

Jozef Gecz

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6168622/publications.pdf>

Version: 2024-02-01

334
papers

24,700
citations

4641

85
h-index

11030

137
g-index

359
all docs

359
docs citations

359
times ranked

25039
citing authors

#	ARTICLE	IF	CITATIONS
1	Refining analyses of copy number variation identifies specific genes associated with developmental delay. <i>Nature Genetics</i> , 2014, 46, 1063-1071.	9.4	583
2	Duplication of the MECP2 Region Is a Frequent Cause of Severe Mental Retardation and Progressive Neurological Symptoms in Males. <i>American Journal of Human Genetics</i> , 2005, 77, 442-453.	2.6	550
3	A recurrent 16p12.1 microdeletion supports a two-hit model for severe developmental delay. <i>Nature Genetics</i> , 2010, 42, 203-209.	9.4	539
4	A systematic, large-scale resequencing screen of X-chromosome coding exons in mental retardation. <i>Nature Genetics</i> , 2009, 41, 535-543.	9.4	528
5	Targeted sequencing identifies 91 neurodevelopmental-disorder risk genes with autism and developmental-disability biases. <i>Nature Genetics</i> , 2017, 49, 515-526.	9.4	443
6	Mutations of CDKL5 Cause a Severe Neurodevelopmental Disorder with Infantile Spasms and Mental Retardation. <i>American Journal of Human Genetics</i> , 2004, 75, 1079-1093.	2.6	414
7	X-linked protocadherin 19 mutations cause female-limited epilepsy and cognitive impairment. <i>Nature Genetics</i> , 2008, 40, 776-781.	9.4	397
8	Mutations in the human ortholog of <i>Aristaless</i> cause X-linked mental retardation and epilepsy. <i>Nature Genetics</i> , 2002, 30, 441-445.	9.4	396
9	Mutations in the <i>JARID1C</i> Gene, Which Is Involved in Transcriptional Regulation and Chromatin Remodeling, Cause X-Linked Mental Retardation. <i>American Journal of Human Genetics</i> , 2005, 76, 227-236.	2.6	349
10	Identification of the gene <i>FMR2</i> , associated with <i>FRAAXE</i> mental retardation. <i>Nature Genetics</i> , 1996, 13, 105-108.	9.4	303
11	A novel X-linked trichothiodystrophy associated with a nonsense mutation in <i>RNF113A</i> . <i>Journal of Medical Genetics</i> , 2015, 52, 269-274.	1.5	302
12	Mutations in <i>DEPDC5</i> cause familial focal epilepsy with variable foci. <i>Nature Genetics</i> , 2013, 45, 546-551.	9.4	301
13	Cerebral palsy: causes, pathways, and the role of genetic variants. <i>American Journal of Obstetrics and Gynecology</i> , 2015, 213, 779-788.	0.7	290
14	<i>PHF6</i> mutations in T-cell acute lymphoblastic leukemia. <i>Nature Genetics</i> , 2010, 42, 338-342.	9.4	282
15	Identification of a MicroRNA that Activates Gene Expression by Repressing Nonsense-Mediated RNA Decay. <i>Molecular Cell</i> , 2011, 42, 500-510.	4.5	267
16	Disruption of the Serine/Threonine Kinase 9 Gene Causes Severe X-Linked Infantile Spasms and Mental Retardation. <i>American Journal of Human Genetics</i> , 2003, 72, 1401-1411.	2.6	265
17	Mutations in the X-Linked Cyclin-Dependent Kinase-“Like 5 (<i>CDKL5/STK9</i>) Gene Are Associated with Severe Neurodevelopmental Retardation. <i>American Journal of Human Genetics</i> , 2004, 75, 1149-1154.	2.6	264
18	<i>ARX</i> , a novel Prd-class-homeobox gene highly expressed in the telencephalon, is mutated in X-linked mental retardation. <i>Human Molecular Genetics</i> , 2002, 11, 981-991.	1.4	248

#	ARTICLE	IF	CITATIONS
19	De Novo Mutations in SLC1A2 and CACNA1A Are Important Causes of Epileptic Encephalopathies. American Journal of Human Genetics, 2016, 99, 287-298.	2.6	247
20	Common genetic variants contribute to risk of rare severe neurodevelopmental disorders. Nature, 2018, 562, 268-271.	13.7	246
21	X-exome sequencing of 405 unresolved families identifies seven novel intellectual disability genes. Molecular Psychiatry, 2016, 21, 133-148.	4.1	243
22	Disorders of sex development: insights from targeted gene sequencing of a large international patient cohort. Genome Biology, 2016, 17, 243.	3.8	241
23	PRRT2 Mutations Cause Benign Familial Infantile Epilepsy and Infantile Convulsions with Choreoathetosis Syndrome. American Journal of Human Genetics, 2012, 90, 152-160.	2.6	234
24	Mutations in DDX3X Are a Common Cause of Unexplained Intellectual Disability with Gender-Specific Effects on Wnt Signaling. American Journal of Human Genetics, 2015, 97, 343-352.	2.6	230
25	Mutations in UPF3B, a member of the nonsense-mediated mRNA decay complex, cause syndromic and nonsyndromic mental retardation. Nature Genetics, 2007, 39, 1127-1133.	9.4	228
26	Mutations in the Small GTPase Gene RAB39B Are Responsible for X-linked Mental Retardation Associated with Autism, Epilepsy, and Macrocephaly. American Journal of Human Genetics, 2010, 86, 185-195.	2.6	220
27	SLC9A6 Mutations Cause X-Linked Mental Retardation, Microcephaly, Epilepsy, and Ataxia, a Phenotype Mimicking Angelman Syndrome. American Journal of Human Genetics, 2008, 82, 1003-1010.	2.6	209
28	COMMD1 is linked to the WASH complex and regulates endosomal trafficking of the copper transporter ATP7A. Molecular Biology of the Cell, 2015, 26, 91-103.	0.9	200
29	Mutations in CUL4B, Which Encodes a Ubiquitin E3 Ligase Subunit, Cause an X-linked Mental Retardation Syndrome Associated with Aggressive Outbursts, Seizures, Relative Macrocephaly, Central Obesity, Hypogonadism, Pes Cavus, and Tremor. American Journal of Human Genetics, 2007, 80, 345-352.	2.6	197
30	Mutations in mammalian target of rapamycin regulator <i>DEPDC5</i> cause focal epilepsy with brain malformations. Annals of Neurology, 2014, 75, 782-787.	2.8	193
31	Mutations in PHF6 are associated with BÅrjesonâ€“Forssmanâ€“Lehmann syndrome. Nature Genetics, 2002, 32, 661-665.	9.4	192
32	The genetic landscape of intellectual disability arising from chromosome X. Trends in Genetics, 2009, 25, 308-316.	2.9	190
33	Identification of the gene (SEDL) causing X-linked spondyloepiphyseal dysplasia tarda. Nature Genetics, 1999, 22, 400-404.	9.4	188
34	Submicroscopic Duplications of the Hydroxysteroid Dehydrogenase HSD17B10 and the E3 Ubiquitin Ligase HUWE1 Are Associated with Mental Retardation. American Journal of Human Genetics, 2008, 82, 432-443.	2.6	187
35	Disruption at the <i>PTCHD1</i> Locus on Xp22.11 in Autism Spectrum Disorder and Intellectual Disability. Science Translational Medicine, 2010, 2, 49ra68.	5.8	178
36	Whole-exome sequencing points to considerable genetic heterogeneity of cerebral palsy. Molecular Psychiatry, 2015, 20, 176-182.	4.1	178

