acta Neuropathologica</i> , 2013, 125, 95-109.	7.7	133
86	C9ORF72 interaction with cofilin modulates actin dynamics in motor neurons. <i>Nature Neuroscience</i> , 2016, 19, 1610-1618.	14.8	131
87	The heat shock response plays an important role in TDP-43 clearance: evidence for dysfunction in amyotrophic lateral sclerosis. <i>Brain</i> , 2016, 139, 1417-1432.	7.6	131
88	Is language impairment more common than executive dysfunction in amyotrophic lateral sclerosis?. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2013, 84, 494-498.	1.9	130
89	ALS-associated mutations in <i>FUS</i> disrupt the axonal distribution and function of SMN. <i>Human Molecular Genetics</i> , 2013, 22, 3690-3704.	2.9	130
90	Mutations in the vesicular trafficking protein annexin A11 are associated with amyotrophic lateral sclerosis. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	129

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91	Combination of neurofilament heavy chain and complement C3 as CSF biomarkers for ALS. <i>Journal of Neurochemistry</i> , 2011, 117, 528-537.	3.9	128
92	Amyotrophic Lateral Sclerosis in South-East England: A Population-Based Study. <i>Neuroepidemiology</i> , 2007, 29, 44-48.	2.3	127
93	Amyotrophic Lateral Sclerosis Associated with Genetic Abnormalities in the Gene Encoding Cu/Zn Superoxide Dismutase: Molecular Pathology of Five New Cases, and Comparison with Previous Reports and 73 Sporadic Cases of ALS. <i>Journal of Neuropathology and Experimental Neurology</i> , 1998, 57, 895-904.	1.7	124
94	A genome-wide association meta-analysis identifies a novel locus at 17q11.2 associated with sporadic amyotrophic lateral sclerosis. <i>Human Molecular Genetics</i> , 2014, 23, 2220-2231.	2.9	123
95	The RNA of the glutamate transporter EAAT2 is variably spliced in amyotrophic lateral sclerosis and normal individuals. <i>Journal of the Neurological Sciences</i> , 1999, 170, 45-50.	0.6	121
96	Familial amyotrophic lateral sclerosis with a mutation in exon 4 of the Cu/Zn superoxide dismutase gene: pathological and immunocytochemical changes. <i>Acta Neuropathologica</i> , 1996, 92, 395-403.	7.7	120
97	A blinded international study on the reliability of genetic testing for GGGGCC-repeat expansions in <i>C9orf72</i> reveals marked differences in results among 14 laboratories. <i>Journal of Medical Genetics</i> , 2014, 51, 419-424.	3.2	118
98	Shared polygenic risk and causal inferences in amyotrophic lateral sclerosis. <i>Annals of Neurology</i> , 2019, 85, 470-481.	5.3	118
99	Downregulation of MicroRNA-9 in iPSC-Derived Neurons of FTD/ALS Patients with TDP-43 Mutations. <i>PLoS ONE</i> , 2013, 8, e76055.	2.5	117
100	The chromosome 9 ALS and FTD locus is probably derived from a single founder. <i>Neurobiology of Aging</i> , 2012, 33, 209.e3-209.e8.	3.1	115
101	Loss and gain of <i>Drosophila</i> TDP-43 impair synaptic efficacy and motor control leading to age-related neurodegeneration by loss-of-function phenotypes. <i>Human Molecular Genetics</i> , 2013, 22, 1539-1557.	2.9	115
102	Dominant mutations in the cation channel gene transient receptor potential vanilloid 4 cause an unusual spectrum of neuropathies. <i>Brain</i> , 2010, 133, 1798-1809.	7.6	113
103	Novel FUS/TLS Mutations and Pathology in Familial and Sporadic Amyotrophic Lateral Sclerosis. <i>Archives of Neurology</i> , 2010, 67, 455-61.	4.5	113
104	Amyotrophic lateral sclerosis-associated mutant VAPBP56S perturbs calcium homeostasis to disrupt axonal transport of mitochondria. <i>Human Molecular Genetics</i> , 2012, 21, 1979-1988.	2.9	112
105	Association of apolipoprotein E ϵ 4 allele with bulbar-onset motor neuron disease. <i>Lancet</i> , 1996, 347, 159-160.	13.7	111
