Leonard Petrucelli

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

150	13,729	61	116
papers	citations	h-index	g-index
161	16,876 ext. citations	14.1	6.11
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
150	TDP-43-associated atrophy in brains with and without frontotemporal lobar degeneration <i>NeuroImage: Clinical</i> , 2022 , 34, 102954	5.3	O
149	TDP-43 represses cryptic exon inclusion in the FTD-ALS gene UNC13A <i>Nature</i> , 2022 ,	50.4	14
148	Amyotrophic lateral sclerosis - insight into susceptibility <i>Nature Reviews Neurology</i> , 2022 ,	15	1
147	Homotypic fibrillization of TMEM106B across diverse neurodegenerative diseases Cell, 2022,	56.2	5
146	Plasma PolyQ-ATXN3 Levels Associate With Cerebellar Degeneration and Behavioral Abnormalities in a New AAV-Based SCA3 Mouse Model <i>Frontiers in Cell and Developmental Biology</i> , 2022 , 10, 863089	5.7	O
145	Modelling amyotrophic lateral sclerosis in rodents Nature Reviews Neuroscience, 2022,	13.5	1
144	Comprehensive cross-sectional and longitudinal analyses of plasma neurofilament light across FTD spectrum disorders <i>Cell Reports Medicine</i> , 2022 , 3, 100607	18	O
143	Cracking the cryptic code in amyotrophic lateral sclerosis and frontotemporal dementia: Towards therapeutic targets and biomarkers <i>Clinical and Translational Medicine</i> , 2022 , 12, e818	5.7	O
142	Comment on: Polyglutamine-Expanded Ataxin-3: A Target Engagement Marker for Spinocerebellar Ataxia Type 3 in Peripheral Blood <i>Movement Disorders</i> , 2022 , 37, 1120-1121	7	
141	Poly(GR) and poly(GA) in cerebrospinal fluid as potential biomarkers for C9ORF72-ALS/FTD <i>Nature Communications</i> , 2022 , 13, 2799	17.4	2
140	HDAC6 Interacts With Poly (GA) and Modulates its Accumulation in c9FTD/ALS Frontiers in Cell and Developmental Biology, 2021 , 9, 809942	5.7	O
139	A Small Molecule Exploits Hidden Structural Features within the RNA Repeat Expansion That Causes c9ALS/FTD and Rescues Pathological Hallmarks. <i>ACS Chemical Neuroscience</i> , 2021 , 12, 4076-4089	95.7	O
138	Ribonuclease recruitment using a small molecule reduced c9ALS/FTD r(GC) repeat expansion in vitro and in vivo ALS models. <i>Science Translational Medicine</i> , 2021 , 13, eabd5991	17.5	6
137	Application of a bioinformatic pipeline to RNA-seq data identifies novel virus-like sequence in human blood. <i>G3: Genes, Genomes, Genetics</i> , 2021 , 11,	3.2	2
136	Deep vein thrombosis and pulmonary embolism among hospitalized coronavirus disease 2019-positive patients predicted for higher mortality and prolonged intensive care unit and hospital stays in a multisite healthcare system. <i>Journal of Vascular Surgery: Venous and Lymphatic</i>	3.2	6
135	Long-read targeted sequencing uncovers clinicopathological associations for C9orf72-linked diseases. <i>Brain</i> , 2021 , 144, 1082-1088	11.2	2
134	Serum neurofilament light protein correlates with unfavorable clinical outcomes in hospitalized patients with COVID-19. <i>Science Translational Medicine</i> , 2021 , 13,	17.5	16