#	ARTICLE	IF	CITATIONS
37	Epilepsy and mental retardation limited to females: an under-recognized disorder. <i>Brain</i> , 2008, 131, 918-927.	3.7	172
38	Evaluation of DNA Methylation Episignatures for Diagnosis and Phenotype Correlations in 42 Mendelian Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2020, 106, 356-370.	2.6	171
39	Infantile spasms, dystonia, and other X-linked phenotypes caused by mutations in Aristaless related homeobox gene, ARX. <i>Brain and Development</i> , 2002, 24, 266-268.	0.6	170
40	A Mutation in the Golgi Qb-SNARE Gene GOSR2 Causes Progressive Myoclonus Epilepsy with Early Ataxia. <i>American Journal of Human Genetics</i> , 2011, 88, 657-663.	2.6	166
41	Rett syndrome: clinical review and genetic update. <i>Journal of Medical Genetics</i> , 2005, 42, 1-7.	1.5	161
42	Targeted Next-Generation Sequencing Analysis of 1,000 Individuals with Intellectual Disability. <i>Human Mutation</i> , 2015, 36, 1197-1204.	1.1	161
43	Early onset seizures and Rett-like features associated with mutations in CDKL5. <i>European Journal of Human Genetics</i> , 2005, 13, 1113-1120.	1.4	160
44	Mutations in the DLC3 Gene Cause Nonsyndromic X-Linked Mental Retardation. <i>American Journal of Human Genetics</i> , 2004, 75, 318-324.	2.6	157
45	XLMR genes: update 2007. <i>European Journal of Human Genetics</i> , 2008, 16, 422-434.	1.4	155
46	The original Lujan syndrome family has a novel missense mutation (p.N1007S) in the MED12 gene. <i>Journal of Medical Genetics</i> , 2007, 44, 472-477.	1.5	153
47	Hotspots of missense mutation identify neurodevelopmental disorder genes and functional domains. <i>Nature Neuroscience</i> , 2017, 20, 1043-1051.	7.1	152
48	ARX spectrum disorders: making inroads into the molecular pathology. <i>Human Mutation</i> , 2010, 31, 889-900.	1.1	151
49	Mutations in ZDHHC9, Which Encodes a Palmitoyltransferase of NRAS and HRAS, Cause X-Linked Mental Retardation Associated with a Marfanoid Habitus. <i>American Journal of Human Genetics</i> , 2007, 80, 982-987.	2.6	150
50	La FAM fatale: USP9X in development and disease. <i>Cellular and Molecular Life Sciences</i> , 2015, 72, 2075-2089.	2.4	145
51	A missense mutation in RPS6KA3 (RSK2) responsible for non-specific mental retardation. <i>Nature Genetics</i> , 1999, 22, 13-14.	9.4	142
52	Disruptive de novo mutations of DYRK1A lead to a syndromic form of autism and ID. <i>Molecular Psychiatry</i> , 2016, 21, 126-132.	4.1	142
53	Mutations in the polyglutamine binding protein 1 gene cause X-linked mental retardation. <i>Nature Genetics</i> , 2003, 35, 313-315.	9.4	139
54	Mutations in the guanine nucleotide exchange factor gene IQSEC2 cause nonsyndromic intellectual disability. <i>Nature Genetics</i> , 2010, 42, 486-488.	9.4	134

#	ARTICLE	IF	CITATIONS
55	Disruption of POGZ Is Associated with Intellectual Disability and Autism Spectrum Disorders. American Journal of Human Genetics, 2016, 98, 541-552.	2.6	132
56	Oligosaccharyltransferase-Subunit Mutations in Nonsyndromic Mental Retardation. American Journal of Human Genetics, 2008, 82, 1150-1157.	2.6	130
57	Structural variation in Xq28: MECP2 duplications in 1% of patients with unexplained XLMR and in 2% of male patients with severe encephalopathy. European Journal of Human Genetics, 2009, 17, 444-453.	1.4	130
58	Nonsense-mediated mRNA decay: Inter-individual variability and human disease. Neuroscience and Biobehavioral Reviews, 2014, 46, 175-186.	2.9	130
59	Functional characterization of GATA3 mutations causing the hypoparathyroidism-deafness-renal (HDR) dysplasia syndrome: insight into mechanisms of DNA binding by the GATA3 transcription factor. Human Molecular Genetics, 2006, 16, 265-275.	1.4	129
60	“North Sea” progressive myoclonus epilepsy: phenotype of subjects with GOSR2 mutation. Brain, 2013, 136, 1146-1154.	3.7	129
61	YY1 Haploinsufficiency Causes an Intellectual Disability Syndrome Featuring Transcriptional and Chromatin Dysfunction. American Journal of Human Genetics, 2017, 100, 907-925.	2.6	125
62	The genetic basis of cerebral palsy. Developmental Medicine and Child Neurology, 2017, 59, 462-469.	1.1	125
63	ARX: a gene for all seasons. Current Opinion in Genetics and Development, 2006, 16, 308-316.	1.5	123
64	Defects in tRNA Anticodon Loop 2'-O-Methylation Are Implicated in Nonsyndromic X-Linked Intellectual Disability due to Mutations in FTSJ1. Human Mutation, 2015, 36, 1176-1187.	1.1	122
65	Contribution of copy number variants involving nonsense-mediated mRNA decay pathway genes to neuro-developmental disorders. Human Molecular Genetics, 2013, 22, 1816-1825.	1.4	120
66	Mutations in the FTSJ1 Gene Coding for a Novel S-Adenosylmethionine-Binding Protein Cause Nonsyndromic X-Linked Mental Retardation. American Journal of Human Genetics, 2004, 75, 305-309.	2.6	117
67	Mutations in USP9X Are Associated with X-Linked Intellectual Disability and Disrupt Neuronal Cell Migration and Growth. American Journal of Human Genetics, 2014, 94, 470-478.	2.6	117
68	Mutations of the UPF3B gene, which encodes a protein widely expressed in neurons, are associated with nonspecific mental retardation with or without autism. Molecular Psychiatry, 2010, 15, 767-776.	4.1	113
69	Biallelic SUN5 Mutations Cause Autosomal-Recessive Acephalic Spermatozoa Syndrome. American Journal of Human Genetics, 2016, 99, 942-949.	2.6	113
70	Variable expression of mental retardation, autism, seizures, and dystonic hand movements in two families with an identical ARX gene mutation. American Journal of Medical Genetics Part A, 2002, 112, 405-411.	2.4	111
71	A Focal Epilepsy and Intellectual Disability Syndrome Is Due to a Mutation in TBC1D24. American Journal of Human Genetics, 2010, 87, 371-375.	2.6	111
72	Unstable TTTTA/TTTCA expansions in MARCH6 are associated with Familial Adult Myoclonic Epilepsy type 3. Nature Communications, 2019, 10, 4919.	5.8	111

#	ARTICLE	IF	CITATIONS
73	The Koolen-de Vries syndrome: a phenotypic comparison of patients with a 17q21.31 microdeletion versus a KANSL1 sequence variant. <i>European Journal of Human Genetics</i> , 2016, 24, 652-659.	1.4	108
74	Mutations in a Novel Gene, NHS, Cause the Pleiotropic Effects of Nance-Horan Syndrome, Including Severe Congenital Cataract, Dental Anomalies, and Mental Retardation. <i>American Journal of Human Genetics</i> , 2003, 73, 1120-1130.	2.6	107
75	Fine-Scale Survey of X Chromosome Copy Number Variants and Indels Underlying Intellectual Disability. <i>American Journal of Human Genetics</i> , 2010, 87, 173-188.	2.6	107
76	A UPF3-mediated regulatory switch that maintains RNA surveillance. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 747-753.	3.6	106
77	CASK mutations are frequent in males and cause X-linked nystagmus and variable XLMR phenotypes. <i>European Journal of Human Genetics</i> , 2010, 18, 544-552.	1.4	105
78	Large-scale targeted sequencing identifies risk genes for neurodevelopmental disorders. <i>Nature Communications</i> , 2020, 11, 4932.	5.8	105
79	Mutation frequencies of X-linked mental retardation genes in families from the EuroMRX consortium. <i>Human Mutation</i> , 2007, 28, 207-208.	1.1	103
80	Mutations in the Gene Encoding the Sigma 2 Subunit of the Adaptor Protein 1 Complex, AP1S2, Cause X-Linked Mental Retardation. <i>American Journal of Human Genetics</i> , 2006, 79, 1119-1124.	2.6	102
81	XNP mutation in a large family with Juberg-Marsidi syndrome. <i>Nature Genetics</i> , 1996, 12, 359-360.	9.4	101
82	The UPF3B gene, implicated in intellectual disability, autism, ADHD and childhood onset schizophrenia regulates neural progenitor cell behaviour and neuronal outgrowth. <i>Human Molecular Genetics</i> , 2013, 22, 4673-4687.	1.4	101
83	CCDC22 deficiency in humans blunts activation of proinflammatory NF- κ B signaling. <i>Journal of Clinical Investigation</i> , 2013, 123, 2244-2256.	3.9	101
84	Mutations in ionotropic AMPA receptor 3 alter channel properties and are associated with moderate cognitive impairment in humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 18163-18168.	3.3	100
85	Abnormal Cell Sorting Underlies the Unique X-Linked Inheritance of PCDH19 Epilepsy. <i>Neuron</i> , 2018, 97, 59-66.e5.	3.8	100
86	Intronic ATTTC repeat expansions in STARD7 in familial adult myoclonic epilepsy linked to chromosome 2. <i>Nature Communications</i> , 2019, 10, 4920.	5.8	99
87	Transcriptome profiling of UPF3B/NMD-deficient lymphoblastoid cells from patients with various forms of intellectual disability. <i>Molecular Psychiatry</i> , 2012, 17, 1103-1115.	4.1	97
88	<i>TBC1D24</i> genotype-phenotype correlation. <i>Neurology</i> , 2016, 87, 77-85.	1.5	97
89	Mutations disrupting neuritogenesis genes confer risk for cerebral palsy. <i>Nature Genetics</i> , 2020, 52, 1046-1056.	9.4	96
90	De Novo Loss-of-Function Mutations in USP9X Cause a Female-Specific Recognizable Syndrome with Developmental Delay and Congenital Malformations. <i>American Journal of Human Genetics</i> , 2016, 98, 373-381.	2.6	95