106	Estimating clinical stage of amyotrophic lateral sclerosis from the ALS Functional Rating Scale. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2014, 15, 279-284.	1.7	111
107	An MND/ALS phenotype associated with <i>C9orf72</i> repeat expansion: Abundant p62-positive, TDP-43-negative inclusions in cerebral cortex, hippocampus and cerebellum but without associated cognitive decline. <i>Neuropathology</i> , 2012, 32, 505-514.	1.2	110
108	TDP-43 is consistently co-localized with ubiquitinated inclusions in sporadic and Guam amyotrophic lateral sclerosis but not in familial amyotrophic lateral sclerosis with and without SOD1 mutations. <i>Neuropathology</i> , 2009, 29, 672-683.	1.2	108

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109	SRSF1-dependent nuclear export inhibition of C9ORF72 repeat transcripts prevents neurodegeneration and associated motor deficits. <i>Nature Communications</i> , 2017, 8, 16063.	12.8	106
110	p38 β stress-activated protein kinase phosphorylates neurofilaments and is associated with neurofilament pathology in amyotrophic lateral sclerosis. <i>Molecular and Cellular Neurosciences</i> , 2004, 26, 354-364.	2.2	104
111	Exome sequencing in amyotrophic lateral sclerosis implicates a novel gene, DNAJC7, encoding a heat-shock protein. <i>Nature Neuroscience</i> , 2019, 22, 1966-1974.	14.8	101
112	Loss of nuclear TDP43 in amyotrophic lateral sclerosis (ALS) causes altered expression of splicing machinery and widespread dysregulation of RNA splicing in motor neurones. <i>Neuropathology and Applied Neurobiology</i> , 2014, 40, 670-685.	3.2	98
113	Review: Glial lineages and myelination in the central nervous system. <i>Journal of Anatomy</i> , 1997, 190, 161-200.	1.5	96
114	Amyotrophic lateral sclerosis with sensory neuropathy: part of a multisystem disorder?. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2006, 78, 750-753.	1.9	96
115	Rasch analysis of the hospital anxiety and depression scale (hads) for use in motor neurone disease. <i>Health and Quality of Life Outcomes</i> , 2011, 9, 82.	2.4	96
116	C9orf72 and UNC13A are shared risk loci for amyotrophic lateral sclerosis and frontotemporal dementia: A genome-wide meta-analysis. <i>Annals of Neurology</i> , 2014, 76, 120-133.	5.3	91
117	D90A-SOD1 mediated amyotrophic lateral sclerosis: A single founder for all cases with evidence for a cis-acting disease modifier in the recessive haplotype. <i>Human Mutation</i> , 2002, 20, 473-473.	2.5	90
118	Mutant C9orf72 human iPSC-derived astrocytes cause non-cell autonomous motor neuron pathophysiology. <i>Glia</i> , 2020, 68, 1046-1064.	4.9	90
119	Phosphorylation of neurofilament heavy chain side-arms by stress activated protein kinase-1b/Jun N-terminal kinase-3. <i>Journal of Cell Science</i> , 2000, 113, 401-407.	2.0	90
120	Novel mutations support a role for Profilin 1 in the pathogenesis of ALS. <i>Neurobiology of Aging</i> , 2015, 36, 1602.e17-1602.e27.	3.1	87
121	Biomarkers in Motor Neuron Disease: A State of the Art Review. <i>Frontiers in Neurology</i> , 2019, 10, 291.	2.4	87
122	ATXN2 trinucleotide repeat length correlates with risk of ALS. <i>Neurobiology of Aging</i> , 2017, 51, 178.e1-178.e9.	3.1	86
123	Astrocyte adenosine deaminase loss increases motor neuron toxicity in amyotrophic lateral sclerosis. <i>Brain</i> , 2019, 142, 586-605.	7.6	84
124	C9ORF72 GGGGCC Expanded Repeats Produce Splicing Dysregulation which Correlates with Disease Severity in Amyotrophic Lateral Sclerosis. <i>PLoS ONE</i> , 2015, 10, e0127376.	2.5	83
125	Prognostic modelling of therapeutic interventions in amyotrophic lateral sclerosis. <i>Amyotrophic Lateral Sclerosis and Other Motor Neuron Disorders: Official Publication of the World Federation of Neurology, Research Group on Motor Neuron Diseases</i> , 2002, 3, 15-21.	1.2	82