p53 is a central regulator driving neurodegeneration caused by C9orf72 poly(PR). Cell, 2021, 184, 689-70% e20 26 133 TIA1 potentiates tau phase separation and promotes generation of toxic oligomeric tau. 132 11.5 22 Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, The AD tau core spontaneously self-assembles and recruits full-length tau to filaments. Cell Reports 10.6 8 131 , **2021**, 34, 108843 NIH funding trends for neurosurgeon-scientists from 1993-2017: Biomedical workforce implications 4.8 130 for neurooncology. Journal of Neuro-Oncology, 2021, 154, 51-62 Urine levels of the polyglutamine ataxin-3 protein are elevated in patients with spinocerebellar 3.6 129 3 ataxia type 3. Parkinsonism and Related Disorders, 2021, 89, 151-154 Interaction of tau with HNRNPA2B1 and N-methyladenosine RNA mediates the progression of 128 17.6 11 tauopathy. *Molecular Cell*, **2021**, 81, 4209-4227.e12 Cellular and pathological heterogeneity of primary tauopathies. Molecular Neurodegeneration, 2021 127 19 11 , 16, 57 Microglial lysosome dysfunction contributes to white matter pathology and TDP-43 proteinopathy 126 10.6 in GRN-associated FTD. Cell Reports, 2021, 36, 109581 Cross-sectional and longitudinal measures of chitinase proteins in amyotrophic lateral sclerosis and expression of CHI3L1 in activated astrocytes. Journal of Neurology, Neurosurgery and Psychiatry, 125 5.5 22 2020, 91, 350-358 Hexanucleotide Repeat Expansions in c9FTD/ALS and SCA36 Confer Selective Patterns of 18 10.6 124 Neurodegeneration In Vivo. Cell Reports, 2020, 31, 107616 Utility of FDG-PET in diagnosis of Alzheimer-related TDP-43 proteinopathy. Neurology, 2020, 95, e23-e346.5 123 11 Premature termination codon readthrough upregulates progranulin expression and improves 8 122 19 lysosomal function in preclinical models of GRN deficiency. Molecular Neurodegeneration, 2020, 15, 21 AlPuts the Alpha in Synuclein. Neuron, 2020, 105, 205-206 121 13.9 Elevated methylation levels, reduced expression levels, and frequent contractions in a clinical 120 19 20 cohort of C9orf72 expansion carriers. Molecular Neurodegeneration, 2020, 15, 7 Posttranslational Modifications Mediate the Structural Diversity of Tauopathy Strains. Cell, 2020, 56.2 156 119 180, 633-644.e12 Reduced C9ORF72 function exacerbates gain of toxicity from ALS/FTD-causing repeat expansion in 118 80 25.5 C9orf72. Nature Neuroscience, 2020, 23, 615-624 Truncated stathmin-2 is a marker of TDP-43 pathology in frontotemporal dementia. Journal of 117 15.9 34 Clinical Investigation, **2020**, 130, 6080-6092 Chimeric Peptide Species Contribute to Divergent Dipeptide Repeat Pathology in c9ALS/FTD and 116 13.9 25 SCA36. Neuron, 2020, 107, 292-305.e6