#	ARTICLE	IF	CITATIONS
91	Mutations of protocadherin 19 in female epilepsy (PCDH19-FE) lead to allopregnanolone deficiency. <i>Human Molecular Genetics</i> , 2015, 24, 5250-5259.	1.4	93
92	The Molecular Basis of X-Linked Spondyloepiphyseal Dysplasia Tarda. <i>American Journal of Human Genetics</i> , 2001, 68, 1386-1397.	2.6	90
93	A Noncoding, Regulatory Mutation Implicates HCFC1 in Nonsyndromic Intellectual Disability. <i>American Journal of Human Genetics</i> , 2012, 91, 694-702.	2.6	89
94	X-linked myoclonic epilepsy with spasticity and intellectual disability. <i>Neurology</i> , 2002, 59, 348-356.	1.5	88
95	Identification and characterization of a missense mutation in the O-linked Ñ ² -N-acetylglucosamine (O-GlcNAc) transferase gene that segregates with X-linked intellectual disability. <i>Journal of Biological Chemistry</i> , 2017, 292, 8948-8963.	1.6	87
96	De Novo Pathogenic Variants in CACNA1E Cause Developmental and Epileptic Encephalopathy with Contractures, Macrocephaly, and Dyskinesias. <i>American Journal of Human Genetics</i> , 2018, 103, 666-678.	2.6	87
97	A systematic review and meta-analysis of 271 PCDH19-variant individuals identifies psychiatric comorbidities, and association of seizure onset and disease severity. <i>Molecular Psychiatry</i> , 2019, 24, 241-251.	4.1	86
98	Mutations in the BRWD3 Gene Cause X-Linked Mental Retardation Associated with Macrocephaly. <i>American Journal of Human Genetics</i> , 2007, 81, 367-374.	2.6	85
99	Characterization of the Human Glutamate Receptor Subunit 3 Gene (GRIA3), a Candidate for Bipolar Disorder and Nonspecific X-Linked Mental Retardation. <i>Genomics</i> , 1999, 62, 356-368.	1.3	84
100	A ubiquitin-dependent signalling axis specific for ALKBH-mediated DNA dealkylation repair. <i>Nature</i> , 2017, 551, 389-393.	13.7	83
101	THOC2 Mutations Implicate mRNA-Export Pathway in X-Linked Intellectual Disability. <i>American Journal of Human Genetics</i> , 2015, 97, 302-310.	2.6	82
102	Genetic or Other Causation Should Not Change the Clinical Diagnosis of Cerebral Palsy. <i>Journal of Child Neurology</i> , 2019, 34, 472-476.	0.7	82
103	<i>FOXP1</i> mutations cause intellectual disability and a recognizable phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 3166-3175.	0.7	79
104	Loss of Usp9x Disrupts Cortical Architecture, Hippocampal Development and TGFÑ ² -Mediated Axonogenesis. <i>PLoS ONE</i> , 2013, 8, e68287.	1.1	77
105	Mouse orthologue of ARX, a gene mutated in several X-linked forms of mental retardation and epilepsy, is a marker of adult neural stem cells and forebrain GABAergic neurons. <i>Developmental Dynamics</i> , 2004, 231, 631-639.	0.8	76
106	Two novel JAK2 exon 12 mutations in JAK2V617F-negative polycythaemia vera patients. <i>Leukemia</i> , 2008, 22, 870-873.	3.3	76
107	Fibroblast growth factor homologous factor 2 (FHF2): gene structure, expression and mapping to the BÅ ^r rjeson-Forssman-Lehmann syndrome region in Xq26 delineated by a duplication breakpoint in a BFLS-like patient. <i>Human Genetics</i> , 1999, 104, 56-63.	1.8	75
108	ZNF674: A New KrÃ ¹ 4ppel-Associated Boxâ€“Containing Zinc-Finger Gene Involved in Nonsyndromic X-Linked Mental Retardation. <i>American Journal of Human Genetics</i> , 2006, 78, 265-278.	2.6	75

#	ARTICLE	IF	CITATIONS
109	Epilepsy and mental retardation limited to females with PCDH19 mutations can present de novo or in single generation families. <i>Journal of Medical Genetics</i> , 2010, 47, 211-216.	1.5	74
110	Evolution of the human X â€“ a smart and sexy chromosome that controls speciation and development. <i>Cytogenetic and Genome Research</i> , 2002, 99, 141-145.	0.6	72
111	Dominant <i>KCNA2</i> mutation causes episodic ataxia and pharmacoresponsive epilepsy. <i>Neurology</i> , 2016, 87, 1975-1984.	1.5	71
112	Mutations in the intellectual disability gene <i>KDM5C</i> reduce protein stability and demethylase activity. <i>Human Molecular Genetics</i> , 2015, 24, 2861-2872.	1.4	69
113	<i>ZC4H2</i> Mutations Are Associated with Arthrogryposis Multiplex Congenita and Intellectual Disability through Impairment of Central and Peripheral Synaptic Plasticity. <i>American Journal of Human Genetics</i> , 2013, 92, 681-695.	2.6	68
114	Severe childhood speech disorder. <i>Neurology</i> , 2020, 94, e2148-e2167.	1.5	68
115	Genes for Cognitive Function: Developments on the X. <i>Genome Research</i> , 2000, 10, 157-163.	2.4	67
116	FRAXE-associated mental retardation protein (<i>FMR2</i>) is an RNA-binding protein with high affinity for G-quartet RNA forming structure. <i>Nucleic Acids Research</i> , 2009, 37, 1269-1279.	6.5	67
117	Identification and characterization of two novel <i>JARID1C</i> mutations: suggestion of an emerging genotypeâ€“phenotype correlation. <i>European Journal of Human Genetics</i> , 2010, 18, 330-335.	1.4	66
118	Seizures Are Regulated by Ubiquitin-specific Peptidase 9 X-linked (<i>USP9X</i>), a De-Ubiquitinase. <i>PLoS Genetics</i> , 2015, 11, e1005022.	1.5	66
119	Nonsyndromic X-linked mental retardation: where are the missing mutations?. <i>Trends in Genetics</i> , 2003, 19, 316-320.	2.9	65
120	Rare copy number variation in cerebral palsy. <i>European Journal of Human Genetics</i> , 2014, 22, 40-45.	1.4	65
121	Gene Structure and Subcellular Localization of <i>FMR2</i> , a Member of a New Family of Putative Transcription Activators. <i>Genomics</i> , 1997, 44, 201-213.	1.3	64
122	Hypomorphic Temperature-Sensitive Alleles of <i>NSDHL</i> Cause CK Syndrome. <i>American Journal of Human Genetics</i> , 2010, 87, 905-914.	2.6	64
123	The clinical picture of the BÅ“rjeson-Forsman-Lehmann syndrome in males and heterozygous females with <i>PHF6</i> mutations. <i>Clinical Genetics</i> , 2004, 65, 226-232.	1.0	63
124	Functional characterization of the AFF (<i>AF4/FMR2</i>) family of RNA-binding proteins: insights into the molecular pathology of FRAXE intellectual disability. <i>Human Molecular Genetics</i> , 2011, 20, 1873-1885.	1.4	63
125	Aristaless-related homeobox gene, the gene responsible for West syndrome and related disorders, is a Groucho/transducin-like enhancer of split dependent transcriptional repressor. <i>Neuroscience</i> , 2007, 146, 236-247.	1.1	62
126	Mutation detection in <i>FGFR2</i> craniosynostosis syndromes. <i>Human Genetics</i> , 1997, 99, 251-255.	1.8	60

