## Paul J Lockhart

List of Publications by Year in descending order

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45317 44069 9,259 146 48 90 citations h-index g-index papers 158 158 158 13119 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Isolation of a partial candidate gene for Menkes disease by positional cloning. Nature Genetics, 1993, 3, 20-25.	21.4	688
2	Refining analyses of copy number variation identifies specific genes associated with developmental delay. Nature Genetics, 2014, 46, 1063-1071.	21.4	583
3	Parkin Protects against the Toxicity Associated with Mutant α-Synuclein. Neuron, 2002, 36, 1007-1019.	8.1	542
4	Targeted sequencing identifies 91 neurodevelopmental-disorder risk genes with autism and developmental-disability biases. Nature Genetics, 2017, 49, 515-526.	21.4	443
5	alpha-synuclein gene haplotypes are associated with Parkinson's disease. Human Molecular Genetics, 2001, 10, 1847-1851.	2.9	314
6	Translation Initiator EIF4G1 Mutations in Familial Parkinson Disease. American Journal of Human Genetics, 2011, 89, 398-406.	6.2	250
7	UCHL1 is a Parkinson's disease susceptibility gene. Annals of Neurology, 2004, 55, 512-521.	5.3	227
8	Mutations in RAB39B Cause X-Linked Intellectual Disability and Early-Onset Parkinson Disease with α-Synuclein Pathology. American Journal of Human Genetics, 2014, 95, 729-735.	6.2	207
9	The PARK8 Locus in Autosomal Dominant Parkinsonism: Confirmation of Linkage and Further Delineation of the Disease-Containing Interval. American Journal of Human Genetics, 2004, 74, 11-19.	6.2	195
10	Metalloprotease SPRTN/DVC1 Orchestrates Replication-Coupled DNA-Protein Crosslink Repair. Molecular Cell, 2016, 64, 704-719.	9.7	193
11	Degeneration in Different Parkinsonian Syndromes Relates to Astrocyte Type and Astrocyte Protein Expression. Journal of Neuropathology and Experimental Neurology, 2009, 68, 1073-1083.	1.7	173
12	Bioinformatics-Based Identification of Expanded Repeats: A Non-reference Intronic Pentamer Expansion in RFC1 Causes CANVAS. American Journal of Human Genetics, 2019, 105, 151-165.	6.2	170
13	Mutations in SPRTN cause early onset hepatocellular carcinoma, genomic instability and progeroid features. Nature Genetics, 2014, 46, 1239-1244.	21.4	165
14	Parkin genetics: one model for Parkinson's disease. Human Molecular Genetics, 2004, 13, 127R-133.	2.9	153
15	Mutations in DARS Cause Hypomyelination with Brain Stem and Spinal Cord Involvement and Leg Spasticity. American Journal of Human Genetics, 2013, 92, 774-780.	6.2	151
16	The Role of GMXCXXC Metal Binding Sites in the Copper-induced Redistribution of the Menkes Protein. Journal of Biological Chemistry, 1999, 274, 11170-11177.	3.4	150
17	Germline and somatic FGFR1 abnormalities in dysembryoplastic neuroepithelial tumors. Acta Neuropathologica, 2016, 131, 847-863.	7.7	143
18	RING finger 1 mutations in Parkin produce altered localization of the protein. Human Molecular Genetics, 2003, 12, 2957-2965.	2.9	138

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19	Profile of families with parkinsonism-predominant spinocerebellar ataxia type 2 (SCA2). Movement Disorders, 2004, 19, 622-629.	3.9	127
20	Quantitative proteomic analysis of mitochondrial proteins: relevance to Lewy body formation and Parkinson's disease. Molecular Brain Research, 2005, 134, 119-138.	2.3	126
21	Human and Mouse Mutations in WDR35 Cause Short-Rib Polydactyly Syndromes Due to Abnormal Ciliogenesis. American Journal of Human Genetics, 2011, 88, 508-515.	6.2	122
22	Mutations in the murine homologue of the Menkes gene in dappled and blotchy mice. Nature Genetics, 1994, 6, 374-378.	21.4	121
23	Familial cortical dysplasia caused by mutation in the mammalian target of rapamycin regulator <i>NPRL3</i> . Annals of Neurology, 2016, 79, 132-137.	5.3	116