115	Toward allele-specific targeting therapy and pharmacodynamic marker for spinocerebellar ataxia type 3. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	15
114	Astrocyte-derived clusterin suppresses amyloid formation in vivo. <i>Molecular Neurodegeneration</i> , 2020 , 15, 71	19	11
113	Plasma neurofilament light predicts mortality in patients with stroke. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	20
112	Clusterin ameliorates tau pathology in vivo by inhibiting fibril formation. <i>Acta Neuropathologica Communications</i> , 2020 , 8, 210	7.3	5
111	Structural Features of Small Molecules Targeting the RNA Repeat Expansion That Causes Genetically Defined ALS/FTD. <i>ACS Chemical Biology</i> , 2020 , 15, 3112-3123	4.9	4
110	poly(GR) aggregation induces TDP-43 proteinopathy. Science Translational Medicine, 2020, 12,	17.5	49
109	Mitophagy alterations in Alzheimer's disease are associated with granulovacuolar degeneration and early tau pathology. <i>Alzheimermand Dementia</i> , 2020 , 17, 417	1.2	13
108	The influence of tau, amyloid, alpha-synuclein, TDP-43, and vascular pathology in clinically normal elderly individuals. <i>Neurobiology of Aging</i> , 2019 , 77, 26-36	5.6	32
107	Enhanced phosphorylation of T153 in soluble tau is a defining biochemical feature of the A152T tau risk variant. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 10	7.3	1
106	Toxic expanded GGGGCC repeat transcription is mediated by the PAF1 complex in C9orf72-associated FTD. <i>Nature Neuroscience</i> , 2019 , 22, 863-874	25.5	38
105	Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. <i>Genome Biology</i> , 2019 , 20, 97	18.3	68
104	eIF4B and eIF4H mediate GR production from expanded G4C2 in a Drosophila model for C9orf72-associated ALS. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 62	7.3	22
103	Genetic Convergence Brings Clarity to the Enigmatic Red Line in ALS. <i>Neuron</i> , 2019 , 101, 1057-1069	13.9	63
102	Tau exhibits unique seeding properties in globular glial tauopathy. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 36	7.3	14
101	ADAR2 mislocalization and widespread RNA editing aberrations in C9orf72-mediated ALS/FTD. <i>Acta Neuropathologica</i> , 2019 , 138, 49-65	14.3	27
100	Genome-wide analyses as part of the international FTLD-TDP whole-genome sequencing consortium reveals novel disease risk factors and increases support for immune dysfunction in FTLD. <i>Acta Neuropathologica</i> , 2019 , 137, 879-899	14.3	50
99	Aberrant deposition of stress granule-resident proteins linked to C9orf72-associated TDP-43 proteinopathy. <i>Molecular Neurodegeneration</i> , 2019 , 14, 9	19	64
98	RPS25 is required for efficient RAN translation of C9orf72 and other neurodegenerative disease-associated nucleotide repeats. <i>Nature Neuroscience</i> , 2019 , 22, 1383-1388	25.5	54

(2018-2019)

97	C-terminal and full length TDP-43 specie differ according to FTLD-TDP lesion type but not genetic mutation. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 100	7.3	9
96	Extensive transcriptomic study emphasizes importance of vesicular transport in C9orf72 expansion carriers. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 150	7.3	18
95	Microglia in frontotemporal lobar degeneration with progranulin or C9ORF72 mutations. <i>Annals of Clinical and Translational Neurology</i> , 2019 , 6, 1782-1796	5.3	11
94	Heterochromatin anomalies and double-stranded RNA accumulation underlie poly(PR) toxicity. <i>Science</i> , 2019 , 363,	33.3	104
93	Pathological, imaging and genetic characteristics support the existence of distinct TDP-43 types in non-FTLD brains. <i>Acta Neuropathologica</i> , 2019 , 137, 227-238	14.3	32
92	The Hairpin Form of r(GC) in c9ALS/FTD Is Repeat-Associated Non-ATG Translated and a Target for Bioactive Small Molecules. <i>Cell Chemical Biology</i> , 2019 , 26, 179-190.e12	8.2	43
91	TIA1 regulates the generation and response to toxic tau oligomers. <i>Acta Neuropathologica</i> , 2019 , 137, 259-277	14.3	39
90	Poly(GP), neurofilament and grey matter deficits in expansion carriers. <i>Annals of Clinical and Translational Neurology</i> , 2018 , 5, 583-597	5.3	29
89	Unaffected mosaic case: RNA foci, dipeptide proteins, but upregulated C9orf72 expression. <i>Neurology</i> , 2018 , 90, e323-e331	6.5	24
88	CUG initiation and frameshifting enable production of dipeptide repeat proteins from ALS/FTD C9ORF72 transcripts. <i>Nature Communications</i> , 2018 , 9, 152	17.4	77
87	TDP-43 pathology disrupts nuclear pore complexes and nucleocytoplasmic transport in ALS/FTD. <i>Nature Neuroscience</i> , 2018 , 21, 228-239	25.5	240
86	A zebrafish model for C9orf72 ALS reveals RNA toxicity as a pathogenic mechanism. <i>Acta Neuropathologica</i> , 2018 , 135, 427-443	14.3	66
85	Potential genetic modifiers of disease risk and age at onset in patients with frontotemporal lobar degeneration and GRN mutations: a genome-wide association study. <i>Lancet Neurology, The</i> , 2018 , 17, 548-558	24.1	60
84	Disease Mechanisms of Repeat Expansions. Cold Spring Harbor Perspectives in Medicine, 2018, 8,	5.4	49
83	Loss of Tmem106b is unable to ameliorate frontotemporal dementia-like phenotypes in an AAV mouse model of C9ORF72-repeat induced toxicity. <i>Acta Neuropathologica Communications</i> , 2018 , 6, 42	7.3	14
82	Replication of progressive supranuclear palsy genome-wide association study identifies SLCO1A2 and DUSP10 as new susceptibility loci. <i>Molecular Neurodegeneration</i> , 2018 , 13, 37	19	28
81	Microglial translational profiling reveals a convergent APOE pathway from aging, amyloid, and tau. <i>Journal of Experimental Medicine</i> , 2018 , 215, 2235-2245	16.6	85
8o	Poly-GR dipeptide repeat polymers correlate with neurodegeneration and Clinicopathological subtypes in C9ORF72-related brain disease. <i>Acta Neuropathologica Communications</i> , 2018 , 6, 63	7.3	51