126	Physical activity as an exogenous risk factor in motor neuron disease (MND): A review of the evidence. <i>Amyotrophic Lateral Sclerosis and Other Motor Neuron Disorders</i> , 2009, 10, 191-204.	2.1	82

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127	Meta-analysis of pharmacogenetic interactions in amyotrophic lateral sclerosis clinical trials. <i>Neurology</i> , 2017, 89, 1915-1922.	1.1	82
128	Transcriptomic indices of fast and slow disease progression in two mouse models of amyotrophic lateral sclerosis. <i>Brain</i> , 2013, 136, 3305-3332.	7.6	81
129	Optimised and Rapid Pre-clinical Screening in the SOD1G93A Transgenic Mouse Model of Amyotrophic Lateral Sclerosis (ALS). <i>PLoS ONE</i> , 2011, 6, e23244.	2.5	80
130	Dipeptide repeat protein inclusions are rare in the spinal cord and almost absent from motor neurons in C9ORF72 mutant amyotrophic lateral sclerosis and are unlikely to cause their degeneration. <i>Acta Neuropathologica Communications</i> , 2015, 3, 38.	5.2	80
131	Broad clinical phenotypes associated with TAR-DNA binding protein (TARDBP) mutations in amyotrophic lateral sclerosis. <i>Neurogenetics</i> , 2010, 11, 217-225.	1.4	79
132	Amyotrophic lateral sclerosis-like superoxide dismutase 1 proteinopathy is associated with neuronal loss in Parkinson's disease brain. <i>Acta Neuropathologica</i> , 2017, 134, 113-127.	7.7	78
133	Superoxide dismutase 1 mutation in a cellular model of amyotrophic lateral sclerosis shifts energy generation from oxidative phosphorylation to glycolysis. <i>Neurobiology of Aging</i> , 2014, 35, 1499-1509.	3.1	77
134	Two Families with Familial Amyotrophic Lateral Sclerosis Are Linked to a Novel Locus on Chromosome 16q. <i>American Journal of Human Genetics</i> , 2003, 73, 390-396.	6.2	76
135	Use of clinical staging in amyotrophic lateral sclerosis for phase 3 clinical trials. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2015, 86, 45-49.	1.9	75
136	A feedback loop between dipeptide-repeat protein, TDP-43 and karyopherin- β mediates C9orf72-related neurodegeneration. <i>Brain</i> , 2018, 141, 2908-2924.	7.6	75
137	C9orf72 intermediate expansions of 24-30 repeats are associated with ALS. <i>Acta Neuropathologica Communications</i> , 2019, 7, 115.	5.2	75
138	ALS2/Alsin Regulates Rac-PAK Signaling and Neurite Outgrowth. <i>Journal of Biological Chemistry</i> , 2005, 280, 34735-34740.	3.4	74
139	Non-invasive ventilation in motor neuron disease: an update of current UK practice. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2012, 83, 371-376.	1.9	74
140	Mitochondrial abnormalities and disruption of the neuromuscular junction precede the clinical phenotype and motor neuron loss in hFUSWT transgenic mice. <i>Human Molecular Genetics</i> , 2018, 27, 463-474.	2.9	74
141	Mixed tau, TDP-43 and p62 pathology in FTLD associated with a C9ORF72 repeat expansion and p.Ala239Thr MAPT (tau) variant. <i>Acta Neuropathologica</i> , 2013, 125, 303-310.	7.7	73
142	Wild type human TDP-43 potentiates ALS-linked mutant TDP-43 driven progressive motor and cortical neuron degeneration with pathological features of ALS. <i>Acta Neuropathologica Communications</i> , 2015, 3, 36.	5.2	73
143	Gene expression signatures in motor neurone disease fibroblasts reveal dysregulation of metabolism, hypoxia response and <i>scn</i> RNA processing functions. <i>Neuropathology and Applied Neurobiology</i> , 2015, 41, 201-226.	3.2	73
144	Drosophila TDP-43 dysfunction in glia and muscle cells cause cytological and behavioural phenotypes that characterize ALS and FTLD. <i>Human Molecular Genetics</i> , 2013, 22, 3883-3893.	2.9	72

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