24	Compound heterozygous <i>FXN</i> mutations and clinical outcome in friedreich ataxia. Annals of Neurology, 2016, 79, 485-495.	5.3	115
25	SCA-2 presenting as parkinsonism in an Alberta family. Neurology, 2002, 59, 1625-1627.	1.1	113
26	Identification of a Novel Gene Linked to Parkin via a Bi-directional Promoter. Journal of Molecular Biology, 2003, 326, 11-19.	4.2	111
27	Large-scale targeted sequencing identifies risk genes for neurodevelopmental disorders. Nature Communications, 2020, 11, 4932.	12.8	105
28	SCA2 may present as levodopa-responsive parkinsonism. Movement Disorders, 2003, 18, 425-429.	3.9	99
29	Correction of the Copper Transport Defect of Menkes Patient Fibroblasts by Expression of the Menkes and Wilson ATPases. Journal of Biological Chemistry, 1998, 273, 31375-31380.	3.4	97
30	Functional association of the parkin gene promoter with idiopathic Parkinson's disease. Human Molecular Genetics, 2002, $11$ , $2787$ - $2792$ .	2.9	95
31	Familial cortical dysplasia type <scp>IIA</scp> caused by a germline mutation in <i><i>&gt;scp&gt;DEPDC5</i>. Annals of Clinical and Translational Neurology, 2015, 2, 575-580.</i>	3.7	95
32	Detecting Expansions of Tandem Repeats in Cohorts Sequenced with Short-Read Sequencing Data. American Journal of Human Genetics, 2018, 103, 858-873.	6.2	93
33	An open-label trial in Friedreich ataxia suggests clinical benefit with high-dose resveratrol, without effect on frataxin levels. Journal of Neurology, 2015, 262, 1344-1353.	3.6	89
34	Oligomeric α-synuclein inhibits tubulin polymerization. Biochemical and Biophysical Research Communications, 2007, 356, 548-553.	2.1	86
35	Functional analysis and intracellular localization of the human menkes protein (MNK) stably expressed from a cDNA construct in Chinese hamster ovary cells (CHO-K1). Human Molecular Genetics, 1998, 7, 1293-1300.	2.9	84
36	Recent advances in the detection of repeat expansions with short-read next-generation sequencing. F1000Research, 2018, 7, 736.	1.6	84

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37	Molecular basis of the brindled mouse mutant (Mo(br)): a murine model of Menkes disease. Human Molecular Genetics, 1997, 6, 1037-1042.	2.9	82
38	ASK1 inhibition: a therapeutic strategy with multi-system benefits. Journal of Molecular Medicine, 2020, 98, 335-348.	3.9	75
39	DJ-1 mutations are a rare cause of recessively inherited early onset parkinsonism mediated by loss of protein function. Journal of Medical Genetics, 2004, 41, 22e-22.	3.2	74
40	Loss of function of SLC25A46 causes lethal congenital pontocerebellar hypoplasia. Brain, 2016, 139, 2877-2890.	7.6	74
41	Reducing the exome search space for Mendelian diseases using genetic linkage analysis of exome genotypes. Genome Biology, 2011, 12, R85.	9.6	72
42	Mutations in DCC cause isolated agenesis of the corpus callosum with incomplete penetrance. Nature Genetics, 2017, 49, 511-514.	21.4	69
43	Hemispheric cortical dysplasia secondary to a mosaic somatic mutation in <i>MTOR</i> . Neurology, 2015, 84, 2029-2032.	1.1	64
44	Expression of the Menkes gene homologue in mouse tissues lack of effect of copper on the mRNA levels. FEBS Letters, 1994, 351, 186-190.	2.8	58
45	It's a double knock-out! The quaking mouse is a spontaneous deletion of parkin and parkin co-regulated gene (PACRG). Movement Disorders, 2004, 19, 101-104.	3.9	58
46	The emerging role of Rab GTPases in the pathogenesis of Parkinson's disease. Movement Disorders, 2018, 33, 196-207.	3.9	55
47	Secondâ€hit <i>&gt; DEPDC5</i> mutation is limited to dysmorphic neurons in cortical dysplasia type IIA. Annals of Clinical and Translational Neurology, 2019, 6, 1338-1344.	3.7	55
48	Intracellular localization and loss of copper responsiveness of Mnk, the murine homologue of the Menkes protein, in cells from blotchy (Mo blo) and brindled (Mo br) mouse mutants. Human Molecular Genetics, 1999, 8, 1069-1075.	2.9	54