79	Dipeptide repeat proteins activate a heat shock response found in C9ORF72-ALS/FTLD patients. <i>Acta Neuropathologica Communications</i> , 2018 , 6, 55	7.3	15
78	Long-read sequencing across the C9orf72 'GGGGCC' repeat expansion: implications for clinical use and genetic discovery efforts in human disease. <i>Molecular Neurodegeneration</i> , 2018 , 13, 46	19	66
77	The Caenorhabditis elegans Ortholog of TDP-43 Regulates the Chromatin Localization of the Heterochromatin Protein 1 Homolog HPL-2. <i>Molecular and Cellular Biology</i> , 2018 , 38,	4.8	8
76	OPTN p.Met468Arg and ATXN2 intermediate length polyQ extension in families with C9orf72 mediated amyotrophic lateral sclerosis and frontotemporal dementia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 75-85	3.5	10
75	Converging pathways in neurodegeneration, from genetics to mechanisms. <i>Nature Neuroscience</i> , 2018 , 21, 1300-1309	25.5	183
74	Association of Apolipoprotein E & With Transactive Response DNA-Binding Protein 43. <i>JAMA Neurology</i> , 2018 , 75, 1347-1354	17.2	42
73	APOE 2 is associated with increased tau pathology in primary tauopathy. <i>Nature Communications</i> , 2018 , 9, 4388	17.4	68
7 2	TRIO gene segregation in a family with cerebellar ataxia. <i>Neurologia I Neurochirurgia Polska</i> , 2018 , 52, 743-749	1	5
71	Tau Protein Disrupts Nucleocytoplasmic Transport in Alzheimer's Disease. <i>Neuron</i> , 2018 , 99, 925-940.e7	13.9	169
70	Poly(GR) impairs protein translation and stress granule dynamics in C9orf72-associated frontotemporal dementia and amyotrophic lateral sclerosis. <i>Nature Medicine</i> , 2018 , 24, 1136-1142	50.5	149
69	Tau aggregation influences cognition and hippocampal atrophy in the absence of beta-amyloid: a clinico-imaging-pathological study of primary age-related tauopathy (PART). <i>Acta Neuropathologica</i> , 2017 , 133, 705-715	14.3	91
68	Spinal poly-GA inclusions in a C9orf72 mouse model trigger motor deficits and inflammation without neuron loss. <i>Acta Neuropathologica</i> , 2017 , 134, 241-254	14.3	70
67	Mutant TDP-43 does not impair mitochondrial bioenergetics in vitro and in vivo. <i>Molecular Neurodegeneration</i> , 2017 , 12, 37	19	28
66	In-depth clinico-pathological examination of RNA foci in a large cohort of C9ORF72 expansion carriers. <i>Acta Neuropathologica</i> , 2017 , 134, 255-269	14.3	57
65	Phosphorylated neurofilament heavy chain: A biomarker of survival for C9ORF72-associated amyotrophic lateral sclerosis. <i>Annals of Neurology</i> , 2017 , 82, 139-146	9.4	58
64	TDP-43 mutations causing amyotrophic lateral sclerosis are associated with altered expression of RNA-binding protein hnRNP K and affect the Nrf2 antioxidant pathway. <i>Human Molecular Genetics</i> , 2017 , 26, 1732-1746	5.6	41
63	Poly(GP) proteins are a useful pharmacodynamic marker for -associated amyotrophic lateral sclerosis. <i>Science Translational Medicine</i> , 2017 , 9,	17.5	128
62	C9orf72 poly GA RAN-translated protein plays a key role in amyotrophic lateral sclerosis via aggregation and toxicity. <i>Human Molecular Genetics</i> , 2017 , 26, 4765-4777	5.6	43