#	ARTICLE	IF	CITATIONS
127	X-linked mild non-syndromic mental retardation with neuropsychiatric problems and the missense mutation A365E in PAK3. , 2003, 120A, 509-517.		60
128	Disruption of a new X linked gene highly expressed in brain in a family with two mentally retarded males. Journal of Medical Genetics, 2004, 41, 736-742.	1.5	60
129	Folate-sensitive fragile site FRA10A is due to an expansion of a CGG repeat in a novel gene, FRA10AC1, encoding a nuclear protein. Genomics, 2004, 84, 69-81.	1.3	60
130	Truncating Variants in NAA15 Are Associated with Variable Levels of Intellectual Disability, Autism Spectrum Disorder, and Congenital Anomalies. American Journal of Human Genetics, 2018, 102, 985-994.	2.6	59
131	Cloning and characterization of a new human Xq13 gene, encoding a putative helicase. Human Molecular Genetics, 1994, 3, 1957-1964.	1.4	58
132	Three new families with X-linked mental retardation caused by the 428-451dup(24bp) mutation in ARX. Clinical Genetics, 2004, 66, 39-45.	1.0	58
133	MCT8 mutation analysis and identification of the first female with Allanâ€“Herndonâ€“Dudley syndrome due to loss of MCT8 expression. European Journal of Human Genetics, 2008, 16, 1029-1037.	1.4	56
134	O-GlcNAc transferase missense mutations linked to X-linked intellectual disability deregulate genes involved in cell fate determination and signaling. Journal of Biological Chemistry, 2018, 293, 10810-10824.	1.6	56
135	Splicing mutation in the ATR-X gene can lead to a dysmorphic mental retardation phenotype without alpha-thalassemia. American Journal of Human Genetics, 1996, 58, 499-505.	2.6	56
136	Fragile XE-associated familial mental retardation protein 2 (FMR2) acts as a potent transcription activator. Journal of Human Genetics, 2001, 46, 251-259.	1.1	55
137	Disruptions of the novel KIAA1202 gene are associated with X-linked mental retardation. Human Genetics, 2006, 118, 578-590.	1.8	55
138	A Upf3b-mutant mouse model with behavioral and neurogenesis defects. Molecular Psychiatry, 2018, 23, 1773-1786.	4.1	54
139	FMR2 Expression in Families with Fraxe Mental Retardation. Human Molecular Genetics, 1997, 6, 435-441.	1.4	53
140	â€“Big issuesâ€“™ in neurodevelopment for children and adults with congenital heart disease. Open Heart, 2019, 6, e000998.	0.9	53
141	Expanding the molecular basis and phenotypic spectrum of X-linked Joubert syndrome associated with OFD1 mutations. European Journal of Human Genetics, 2012, 20, 806-809.	1.4	52
142	Pcdh19 Loss-of-Function Increases Neuronal Migration In Vitro but is Dispensable for Brain Development in Mice. Scientific Reports, 2016, 6, 26765.	1.6	52
143	New insights into Brunner syndrome and potential for targeted therapy. Clinical Genetics, 2016, 89, 120-127.	1.0	52
144	CCDC22: a novel candidate gene for syndromic X-linked intellectual disability. Molecular Psychiatry, 2012, 17, 4-7.	4.1	50

#	ARTICLE	IF	CITATIONS
145	Rare deleterious mutations of HNRNP genes result in shared neurodevelopmental disorders. <i>Genome Medicine</i> , 2021, 13, 63.	3.6	50
146	The FMR2 gene, FRAXE and non-specific X-linked mental retardation: clinical and molecular aspects. <i>Annals of Human Genetics</i> , 2000, 64, 95-106.	0.3	49
147	A Recurrent RNA-Splicing Mutation in the SEDL Gene Causes X-Linked Spondyloepiphyseal Dysplasia Tarda. <i>American Journal of Human Genetics</i> , 2001, 68, 1398-1407.	2.6	49
148	HCFC1 loss-of-function mutations disrupt neuronal and neural progenitor cells of the developing brain. <i>Human Molecular Genetics</i> , 2015, 24, 3335-3347.	1.4	47
149	Cloning and expression of the murine homologue of a putative human X-linked nuclear protein gene closely linked to PCK1 in Xq13.3. <i>Human Molecular Genetics</i> , 1994, 3, 39-44.	1.4	46
150	Loss of SLC38A5 and FTSJ1 at Xp11.23 in three brothers with non-syndromic mental retardation due to a microdeletion in an unstable genomic region. <i>Human Genetics</i> , 2007, 121, 539-547.	1.8	46
151	Interchromosomal Insertional Translocation at Xq26.3 Alters SOX3 Expression in an Individual With XX Male Sex Reversal. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E815-E820.	1.8	46
152	Gene Structure and Expression Study of the SEDL Gene for Spondyloepiphyseal Dysplasia Tarda. <i>Genomics</i> , 2000, 69, 242-251.	1.3	45
153	The Börjeson-Forssman-Lehman syndrome (BFLS, MIM #301900). <i>European Journal of Human Genetics</i> , 2006, 14, 1233-1237.	1.4	45
154	Protein and gene expression analysis of Phf6, the gene mutated in the Börjeson-Forssman-Lehmann Syndrome of intellectual disability and obesity. <i>Gene Expression Patterns</i> , 2007, 7, 858-871.	0.3	45
155	Eight further individuals with intellectual disability and epilepsy carrying bi-allelic CNTNAP2 aberrations allow delineation of the mutational and phenotypic spectrum. <i>Journal of Medical Genetics</i> , 2016, 53, 820-827.	1.5	45
156	De novo and inherited mutations in the X-linked gene CLCN4 are associated with syndromic intellectual disability and behavior and seizure disorders in males and females. <i>Molecular Psychiatry</i> , 2018, 23, 222-230.	4.1	45
157	Mutation screening in Borjeson-Forssman-Lehmann syndrome: identification of a novel de novo PHF6 mutation in a female patient. <i>Journal of Medical Genetics</i> , 2005, 43, 238-243.	1.5	43
158	Xp11.2 microduplications including IQSEC2, TSPYL2 and KDM5C genes in patients with neurodevelopmental disorders. <i>European Journal of Human Genetics</i> , 2016, 24, 373-380.	1.4	43
159	Disruptive mutations in TANC2 define a neurodevelopmental syndrome associated with psychiatric disorders. <i>Nature Communications</i> , 2019, 10, 4679.	5.8	43
160	Inhibition of Upf2-Dependent Nonsense-Mediated Decay Leads to Behavioral and Neurophysiological Abnormalities by Activating the Immune Response. <i>Neuron</i> , 2019, 104, 665-679.e8.	3.8	43
161	Molecular pathology of expanded polyalanine tract mutations in the Aristaless-related homeobox gene. <i>Genomics</i> , 2007, 90, 59-71.	1.3	42
162	Novel mutations in NLGN3 causing autism spectrum disorder and cognitive impairment. <i>Human Mutation</i> , 2019, 40, 2021-2032.	1.1	42