49	Deletion of the Parkin co-regulated gene causes defects in ependymal ciliary motility and hydrocephalus in the quakingviable mutant mouse. Human Molecular Genetics, 2010, 19, 1593-1602.	2.9	52
50	ALPK3-deficient cardiomyocytes generated from patient-derived induced pluripotent stem cells and mutant human embryonic stem cells display abnormal calcium handling and establish that ALPK3 deficiency underlies familial cardiomyopathy. European Heart Journal, 2016, 37, 2586-2590.	2.2	49
51	Refinement of the PARK3 locus on chromosome 2p13 and the analysis of 14 candidate genes. European Journal of Human Genetics, 2001, 9, 659-666.	2.8	46
52	Cerebrospinal fluid liquid biopsy for detecting somatic mosaicism in brain. Brain Communications, 2021, 3, fcaa235.	3.3	42
53	<i>DCC</i> mutation update: Congenital mirror movements, isolated agenesis of the corpus callosum, and developmental split brain syndrome. Human Mutation, 2018, 39, 23-39.	2.5	41
54	Prevalence of <i>RFC1</i> -mediated spinocerebellar ataxia in a North American ataxia cohort. Neurology: Genetics, 2020, 6, e440.	1.9	40

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55	<i>ARID1B</i> -mediated disorders: Mutations and possible mechanisms. Intractable and Rare Diseases Research, 2015, 4, 17-23.	0.9	38
56	Diagnostic and cost utility of whole exome sequencing in peripheral neuropathy. Annals of Clinical and Translational Neurology, 2017, 4, 318-325.	3.7	36
57	Genetic abnormalities in a large cohort of Coffin–Siris syndrome patients. Journal of Human Genetics, 2019, 64, 1173-1186.	2.3	36
58	Parkin Co-regulated Gene (PACRG) is regulated by the ubiquitin–proteasomal system and is present in the pathological features of parkinsonian diseases. Neurobiology of Disease, 2007, 27, 238-247.	4.4	32
59	DEPDC5 and NPRL3 modulate cell size, filopodial outgrowth, and localization of mTOR in neural progenitor cells and neurons. Neurobiology of Disease, 2018, 114, 184-193.	4.4	32
60	Clinical spectrum of the pentanucleotide repeat expansion in the <i>RFC1</i> gene in ataxia syndromes. Neurology, 2020, 95, e2912-e2923.	1.1	32
61	Identification and characterization of the human parkin gene promoter. Journal of Neurochemistry, 2001, 78, 1146-1152.	3.9	31
62	MCM3AP in recessive Charcot-Marie-Tooth neuropathy and mild intellectual disability. Brain, 2017, 140, 2093-2103.	7.6	31
63	<i>ADGRV1</i> is implicated in myoclonic epilepsy. Epilepsia, 2018, 59, 381-388.	5.1	31
64	Mutations in SH3PXD2B cause Borrone dermato-cardio-skeletal syndrome. European Journal of Human Genetics, 2014, 22, 741-747.	2.8	30
65	Genetic characterization identifies bottom-of-sulcus dysplasia as an mTORopathy. Neurology, 2020, 95, e2542-e2551.	1.1	30
66	Cell and Gene Therapy for Friedreich Ataxia: Progress to Date. Human Gene Therapy, 2014, 25, 684-693.	2.7	29
67	Complete callosal agenesis, pontocerebellar hypoplasia, and axonal neuropathy due to AMPD2 loss. Neurology: Genetics, 2015, 1, e16.	1.9	29
68	Somatic <i>GNAQ</i> mutation in the <i>forme fruste</i> of Sturge-Weber syndrome. Neurology: Genetics, 2018, 4, e236.	1.9	29
69	Parkin Co-Regulated Gene is involved in aggresome formation and autophagy in response to proteasomal impairment. Experimental Cell Research, 2012, 318, 2059-2070.	2.6	28
70	Generation of iPSC lines from peripheral blood mononuclear cells from 5 healthy adults. Stem Cell Research, 2019, 34, 101380.	0.7	28
71	Parental health spillover effects of paediatric rare genetic conditions. Quality of Life Research, 2020, 29, 2445-2454.	3.1	28
72	Lack of mutations in DJ-1 in a cohort of Taiwanese ethnic Chinese with early-onset parkinsonism. Movement Disorders, 2004, 19, 1065-1069.	3.9	27

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73	Functional analysis of a hypomorphic allele shows that MMP14 catalytic activity is the prime determinant of the Winchester syndrome phenotype. Human Molecular Genetics, 2018, 27, 2775-2788.	2.9	25