(2016-2017)

61	ARHGEF28 p.Lys280Metfs40Ter in an amyotrophic lateral sclerosis family with a C9orf72 expansion. <i>Neurology: Genetics</i> , 2017 , 3, e190	3.8	4
60	The lysosomal protein cathepsin L is a progranulin protease. <i>Molecular Neurodegeneration</i> , 2017 , 12, 55	19	54
59	An acetylation-phosphorylation switch that regulates tau aggregation propensity and function. <i>Journal of Biological Chemistry</i> , 2017 , 292, 15277-15286	5.4	78
58	TIA1 Mutations in Amyotrophic Lateral Sclerosis and Frontotemporal Dementia Promote Phase Separation and Alter Stress Granule Dynamics. <i>Neuron</i> , 2017 , 95, 808-816.e9	13.9	341
57	Repetitive element transcripts are elevated in the brain of C9orf72 ALS/FTLD patients. <i>Human Molecular Genetics</i> , 2017 , 26, 3421-3431	5.6	63
56	Loss of clusterin shifts amyloid deposition to the cerebrovasculature via disruption of perivascular drainage pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E6962-E6971	11.5	66
55	Abnormal expression of homeobox genes and transthyretin in expansion carriers. <i>Neurology: Genetics</i> , 2017 , 3, e161	3.8	9
54	FTDP-17 with Pick body-like inclusions associated with a novel tau mutation, p.E372G. <i>Brain Pathology</i> , 2017 , 27, 612-626	6	11
53	An autoradiographic evaluation of AV-1451 Tau PET in dementia. <i>Acta Neuropathologica Communications</i> , 2016 , 4, 58	7.3	305
52	Timing and significance of pathological features in C9orf72 expansion-associated frontotemporal dementia. <i>Brain</i> , 2016 , 139, 3202-3216	11.2	90
51	Monitoring peripheral nerve degeneration in ALS by label-free stimulated Raman scattering imaging. <i>Nature Communications</i> , 2016 , 7, 13283	17.4	56
50	ALS and FTD: an epigenetic perspective. <i>Acta Neuropathologica</i> , 2016 , 132, 487-502	14.3	51
49	Interaction of tau with the RNA-Binding Protein TIA1 Regulates tau Pathophysiology and Toxicity. <i>Cell Reports</i> , 2016 , 15, 1455-1466	10.6	176
48	Updated TDP-43 in Alzheimer's disease staging scheme. <i>Acta Neuropathologica</i> , 2016 , 131, 571-85	14.3	168
47	TDP-43 functions within a network of hnRNP proteins to inhibit the production of a truncated human SORT1 receptor. <i>Human Molecular Genetics</i> , 2016 , 25, 534-45	5.6	52
46	C9ORF72 poly(GA) aggregates sequester and impair HR23 and nucleocytoplasmic transport proteins. <i>Nature Neuroscience</i> , 2016 , 19, 668-677	25.5	201
45	C9orf72 promoter hypermethylation is reduced while hydroxymethylation is acquired during reprogramming of ALS patient cells. <i>Experimental Neurology</i> , 2016 , 277, 171-177	5.7	16
44	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, 535-50	13.9	331