#	ARTICLE	IF	CITATIONS
163	Partial Loss of USP9X Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor β Signaling. <i>Biological Psychiatry</i> , 2020, 87, 100-112.	0.7	42
164	Novel diagnostic DNA methylation epesignatures expand and refine the epigenetic landscapes of Mendelian disorders. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100075.	1.0	42
165	FRA2A Is a CGG Repeat Expansion Associated with Silencing of AFF3. <i>PLoS Genetics</i> , 2014, 10, e1004242.	1.5	41
166	Nanceâ€™s Horan syndrome protein, NHS, associates with epithelial cell junctions. <i>Human Molecular Genetics</i> , 2006, 15, 1972-1983.	1.4	40
167	Reduced steroidogenesis in patients with <sc>PCDH</sc>19â€™female limited epilepsy. <i>Epilepsia</i> , 2017, 58, e91-e95.	2.6	40
168	PHF6 regulates hematopoietic stem and progenitor cells and its loss synergizes with expression of TLX3 to cause leukemia. <i>Blood</i> , 2019, 133, 1729-1741.	0.6	40
169	A Survey of Rare Epigenetic Variation in 23,116 Human Genomes Identifies Disease-Relevant Epivariations and CGG Expansions. <i>American Journal of Human Genetics</i> , 2020, 107, 654-669.	2.6	40
170	Identification of three novel SEDL mutations, including mutation in the rare, non-canonical splice site of exon 4. <i>Clinical Genetics</i> , 2003, 64, 235-242.	1.0	39
171	A Regulatory Path Associated with X-Linked Intellectual Disability and Epilepsy Links KDM5C to the Polyalanine Expansions in ARX. <i>American Journal of Human Genetics</i> , 2013, 92, 114-125.	2.6	39
172	HUWE1 mutations in Juberg-Marsidi and Brooks syndromes: the results of an X-chromosome exome sequencing study. <i>BMJ Open</i> , 2016, 6, e009537.	0.8	39
173	Targeted knockout of a chemokine-like gene increases anxiety and fear responses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E1041-E1050.	3.3	39
174	PCDH19 regulation of neural progenitor cell differentiation suggests asynchrony of neurogenesis as a mechanism contributing to PCDH19 Girls Clustering Epilepsy. <i>Neurobiology of Disease</i> , 2018, 116, 106-119.	2.1	39
175	An intellectual disability syndrome with single-nucleotide variants in O-GlcNAc transferase. <i>European Journal of Human Genetics</i> , 2020, 28, 706-714.	1.4	38
176	Identification of a mutation in the XNP/ATR-X gene in a family reported as Smith-Fineman-Myers syndrome. , 2000, 91, 83-85.		37
177	Overlapping submicroscopic deletions in Xq28 in two unrelated boys with developmental disorders: identification of a gene near FRAXE. <i>American Journal of Human Genetics</i> , 1995, 56, 907-14.	2.6	37
178	A distinctive gene expression fingerprint in mentally retarded male patients reflects disease-causing defects in the histone demethylase KDM5C. <i>PathoGenetics</i> , 2010, 3, 2.	5.7	35
179	Disruptive variants of <i>CSDE1</i> associate with autism and interfere with neuronal development and synaptic transmission. <i>Science Advances</i> , 2019, 5, eaax2166.	4.7	35
180	Cyclin-Dependent Kinase-Like 5 (<i>CDKL5</i>) Mutation Screening in Rett Syndrome and Related Disorders. <i>Twin Research and Human Genetics</i> , 2010, 13, 168-178.	0.3	34

#	ARTICLE	IF	CITATIONS
181	Mutations in the nuclear localization sequence of the Aristaless related homeobox; sequestration of mutant ARX with IPO13 disrupts normal subcellular distribution of the transcription factor and retards cell division. <i>PathoGenetics</i> , 2010, 3, 1.	5.7	33
182	Human wild-type SEDL protein functionally complements yeast Trs20p but some naturally occurring SEDL mutants do not. <i>Gene</i> , 2003, 320, 137-144.	1.0	32
183	A genotype-first approach identifies an intellectual disability-overweight syndrome caused by PHIP haploinsufficiency. <i>European Journal of Human Genetics</i> , 2018, 26, 54-63.	1.4	32
184	Subtle functional defects in the Arf-specific guanine nucleotide exchange factor IQSEC2 cause non-syndromic X-linked intellectual disability. <i>Small GTPases</i> , 2010, 1, 98-103.	0.7	31
185	Novel IL1RAPL1 mutations associated with intellectual disability impair synaptogenesis. <i>Human Molecular Genetics</i> , 2015, 24, 1106-1118.	1.4	31
186	Pathogenic copy number variants that affect gene expression contribute to genomic burden in cerebral palsy. <i>Npj Genomic Medicine</i> , 2018, 3, 33.	1.7	31
187	Broadening the phenotype associated with mutations in UPF3B: Two further cases with renal dysplasia and variable developmental delay. <i>European Journal of Medical Genetics</i> , 2012, 55, 476-479.	0.7	30
188	Identification of an <i>IGSF1</i> -specific deletion in a five-generation pedigree with X-linked Central Hypothyroidism without macroorchidism. <i>Clinical Endocrinology</i> , 2016, 85, 609-615.	1.2	30
189	Ohtahara syndrome in a family with an ARX protein truncation mutation (c.81C>G/p.Y27X). <i>European Journal of Human Genetics</i> , 2010, 18, 157-162.	1.4	29
190	Identity by descent fine mapping of familial adult myoclonus epilepsy (FAME) to 2p11.2–2q11.2. <i>Human Genetics</i> , 2016, 135, 1117-1125.	1.8	29
191	Choreoathetosis, congenital hypothyroidism and neonatal respiratory distress syndrome with intact <i>NKX2-1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 3168-3173.	0.7	28
192	TBC1D24 mutation associated with focal epilepsy, cognitive impairment and a distinctive cerebro-cerebellar malformation. <i>Epilepsy Research</i> , 2013, 105, 240-244.	0.8	28
193	Increased <i>STAG2</i> dosage defines a novel cohesinopathy with intellectual disability and behavioral problems. <i>Human Molecular Genetics</i> , 2015, 24, 7171-7181.	1.4	28
194	Protocadherin 19 (PCDH19) interacts with paraspeckle protein NONO to co-regulate gene expression with estrogen receptor alpha (ER α). <i>Human Molecular Genetics</i> , 2017, 26, 2042-2052.	1.4	28
195	Severe neurocognitive and growth disorders due to variation in <i>THOC2</i> , an essential component of nuclear mRNA export machinery. <i>Human Mutation</i> , 2018, 39, 1126-1138.	1.1	28
196	In-depth analysis reveals complex molecular aetiology in a cohort of idiopathic cerebral palsy. <i>Brain</i> , 2022, 145, 119-141.	3.7	28
197	Abnormal expression of the KLF8 (ZNF741) gene in a female patient with an X;autosome translocation t(X;21)(p11.2;q22.3) and non-syndromic mental retardation. <i>Journal of Medical Genetics</i> , 2002, 39, 113-117.	1.5	27
198	USP9X deubiquitylating enzyme maintains RAPTOR protein levels, mTORC1 signalling and proliferation in neural progenitors. <i>Scientific Reports</i> , 2017, 7, 391.	1.6	27