74	Expanding the clinical and genetic spectrum of ALPK3 variants: Phenotypes identified in pediatric cardiomyopathy patients and adults with heterozygous variants. American Heart Journal, 2020, 225, 108-119.	2.7	25
75	Spinocerebellar ataxia type 14: study of a family with an exon 5 mutation in the PRKCG gene. Journal of Neurology, Neurosurgery and Psychiatry, 2005, 76, 1720-1722.	1.9	24
76	Genotype–phenotype correlates in Taiwanese patients with earlyâ€onset recessive parkinsonism. Movement Disorders, 2009, 24, 104-108.	3.9	24
77	Gradient of brain mosaic <i>RHEB</i> variants causes a continuum of cortical dysplasia. Annals of Clinical and Translational Neurology, 2021, 8, 485-490.	3.7	24
78	Mutant torsinA interacts with tyrosine hydroxylase in cultured cells. Neuroscience, 2009, 164, 1127-1137.	2.3	23
79	Pathogenic Variants in GPC4 Cause Keipert Syndrome. American Journal of Human Genetics, 2019, 104, 914-924.	6.2	23
80	The human sideroflexin 5 (SFXN5) gene: sequence, expression analysis and exclusion as a candidate for PARK3. Gene, 2002, 285, 229-237.	2.2	22
81	Genetic heterogeneity of polymicrogyria: study of 123 patients using deep sequencing. Brain Communications, 2021, 3, fcaa221.	3.3	22
82	DCC Is Required for the Development of Nociceptive Topognosis in Mice and Humans. Cell Reports, 2018, 22, 1105-1114.	6.4	21
83	Pathogenic Variants in CEP85L Cause Sporadic and Familial Posterior Predominant Lissencephaly. Neuron, 2020, 106, 237-245.e8.	8.1	21
84	Cloning and expression analysis of the sheep ceruloplasmin cDNA. Gene, 1999, 236, 251-257.	2.2	19
85	Identification of the Human Ubiquitin Specific Protease 31 (USP31) Gene: Structure, Sequence and Expression Analysis. DNA Sequence, 2004, 15, 9-14.	0.7	19
86	De novo 325 kb microdeletion in chromosome band 10q25.3 including ATRNL1 in a boy with cognitive impairment, autism and dysmorphic features. European Journal of Medical Genetics, 2010, 53, 337-339.	1.3	19
87	Distribution of Parkinson's disease associated RAB39B in mouse brain tissue. Molecular Brain, 2020, 13, 52.	2.6	19
88	Accurate Determination of Ataxin-2 Polyglutamine Expansion in Patients with Intermediate-Range Repeats. Genetic Testing and Molecular Biomarkers, 2002, 6, 217-220.	1.7	18
89	Heterozygous mutations in <i>HSD17B4</i> cause juvenile peroxisomal D-bifunctional protein deficiency. Neurology: Genetics, 2016, 2, e114.	1.9	18
90	Cerebral hypomyelination associated with biallelic variants of <i>FIG4</i> . Human Mutation, 2019, 40, 619-630.	2.5	18

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91	Polyalanine expansion mutations in the X-linked hypopituitarism gene SOX3 result in aggresome formation and impaired transactivation. Frontiers in Bioscience - Landmark, 2007, 12, 2085.	3.0	18
92	Expression and localization of the Parkin Co-Regulated Gene in mouse CNS suggests a role in ependymal cilia function. Neuroscience Letters, 2009, 460, 97-101.	2.1	17
93	Neuropathology of childhoodâ€onset basal ganglia degeneration caused by mutation of <i>VAC14</i> Annals of Clinical and Translational Neurology, 2017, 4, 859-864.	3.7	17
94	Eukaryotic Expression Vectors That Replicate to Low Copy Number in Bacteria: Transient Expression of the Menkes Protein. Plasmid, 1998, 39, 245-251.	1.4	16
95	Cloning, mapping and expression analysis of the sheep Wilson disease gene homologue. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2000, 1491, 229-239.	2.4	16
96	Expanding the phenotypic spectrum of ARID1B-mediated disorders and identification of altered cell-cycle dynamics due to ARID1B haploinsufficiency. Orphanet Journal of Rare Diseases, 2014, 9, 43.	2.7	16
97	Generation and characterisation of a parkin-Pacrg knockout mouse line and a Pacrg knockout mouse line. Scientific Reports, 2018, 8, 7528.	3.3	16