43	Poly(GR) in C9ORF72-Related ALS/FTD Compromises Mitochondrial Function and Increases Oxidative Stress and DNA Damage in iPSC-Derived Motor Neurons. <i>Neuron</i> , 2016 , 92, 383-391	13.9	220
42	Spt4 selectively regulates the expression of C9orf72 sense and antisense mutant transcripts. <i>Science</i> , 2016 , 353, 708-12	33.3	92
41	Insights into the pathogenic mechanisms of Chromosome 9 open reading frame 72 (C9orf72) repeat expansions. <i>Journal of Neurochemistry</i> , 2016 , 138 Suppl 1, 145-62	6	54
40	Understanding biomarkers of neurodegeneration: Novel approaches to detecting tau pathology. <i>Nature Medicine</i> , 2015 , 21, 219-20	50.5	14
39	Distinct brain transcriptome profiles in C9orf72-associated and sporadic ALS. <i>Nature Neuroscience</i> , 2015 , 18, 1175-82	25.5	235
38	Neurodegeneration. C9ORF72 repeat expansions in mice cause TDP-43 pathology, neuronal loss, and behavioral deficits. <i>Science</i> , 2015 , 348, 1151-4	33.3	279
37	Differential Toxicity of Nuclear RNA Foci versus Dipeptide Repeat Proteins in a Drosophila Model of C9ORF72 FTD/ALS. <i>Neuron</i> , 2015 , 87, 1207-1214	13.9	149
36	Novel clinical associations with specific C9ORF72 transcripts in patients with repeat expansions in C9ORF72. <i>Acta Neuropathologica</i> , 2015 , 130, 863-76	14.3	81
35	GGGGCC repeat expansion in C9orf72 compromises nucleocytoplasmic transport. <i>Nature</i> , 2015 , 525, 129-33	50.4	540
34	Cerebellar c9RAN proteins associate with clinical and neuropathological characteristics of C9ORF72 repeat expansion carriers. <i>Acta Neuropathologica</i> , 2015 , 130, 559-73	14.3	72
33	Quantitative analysis and clinico-pathological correlations of different dipeptide repeat protein pathologies in C9ORF72 mutation carriers. <i>Acta Neuropathologica</i> , 2015 , 130, 845-61	14.3	155
32	C9orf72 BAC Transgenic Mice Display Typical Pathologic Features of ALS/FTD. <i>Neuron</i> , 2015 , 88, 892-90	1 13.9	201
31	Human C9ORF72 Hexanucleotide Expansion Reproduces RNA Foci and Dipeptide Repeat Proteins but Not Neurodegeneration in BAC Transgenic Mice. <i>Neuron</i> , 2015 , 88, 902-909	13.9	183
30	Linking the VPS35 and EIF4G1 pathways in Parkinson's disease. <i>Neuron</i> , 2015 , 85, 1-3	13.9	29
29	Mechanisms of toxicity in C9FTLD/ALS. <i>Acta Neuropathologica</i> , 2014 , 127, 359-76	14.3	114
28	Characterization of DNA hypermethylation in the cerebellum of c9FTD/ALS patients. <i>Brain Research</i> , 2014 , 1584, 15-21	3.7	63
27	Severe amygdala dysfunction in a MAPT transgenic mouse model of frontotemporal dementia. <i>Neurobiology of Aging</i> , 2014 , 35, 1769-77	5.6	37
26	Ataxin-2 as potential disease modifier in C9ORF72 expansion carriers. <i>Neurobiology of Aging</i> , 2014 , 35, 2421.e13-7	5.6	62

(2012-2014)