#	ARTICLE	IF	CITATIONS
199	Novel PHF6 mutation p.D333del causes Borjeson-Forssman-Lehmann syndrome. <i>Journal of Medical Genetics</i> , 2003, 40, 50e-50.	1.5	26
200	Pathogenic variants in E3 ubiquitin ligase RLIM/RNF12 lead to a syndromic X-linked intellectual disability and behavior disorder. <i>Molecular Psychiatry</i> , 2019, 24, 1748-1768.	4.1	26
201	Novel causative mutations in patients with Nance-Horan syndrome and altered localization of the mutant NHS-A protein isoform. <i>Molecular Vision</i> , 2008, 14, 1856-64.	1.1	26
202	Glutamate receptors and learning and memory. <i>Nature Genetics</i> , 2010, 42, 925-926.	9.4	25
203	Homozygous mutation of STXBP5L explains an autosomal recessive infantile-onset neurodegenerative disorder. <i>Human Molecular Genetics</i> , 2015, 24, 2000-2010.	1.4	25
204	A recurrent missense variant in SLC9A7 causes nonsyndromic X-linked intellectual disability with alteration of Golgi acidification and aberrant glycosylation. <i>Human Molecular Genetics</i> , 2019, 28, 598-614.	1.4	25
205	Spondyloepiphyseal dysplasia tarda (SEDL, MIM #313400). <i>European Journal of Human Genetics</i> , 2003, 11, 639-642.	1.4	24
206	1024C>T (R342X) is a recurrent PHF6 mutation also found in the original Borjeson-Forssman-Lehmann syndrome family. <i>European Journal of Human Genetics</i> , 2004, 12, 787-789.	1.4	24
207	MED12-related XLID disorders are dose-dependent of immediate early genes (IEGs) expression. <i>Human Molecular Genetics</i> , 2017, 26, 2062-2075.	1.4	24
208	Disentangling the paradox of the PCDH19 clustering epilepsy, a disorder of cellular mosaics. <i>Current Opinion in Genetics and Development</i> , 2020, 65, 169-175.	1.5	24
209	Mutation screening of brain-expressed X-chromosomal miRNA genes in 464 patients with nonsyndromic X-linked mental retardation. <i>European Journal of Human Genetics</i> , 2007, 15, 375-378.	1.4	23
210	Lessons learnt from large-scale exon re-sequencing of the X chromosome. <i>Human Molecular Genetics</i> , 2009, 18, R60-R64.	1.4	23
211	Screening and cell-based assessment of mutations in the Aristaless-related homeobox (ARX) gene. <i>Clinical Genetics</i> , 2011, 80, 510-522.	1.0	23
212	Multiplex families with epilepsy. <i>Neurology</i> , 2016, 86, 713-722.	1.5	23
213	A Recurrent De Novo Nonsense Variant in ZSWIM6 Results in Severe Intellectual Disability without Frontonasal or Limb Malformations. <i>American Journal of Human Genetics</i> , 2017, 101, 995-1005.	2.6	23
214	Clinical and functional characterization of recurrent missense variants implicated in THOC6-related intellectual disability. <i>Human Molecular Genetics</i> , 2019, 28, 952-960.	1.4	23
215	Familial adult myoclonic epilepsy type 1 SAMD12 TTCA repeat expansion arose 17,000 years ago and is present in Sri Lankan and Indian families. <i>European Journal of Human Genetics</i> , 2020, 28, 973-978.	1.4	23
216	The FMR2 gene, FRAXE and non-specific X-linked mental retardation: clinical and molecular aspects. <i>Annals of Human Genetics</i> , 2000, 64, 95-106.	0.3	23

#	ARTICLE	IF	CITATIONS
217	Assignment of a polycomb-like chromobox gene (CBX2) to human chromosome 17q25. <i>Genomics</i> , 1995, 26, 130-133.	1.3	22
218	Characterisation and expression of a large, 13.7 kb FMR2 isoform.. <i>European Journal of Human Genetics</i> , 1999, 7, 157-162.	1.4	22
219	NHS-A isoform of the NHS gene is a novel interactor of ZO-1. <i>Experimental Cell Research</i> , 2009, 315, 2358-2372.	1.2	22
220	Analysis of 182 cerebral palsy transcriptomes points to dysregulation of trophic signalling pathways and overlap with autism. <i>Translational Psychiatry</i> , 2018, 8, 88.	2.4	22
221	Targeted resequencing identifies genes with recurrent variation in cerebral palsy. <i>Npj Genomic Medicine</i> , 2019, 4, 27.	1.7	22
222	A new microdeletion syndrome involving TBC1D24, ATP6VOC, and PDPK1 causes epilepsy, microcephaly, and developmental delay. <i>Genetics in Medicine</i> , 2019, 21, 1058-1064.	1.1	22
223	A standardized patient-centered characterization of the phenotypic spectrum of PCDH19 girls clustering epilepsy. <i>Translational Psychiatry</i> , 2020, 10, 127.	2.4	22
224	A novel gene, FAM11A, associated with the FRAXF CpG island is transcriptionally silent in FRAXF full mutation. <i>European Journal of Human Genetics</i> , 2002, 10, 767-772.	1.4	21
225	The molecular basis of intellectual disability: novel genes with naturally occurring mutations causing altered gene expression in the brain. <i>Frontiers in Bioscience - Landmark</i> , 2004, 9, 1.	3.0	21
226	Hybridisation-based resequencing of 17 X-linked intellectual disability genes in 135 patients reveals novel mutations in ATRX, SLC6A8 and PQBP1. <i>European Journal of Human Genetics</i> , 2011, 19, 717-720.	1.4	21
227	Reduced polyalanine-expanded Arx mutant protein in developing mouse subpallium alters Lmo1 transcriptional regulation. <i>Human Molecular Genetics</i> , 2014, 23, 1084-1094.	1.4	20
228	XLMR in MRX families 29, 32, 33 and 38 results from the dup24 mutation in the ARX (Aristaless related) Tj ETQq0 0,0 rgBT /Overlock 10	2.1	19
229	A novel contiguous gene deletion of <i>AVPR2</i> and <i>ARHGAP4</i> genes in male dizygotic twins with nephrogenic diabetes insipidus and intellectual disability. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 2511-2518.	0.7	19
230	ARX homeodomain mutations abolish DNA binding and lead to a loss of transcriptional repression. <i>Human Molecular Genetics</i> , 2012, 21, 1639-1647.	1.4	19
231	Human disease genes website series: An international, open and dynamic library for up-to-date clinical information. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1039-1046.	0.7	19
232	Construction of a YAC contig spanning the Xq13.3 subband. <i>Genomics</i> , 1995, 26, 115-122.	1.3	18
233	Brain cysts associated with mutation in the Aristaless related homeobox gene, ARX. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2003, 74, 536-538.	0.9	18
234	Refined mapping of X-linked reticulate pigmentary disorder and sequencing of candidate genes. <i>Human Genetics</i> , 2008, 123, 469-476.	1.8	18

#	ARTICLE	IF	CITATIONS
235	Histone demethylase KDM5C is a SAHA-sensitive central hub at the crossroads of transcriptional axes involved in multiple neurodevelopmental disorders. <i>Human Molecular Genetics</i> , 2019, 28, 4089-4102.	1.4	18
236	Disrupted Excitatory Synaptic Contacts and Altered Neuronal Network Activity Underpins the Neurological Phenotype in PCDH19-Clustering Epilepsy (PCDH19-CE). <i>Molecular Neurobiology</i> , 2021, 58, 2005-2018.	1.9	18
237	Polyalanine Tract Disorders and Neurocognitive Phenotypes. <i>Advances in Experimental Medicine and Biology</i> , 2012, 769, 185-203.	0.8	18
238	Heterozygous loss of function of <i>IQSEC2</i> leads to increased activated Arf6 and severe neurocognitive seizure phenotype in females. <i>Life Science Alliance</i> , 2019, 2, e201900386.	1.3	18
239	Physical and transcriptional mapping of DXS56-PGK1 1 Mb region: identification of three new transcripts. <i>Human Molecular Genetics</i> , 1993, 2, 1389-1396.	1.4	17
240	Clinical study of two brothers with a novel 33 bp duplication in the <i>ARX</i> gene. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 1482-1486.	0.7	17
241	Inherited balanced translocation t(9;17)(q33.2;q25.3) concomitant with a 16p13.1 duplication in a patient with schizophrenia. , 2011, 156, 204-214.		17
242	Identical by descent L1CAM mutation in two apparently unrelated families with intellectual disability without L1 syndrome. <i>European Journal of Medical Genetics</i> , 2015, 58, 364-368.	0.7	17
243	A mutation in <i>COL4A2</i> causes autosomal dominant porencephaly with cataracts. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1059-1063.	0.7	17
244	A mouse model for intellectual disability caused by mutations in the X-linked 2â€²â€² methyltransferase <i>Ftsj1</i> gene. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2019, 1865, 2083-2093.	1.8	17
245	Missense variant contribution to USP9X-female syndrome. <i>Npj Genomic Medicine</i> , 2020, 5, 53.	1.7	17
246	FMR3 is a novel gene associated with FRAXE CpG island and transcriptionally silent in FRAXE full mutations. <i>Journal of Medical Genetics</i> , 2000, 37, 782-784.	1.5	16
247	Cerebral palsy and genomics: an international consortium. <i>Developmental Medicine and Child Neurology</i> , 2018, 60, 209-210.	1.1	16
248	Levetiracetam efficacy in PCDH19 Girls Clustering Epilepsy. <i>European Journal of Paediatric Neurology</i> , 2020, 24, 142-147.	0.7	16
249	Definition and diagnosis of cerebral palsy in genetic studies: a systematic review. <i>Developmental Medicine and Child Neurology</i> , 2020, 62, 1024-1030.	1.1	16
250	Yield of clinically reportable genetic variants in unselected cerebral palsy by whole genome sequencing. <i>Npj Genomic Medicine</i> , 2021, 6, 74.	1.7	16
251	Restoring reproductive confidence in families with Xâ€²linked mental retardation by finding the causal mutation. <i>Clinical Genetics</i> , 2008, 73, 188-190.	1.0	15
252	BDNF and DYRK1A Are Variable and Inversely Correlated in Lymphoblastoid Cell Lines from Down Syndrome Patients. <i>Molecular Neurobiology</i> , 2012, 46, 297-303.	1.9	15