98	Germline variants in tumor suppressor FBXW7 lead to impaired ubiquitination and a neurodevelopmental syndrome. American Journal of Human Genetics, 2022, 109, 601-617.	6.2	16
99	Molecular analysis of the PArkin co-regulated gene and association with male infertility. Fertility and Sterility, 2010, 93, 2262-2268.	1.0	15
100	The COMT Val158 allele is associated with impaired delayed-match-to-sample performance in ADHD. Behavioral and Brain Functions, 2012, 8, 25.	3.3	15
101	Multiplication of the α-Synuclein Gene Is Not a Common Disease Mechanism in Lewy Body Disease. Journal of Molecular Neuroscience, 2004, 24, 337-342.	2.3	14
102	Clinical and Neuropathological Features Associated With Loss of RAB39B. Movement Disorders, 2020, 35, 687-693.	3.9	14
103	Cortical Dysplasia and the mTOR Pathway: How the Study of Human Brain Tissue Has Led to Insights into Epileptogenesis. International Journal of Molecular Sciences, 2022, 23, 1344.	4.1	14
104	Identification of the copper chaperone SAH in Ovis aries: expression analysis and in vitro interaction of SAH with ATP7B. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2000, 1490, 11-20.	2.4	12
105	Biochemical characterization of torsinB. Molecular Brain Research, 2004, 127, 1-9.	2.3	12
106	Rapid Diagnosis of Spinocerebellar Ataxia 36 in a <scp>Threeâ€Generation</scp> Family Using <scp>Shortâ€Read Wholeâ€Genome</scp> Sequencing Data. Movement Disorders, 2020, 35, 1675-1679.	3.9	12
107	Regional and cellular localisation of Parkin Co-Regulated Gene in developing and adult mouse brain. Brain Research, 2008, 1201, 177-186.	2.2	11
108	A novel <i>AMPD2</i> mutation outside the AMP deaminase domain causes pontocerebellar hypoplasia type 9. American Journal of Medical Genetics, Part A, 2017, 173, 820-823.	1.2	11

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109	Genetic Analysis of RAB39B in an Early-Onset Parkinson's Disease Cohort. Frontiers in Neurology, 2020, 11, 523.	2.4	11
110	Callosal agenesis and congenital mirror movements: outcomes associated with <i>DCC</i> mutations. Developmental Medicine and Child Neurology, 2020, 62, 758-762.	2.1	11
111	Implementation of ironXS: a study of the acceptability and feasibility of genetic screening for hereditary hemochromatosis in high schools. Clinical Genetics, 2010, 77, 241-248.	2.0	10
112	Severe Leukoencephalopathy with Clinical Recovery Caused by Recessive BOLA3 Mutations. JIMD Reports, 2018, 43, 63-70.	1.5	10
113	Intravenously delivered aminoglycoside antibiotics, tobramycin and amikacin, are not ototoxic in mice. Hearing Research, 2020, 386, 107870.	2.0	10
114	Functional analysis of the sheep Wilson disease protein (sATP7B) in CHO cells. European Journal of Cell Biology, 2001, 80, 349-357.	3.6	9
115	Long Range Regulation of Human FXN Gene Expression. PLoS ONE, 2011, 6, e22001.	2.5	9
116	Identification of a Novel RNF213 Variant in a Family with Heterogeneous Intracerebral Vasculopathy. International Journal of Stroke, 2014, 9, E26-E27.	5.9	9
117	HFE p.C282Y heterozygosity is associated with earlier disease onset in Friedreich ataxia. Movement Disorders, 2014, 29, 940-943.	3.9	9
118	Genetic Analysis of Patients Who Experienced Awareness with Recall while under General Anesthesia. Anesthesiology, 2019, 131, 974-982.	2.5	9
119	Chudley-McCullough Syndrome: A Recognizable Clinical Entity Characterized by Deafness and Typical Brain Malformations. Journal of Child Neurology, 2021, 36, 152-158.	1.4	9
120	Analysis of PArkin Co-Regulated Gene in a Taiwanese–Ethnic Chinese cohort with early-onset Parkinson's disease. Parkinsonism and Related Disorders, 2009, 15, 417-421.	2.2	8
121	Sporadic hypothalamic hamartoma is a ciliopathy with somatic and bi-allelic contributions. Human Molecular Genetics, 2022, 31, 2307-2316.	2.9	8
122	Heterozygous <scp><i>PNPT1</i></scp> Variants Cause Spinocerebellar Ataxia Type 25. Annals of Neurology, 2022, 92, 122-137.	5.3	8