25	Discovery of a biomarker and lead small molecules to target r(GGGGCC)-associated defects in c9FTD/ALS. <i>Neuron</i> , 2014 , 83, 1043-50	13.9	232
24	Epigenetic modifications of theC9ORF72gene: a potential biomarker of disease?. <i>Future Neurology</i> , 2014 , 9, 123-126	1.5	
23	ER-mitochondria associations are regulated by the VAPB-PTPIP51 interaction and are disrupted by ALS/FTD-associated TDP-43. <i>Nature Communications</i> , 2014 , 5, 3996	17.4	341
22	Alterations in microRNA-124 and AMPA receptors contribute to social behavioral deficits in frontotemporal dementia. <i>Nature Medicine</i> , 2014 , 20, 1444-51	50.5	125
21	TDP-1, the Caenorhabditis elegans ortholog of TDP-43, limits the accumulation of double-stranded RNA. <i>EMBO Journal</i> , 2014 , 33, 2947-66	13	44
20	Aggregation-prone c9FTD/ALS poly(GA) RAN-translated proteins cause neurotoxicity by inducing ER stress. <i>Acta Neuropathologica</i> , 2014 , 128, 505-24	14.3	227
19	Expanded C9ORF72 hexanucleotide repeat in depressive pseudodementia. <i>JAMA Neurology</i> , 2014 , 71, 775-81	17.2	24
18	Divergent phenotypes in mutant TDP-43 transgenic mice highlight potential confounds in TDP-43 transgenic modeling. <i>PLoS ONE</i> , 2014 , 9, e86513	3.7	23
17	Homozygosity for the C9orf72 GGGGCC repeat expansion in frontotemporal dementia. <i>Acta Neuropathologica</i> , 2013 , 126, 401-9	14.3	119
16	Targeting RNA foci in iPSC-derived motor neurons from ALS patients with a C9ORF72 repeat expansion. <i>Science Translational Medicine</i> , 2013 , 5, 208ra149	17.5	488
15	Association between repeat sizes and clinical and pathological characteristics in carriers of C9ORF72 repeat expansions (Xpansize-72): a cross-sectional cohort study. <i>Lancet Neurology, The</i> , 2013 , 12, 978-88	24.1	200
14	Unconventional translation of C9ORF72 GGGGCC expansion generates insoluble polypeptides specific to c9FTD/ALS. <i>Neuron</i> , 2013 , 77, 639-46	13.9	783
13	RNA toxicity from the ALS/FTD C9ORF72 expansion is mitigated by antisense intervention. <i>Neuron</i> , 2013 , 80, 415-28	13.9	650
12	Antisense transcripts of the expanded C9ORF72 hexanucleotide repeat form nuclear RNA foci and undergo repeat-associated non-ATG translation in c9FTD/ALS. <i>Acta Neuropathologica</i> , 2013 , 126, 829-4	44 ^{14.3}	392
11	Reduced C9orf72 gene expression in c9FTD/ALS is caused by histone trimethylation, an epigenetic event detectable in blood. <i>Acta Neuropathologica</i> , 2013 , 126, 895-905	14.3	217
10	C9ORF72 repeat expansions in cases with previously identified pathogenic mutations. <i>Neurology</i> , 2013 , 81, 1332-41	6.5	75
9	The dual functions of the extreme N-terminus of TDP-43 in regulating its biological activity and inclusion formation. <i>Human Molecular Genetics</i> , 2013 , 22, 3112-22	5.6	119
8	Misregulation of human sortilin splicing leads to the generation of a nonfunctional progranulin receptor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 21510-5	11.5	61

7	Wild-type human TDP-43 expression causes TDP-43 phosphorylation, mitochondrial aggregation, motor deficits, and early mortality in transgenic mice. <i>Journal of Neuroscience</i> , 2010 , 30, 10851-9	6.6	373
6	O1-07-01: Accelerated lipofuscinosis and ubiquitination in granulin knockout mice suggests a role for progranulin in successful aging 2010 , 6, S83-S83		
5	Aberrant cleavage of TDP-43 enhances aggregation and cellular toxicity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 7607-12	11.5	433
4	Novel mutations in TARDBP (TDP-43) in patients with familial amyotrophic lateral sclerosis. <i>PLoS Genetics</i> , 2008 , 4, e1000193	6	339
3	Identification and characterization of the human parkin gene promoter. <i>Journal of Neurochemistry</i> , 2001 , 78, 1146-52	6	25
2	Lewy bodies and parkinsonism in families with parkin mutations. <i>Annals of Neurology</i> , 2001 , 50, 293-30	0 9.4	425
1	Long-read sequencing across the C9orf72 LGGGCCL epeat expansion: implications for clinical use and genetic discovery efforts in human disease		1