#	ARTICLE	IF	CITATIONS
253	NKX2-1 mutation in a family diagnosed with ataxic dyskinetic cerebral palsy. <i>European Journal of Medical Genetics</i> , 2013, 56, 506-509.	0.7	15
254	PCDH19 Pathogenic Variants in Males: Expanding the Phenotypic Spectrum. <i>Advances in Experimental Medicine and Biology</i> , 2020, 1298, 177-187.	0.8	15
255	Copy number variants in patients with intellectual disability affect the regulation of ARX transcription factor gene. <i>Human Genetics</i> , 2015, 134, 1163-1182.	1.8	14
256	Dysregulations of sonic hedgehog signaling in MED12-related X-linked intellectual disability disorders. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e00569.	0.6	14
257	EXOME REPORT: Novel mutation in ATP6V1B2 segregating with autosomal dominant epilepsy, intellectual disability and mild gingival and nail abnormalities. <i>European Journal of Medical Genetics</i> , 2020, 63, 103799.	0.7	14
258	Evidence for a Dual-Pathway, 2-Hit Genetic Model for Focal Cortical Dysplasia and Epilepsy. <i>Neurology: Genetics</i> , 2022, 8, e652.	0.9	14
259	Two unrelated patients with inversions of the X chromosome and non-specific mental retardation: physical and transcriptional mapping of their common breakpoint region in Xq13.1. <i>Journal of Medical Genetics</i> , 1999, 36, 754-758.	1.5	13
260	Borjeson-Forssman-Lehmann Syndrome and Multiple Pituitary Hormone Deficiency. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2003, 16, 1295-300.	0.4	13
261	A novel de novo 27â€‰%bp duplication of the <i>ARX</i> gene, resulting from postzygotic mosaicism and leading to three severely affected males in two generations. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 1655-1660.	0.7	13
262	Association of <i>SLC32A1</i> Missense Variants With Genetic Epilepsy With Febrile Seizures Plus. <i>Neurology</i> , 2021, 96, e2251-e2260.	1.5	13
263	X-linked intellectual disability: Phenotypic expression in carrier females. <i>Clinical Genetics</i> , 2020, 97, 418-425.	1.0	12
264	Expanding Clinical Presentations Due to Variations in THOC2 mRNA Nuclear Export Factor. <i>Frontiers in Molecular Neuroscience</i> , 2020, 13, 12.	1.4	12
265	Regulating transcriptional activity by phosphorylation: A new mechanism for the ARX homeodomain transcription factor. <i>PLoS ONE</i> , 2018, 13, e0206914.	1.1	11
266	Bi-allelic variants in SPATA5L1 lead to intellectual disability, spastic-dystonic cerebral palsy, epilepsy, and hearing loss. <i>American Journal of Human Genetics</i> , 2021, 108, 2006-2016.	2.6	11
267	PCR amplification of large VNTR alleles of D17S5 (YNZ22) locus. <i>Nucleic Acids Research</i> , 1991, 19, 5806-5806.	6.5	10
268	Loss of FMR2 further emphasizes the link between deregulation of immediate early response genes FOS and JUN and intellectual disability. <i>Human Molecular Genetics</i> , 2013, 22, 2984-2991.	1.4	10
269	A non-coding variant in the 5' UTR of DLG3 attenuates protein translation to cause non-syndromic intellectual disability. <i>European Journal of Human Genetics</i> , 2016, 24, 1612-1616.	1.4	10
270	Familial epilepsy with anterior polymicrogyria as a presentation of COL18A1 mutations. <i>European Journal of Medical Genetics</i> , 2017, 60, 437-443.	0.7	10

#	ARTICLE	IF	CITATIONS
271	Variant in the X-chromosome spliceosomal gene GPKOW causes male-lethal microcephaly with intrauterine growth restriction. <i>European Journal of Human Genetics</i> , 2017, 25, 1078-1082.	1.4	10
272	Whole genome sequencing in transposition of the great arteries and associations with clinically relevant heart, brain and laterality genes. <i>American Heart Journal</i> , 2022, 244, 1-13.	1.2	10
273	A novel locus for X-linked congenital cataract on Xq24. <i>Molecular Vision</i> , 2008, 14, 721-6.	1.1	10
274	Is there a Mendelian transmission ratio distortion of the c.429_452dup(24bp) polyalanine tract ARX mutation?. <i>European Journal of Human Genetics</i> , 2012, 20, 1311-1314.	1.4	9
275	Embryonic forebrain transcriptome of mice with polyalanine expansion mutations in the <i>ARX</i> homeobox gene. <i>Human Molecular Genetics</i> , 2016, 25, ddw360.	1.4	9
276	A synonymous <i>UPF3B</i> variant causing a speech disorder implicates NMD as a regulator of neurodevelopmental disorder gene networks. <i>Human Molecular Genetics</i> , 2020, 29, 2568-2578.	1.4	9
277	Characterization of ARHGGEF6, a guanine nucleotide exchange factor for Rho GTPases and a candidate gene for X-linked mental retardation: Mutation screening in <i>Brj1/2</i> rjeson-Forsman-Lehmann syndrome and MRX27. <i>American Journal of Medical Genetics Part A</i> , 2001, 100, 43-48.	2.4	8
278	Partial Androgen Insensitivity Syndrome and t(X;5): Are There Upstream Regulatory Elements of the Androgen Receptor Gene?. <i>Hormone Research in Paediatrics</i> , 2004, 62, 208-214.	0.8	8
279	Screening of 20 patients with X-linked mental retardation using chromosome X-specific array-MAPH. <i>European Journal of Medical Genetics</i> , 2007, 50, 399-410.	0.7	8
280	Genetics of the epilepsies: Genetic twists in the channels and other tales. <i>Epilepsia</i> , 2010, 51, 33-36.	2.6	8
281	Oligonucleotide correction of an intronic TIMMDC1 variant in cells of patients with severe neurodegenerative disorder. <i>Npj Genomic Medicine</i> , 2022, 7, 9.	1.7	8
282	Bi-allelic variants in neuronal cell adhesion molecule cause a neurodevelopmental disorder characterized by developmental delay, hypotonia, neuropathy/spasticity. <i>American Journal of Human Genetics</i> , 2022, 109, 518-532.	2.6	8
283	The DUB Club: Deubiquitinating Enzymes and Neurodevelopmental Disorders. <i>Biological Psychiatry</i> , 2022, 92, 614-625.	0.7	8
284	Fine mapping and cloning of the breakpoint associated with menkes syndrome in a female patient. <i>Genomics</i> , 1992, 14, 557-561.	1.3	7
285	Is FGF13 a major contributor to genetic epilepsy with febrile seizures plus?. <i>Epilepsy Research</i> , 2016, 128, 48-51.	0.8	7
286	A genomic cause of cerebral palsy should not change the clinical classification. <i>Annals of Clinical and Translational Neurology</i> , 2018, 5, 1011-1011.	1.7	7
287	Lack of FMR3 expression in a male with non-syndromic mental retardation and a microdeletion immediately distal to FRAXE CCG repeat. <i>Neuroscience Letters</i> , 2006, 397, 245-248.	1.0	6
288	RLIM Is a Candidate Dosage-Sensitive Gene for Individuals with Varying Duplications of Xq13, Intellectual Disability, and Distinct Facial Features. <i>American Journal of Human Genetics</i> , 2020, 107, 1157-1169.	2.6	6