123	Correction of the copper transport defect of Menkes patient fibroblasts by expression of two forms of the sheep Wilson ATPase. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2002, 1588, 189-194.	3.8	7
124	ironXS: high-school screening for hereditary haemochromatosis is acceptable and feasible. European Journal of Human Genetics, 2012, 20, 505-509.	2.8	7
125	Generation of RAB39B knockout isogenic human embryonic stem cell lines to model RAB39B-mediated Parkinson's disease. Stem Cell Research, 2018, 28, 161-164.	0.7	7
126	Familial early onset Parkinson's disease caused by a homozygous frameshift variant in PARK7: Clinical features and literature update. Parkinsonism and Related Disorders, 2019, 64, 308-311.	2.2	7

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127	Polymicrogyria associated with 17p13.3p13.2 duplication: Case report and review of the literature. European Journal of Medical Genetics, 2020, 63, 103774.	1.3	7
128	A DNA repair disorder caused by de novo monoallelic DDB1 variants is associated with a neurodevelopmental syndrome. American Journal of Human Genetics, 2021, 108, 749-756.	6.2	6
129	CUGC for pontocerebellar hypoplasia type 9 and spastic paraplegia-63. European Journal of Human Genetics, 2019, 27, 161-166.	2.8	5
130	DCC regulates astroglial development essential for telencephalic morphogenesis and corpus callosum formation. ELife, 2021, 10, .	6.0	5
131	Advancing the diagnosis of repeat expansion disorders. Lancet Neurology, The, 2022, 21, 205-207.	10.2	5
132	Generation of seven iPSC lines from peripheral blood mononuclear cells suitable to investigate Autism Spectrum Disorder. Stem Cell Research, 2019, 39, 101516.	0.7	4
133	Resection of tuber centers only for seizure control in tuberous sclerosis complex. Epilepsy Research, 2021, 171, 106572.	1.6	4
134	Functional Analysis of the Menkes Protein (MNK) Expressed from a cDNA Construct. Advances in Experimental Medicine and Biology, 1999, 448, 67-82.	1.6	4
135	Lack of evidence for association of a parkin promoter polymorphism with early-onset Parkinson's disease in a Chinese population. Parkinsonism and Related Disorders, 2009, 15, 149-152.	2.2	3
136	Rasmussen encephalitis tissue transfer program. Epilepsia, 2016, 57, 1005-1007.	5.1	3
137	Tracing Autism Traits in Large Multiplex Families to Identify Endophenotypes of the Broader Autism Phenotype. International Journal of Molecular Sciences, 2020, 21, 7965.	4.1	3
138	Clonally Focused Public and Private T Cells in Resected Brain Tissue From Surgeries to Treat Children With Intractable Seizures. Frontiers in Immunology, 2021, 12, 664344.	4.8	3
139	ASK1 is a novel molecular target for preventing aminoglycoside-induced hair cell death. Journal of Molecular Medicine, 2022, 100, 797-813.	3.9	3
140	Teaching Neurolmages: Imaging features of DCC-mediated mirror movements and isolated agenesis of the corpus callosum. Neurology, 2018, 91, e886-e887.	1.1	2
141	Expanding the clinical and radiological phenotypes of leukoencephalopathy due to biallelic <scp><i>HMBS</i></scp> mutations. American Journal of Medical Genetics, Part A, 2021, 185, 2941-2950.	1.2	2
142	A family study implicates <i>GBE1</i> in the etiology of autism spectrum disorder. Human Mutation, 2022, 43, 16-29.	2.5	2
143	Identification and validation of control cell lines for accurate parkin dosage analysis. Journal of Neuroscience Methods, 2009, 176, 68-71.	2.5	1
144	Generation of four iPSC lines from Neurofibromatosis Type 1 patients. Stem Cell Research, 2020, 49, 102013.	0.7	1

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145	Identification of a Novel Gene Linked to Parkin via a Bidirectional Promoter. Annals of the New York Academy of Sciences, 2003, 991, 311-314.	3.8	O
146	Clinical seizure manifestations in the absence of synaptic connections. Epileptic Disorders, 2021, 23, 167-172.	1.3	0