#	ARTICLE	IF	CITATIONS
289	Chromatin-Binding Protein PHF6 Regulates Activity-Dependent Transcriptional Networks to Promote Hunger Response. <i>Cell Reports</i> , 2020, 30, 3717-3728.e6.	2.9	6
290	X-Linked Lissencephaly With Absent Corpus Callosum and Abnormal Genitalia. <i>Child Neurology Open</i> , 2017, 4, 2329048X1773862.	0.5	6
291	SLITRK2 variants associated with neurodevelopmental disorders impair excitatory synaptic function and cognition in mice. <i>Nature Communications</i> , 2022, 13, .	5.8	6
292	TM4SF10 gene sequencing in XLMR patients identifies common polymorphisms but no disease-associated mutation. <i>BMC Medical Genetics</i> , 2004, 5, 22.	2.1	5
293	Exclusion of biglycan mutations in a cohort of patients with neuromuscular disorders. <i>Neuromuscular Disorders</i> , 2008, 18, 606-609.	0.3	5
294	Recurrent deletion of <i>ZNF630</i> at Xp11.23 is not associated with mental retardation. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 638-645.	0.7	5
295	PCR Amplification and Sequence Analysis of GC-Rich Sequences: Aristaless-Related Homeobox Example. <i>Methods in Molecular Biology</i> , 2013, 1017, 105-120.	0.4	5
296	Transgenic mice with an R342X mutation in <i>Phf6</i> display clinical features of BÅrjesson-Forssman-Lehmann Syndrome. <i>Human Molecular Genetics</i> , 2021, 30, 575-594.	1.4	5
297	Protocadherin 19 Clustering Epilepsy and Neurosteroids: Opportunities for Intervention. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9769.	1.8	5
298	Identification of a <i>SEDL</i> gene mutation in an individual with Leber hereditary optic neuropathy and spondyloepiphyseal dysplasia. , 2004, 129A, 206-207.		4
299	Two novel intragenic variants in the <i>FMR1</i> gene in patients with suspect clinical diagnosis of Fragile X syndrome and no CGG repeat expansion. <i>European Journal of Medical Genetics</i> , 2020, 63, 104010.	0.7	4
300	Cerebral palsy with autism and ADHD: time to pay attention. <i>Developmental Medicine and Child Neurology</i> , 2021, 63, 247-248.	1.1	4
301	A 127â€kb truncating deletion of <i>PGRMC1</i> is a novel cause of X-linked isolated paediatric cataract. <i>European Journal of Human Genetics</i> , 2021, 29, 1206-1215.	1.4	4
302	Downregulation of the GHRH/GH/IGF-1 axis in a mouse model of BÅrjesson-Forssman-Lehman Syndrome. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	4
303	Common data elements to standardize genomics studies in cerebral palsy. <i>Developmental Medicine and Child Neurology</i> , 2022, 64, 1470-1476.	1.1	4
304	A novel syndrome of paediatric cataract, dysmorphism, ectodermal features, and developmental delay in Australian Aboriginal family maps to 1p35.3-p36.32. <i>BMC Medical Genetics</i> , 2010, 11, 165.	2.1	3
305	Developmental disorders: deciphering exomes on a grand scale. <i>Lancet, The</i> , 2015, 385, 1266-1267.	6.3	3
306	Further delineation of <i>BCAP31</i> -linked intellectual disability: description of 17 new families with LoF and missense variants. <i>European Journal of Human Genetics</i> , 2021, 29, 1405-1417.	1.4	3

#	ARTICLE	IF	CITATIONS
307	Simultaneous Screening of the FRAXA and FRAXE Loci for Rapid Detection of FMR1 CCG and/or AFF2 CCG Repeat Expansions by Triplet-Primed PCR. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 941-951.	1.2	3
308	UPF3B Gene and Nonsense-Mediated mRNA Decay in Autism Spectrum Disorders. , 2014, , 1663-1678.		3
309	Deletion of F508 and haplotype analysis of CFTR gene region in Slovak CF patients. <i>Human Genetics</i> , 1992, 89, 305-306.	1.8	2
310	Molecular genetics of X-linked mental retardation: a complex picture emerging. <i>Expert Review of Molecular Diagnostics</i> , 2001, 1, 220-225.	1.5	2
311	A novel genetic syndrome characterized by pediatric cataract, dysmorphism, ectodermal features, and developmental delay in an indigenous Australian family. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 633-639.	0.7	2
312	Blinders, phenotype, and fashionable genetic analysis Setting the record straight for epilepsy!. <i>Epilepsia</i> , 2011, 52, 1757-1758.	2.6	2
313	Avascular necrosis of bone in childhood cancer patients: a possible role of genetic susceptibility. <i>Bratislava Medical Journal</i> , 2015, 116, 289-295.	0.4	2
314	Robust imaging and gene delivery to study human lymphoblastoid cell lines. <i>Journal of Human Genetics</i> , 2018, 63, 945-955.	1.1	2
315	Constraint and conservation of paired-type homeodomains predicts the clinical outcome of missense variants of uncertain significance. <i>Human Mutation</i> , 2020, 41, 1407-1424.	1.1	2
316	TaqI Digestion of PCR Product Increases the Informativity of St14 Vntr for the Diagnosis of Hemophilia A. <i>Disease Markers</i> , 1993, 11, 139-141.	0.6	1
317	Great expectations: using massively parallel sequencing to solve inherited disorders. <i>Expert Review of Molecular Diagnostics</i> , 2010, 10, 833-836.	1.5	1
318	New mutations and sporadic intellectual disability. <i>Lancet, The</i> , 2012, 380, 1630-1631.	6.3	1
319	Challenges of Sticky-Co-immunoprecipitation: Polyalanine Tract Protein-Protein Interactions. <i>Methods in Molecular Biology</i> , 2013, 1017, 121-133.	0.4	1
320	Protocadherin Mutations in Neurodevelopmental Disorders. , 2016, , 221-231.		1
321	Integrated in silico and experimental assessment of disease relevance of PCDH19 missense variants. <i>Human Mutation</i> , 2021, 42, 1030-1041.	1.1	1
322	People with Cerebral Palsy and Their Family's Preferences about Genomics Research. <i>Public Health Genomics</i> , 2022, 25, 22-31.	0.6	1
323	NMD-deficient Upf3b null mice display behavioral and neuropathological defects. <i>FASEB Journal</i> , 2012, 26, 747.5.	0.2	1
324	Distribution of ApoBII, MCT118 (D1S80), YNZ22 (D17S30), and COL2A1 Amp-FLPs (amplified fragment length) in Human Gene Frequencies, 1994, 8, 121-7.	0.1	1

#	ARTICLE	IF	CITATIONS
325	Spectrum of Neurological Phenotypes Caused by ARX Mutations. Acta Neurologica Scandinavica, 2003, 107, 432-432.	1.0	0
326	Phenotypeâ€“genotype complexities: opening DOORS. Lancet Neurology, The, 2014, 13, 24-25.	4.9	0
327	Cerebral Palsy: Causes, Pathways, and the Role of Genetic Variants. Obstetric Anesthesia Digest, 2016, 36, 185-185.	0.0	0
328	Reply. American Journal of Obstetrics and Gynecology, 2016, 214, 671.	0.7	0
329	Non-Syndromic 46,XY Disorders of Sex Development. Acta Medica Martiniana, 2018, 18, 35-41.	0.4	0
330	Different types of diseaseâ€“causing noncoding variants revealed by genomic and gene expression analyses in families with Xâ€“linked intellectual disability. Human Mutation, 2021, 42, 835-847.	1.1	0
331	Frequency of Cystic Fibrosis Mutations and Associated Haplotype Distribution in Slovak CF Patients. Advances in Experimental Medicine and Biology, 1991, 290, 383-385.	0.8	0
332	Direct cDNA Selection Using Human and Mouse cDNAs: Application to Xq13.3 Chromosomal Region. , 1994, , 81-90.		0
333	Developmental Abnormalities Due to Mutations in the Aristaless-Related Homeobox Gene. , 2016, , 761-766.		0
334	Frequency and distribution of deletions in dystrophin gene in Duchenne muscular dystrophy patients from an east-European Slavonic population. Gene Geography: A Computerized Bulletin on Human Gene Frequencies, 1991, 5, 137-40.	0.1	0