

Maha Zaki

List of Publications by Year in descending order

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Version: 2024-02-01

160
papers

7,417
citations

66315

42
h-index

69214

77
g-index

165
all docs

165
docs citations

165
times ranked

12948
citing authors

#	ARTICLE	IF	CITATIONS
1	Biallelic variants in <i>SLC38A3</i> encoding a glutamine transporter cause epileptic encephalopathy. <i>Brain</i> , 2022, 145, 909-924.	3.7	17
2	Biallelic <i>FRA10AC1</i> variants cause a neurodevelopmental disorder with growth retardation. <i>Brain</i> , 2022, 145, 1551-1563.	3.7	9
3	Clinico-radiological features, molecular spectrum, and identification of prognostic factors in developmental and epileptic encephalopathy due to inosine triphosphate pyrophosphatase (ITPase) deficiency. <i>Human Mutation</i> , 2022, 43, 403-419.	1.1	9
4	Bi-allelic variants in <i>CHKA</i> cause a neurodevelopmental disorder with epilepsy and microcephaly. <i>Brain</i> , 2022, 145, 1916-1923.	3.7	3
5	<i>El-Hattab-Alkuraya</i> syndrome caused by biallelic <i>WDR45B</i> pathogenic variants: Further delineation of the phenotype and genotype. <i>Clinical Genetics</i> , 2022, 101, 530-540.	1.0	7
6	Variable predicted pathogenic mechanisms for novel <i>MECP2</i> variants in RTT patients. <i>Journal of Genetic Engineering and Biotechnology</i> , 2022, 20, 44.	1.5	0
7	Biallelic Variants in the Ectonucleotidase <i>ENTPD1</i> Cause a Complex Neurodevelopmental Disorder with Intellectual Disability, Distinct White Matter Abnormalities, and Spastic Paraplegia. <i>Annals of Neurology</i> , 2022, 92, 304-321.	2.8	2
8	Monoallelic and biallelic mutations in <i>RELN</i> underlie a graded series of neurodevelopmental disorders. <i>Brain</i> , 2022, 145, 3274-3287.	3.7	6
9	Biallelic variants in <i>ZNF142</i> lead to a syndromic neurodevelopmental disorder. <i>Clinical Genetics</i> , 2022, 102, 98-109.	1.0	6
10	Inhibition of G-protein signalling in cardiac dysfunction of intellectual developmental disorder with cardiac arrhythmia (IDCCA) syndrome. <i>Journal of Medical Genetics</i> , 2021, 58, 815-831.	1.5	3
11	Biallelic variants in <i>ADARB1</i> , encoding a dsRNA-specific adenosine deaminase, cause a severe developmental and epileptic encephalopathy. <i>Journal of Medical Genetics</i> , 2021, 58, 495-504.	1.5	14
12	Biallelic variants in <i>HPDL</i> , encoding 4-hydroxyphenylpyruvate dioxygenase-like protein, lead to an infantile neurodegenerative condition. <i>Genetics in Medicine</i> , 2021, 23, 524-533.	1.1	17
13	Mutations in Spliceosomal Genes <i>PPIL1</i> and <i>PRP17</i> Cause Neurodegenerative Pontocerebellar Hypoplasia with Microcephaly. <i>Neuron</i> , 2021, 109, 241-256.e9.	3.8	31
14	Clinical, Biochemical, and Molecular Characterization of Metachromatic Leukodystrophy Among Egyptian Pediatric Patients: Expansion of the <i>ARSA</i> Mutational Spectrum. <i>Journal of Molecular Neuroscience</i> , 2021, 71, 1112-1130.	1.1	1
15	<i>UBR7</i> functions with <i>UBR5</i> in the Notch signaling pathway and is involved in a neurodevelopmental syndrome with epilepsy, ptosis, and hypothyroidism. <i>American Journal of Human Genetics</i> , 2021, 108, 134-147.	2.6	15
16	Clinical and genetic characterization of ten Egyptian patients with Wolf-Hirschhorn syndrome and review of literature. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1546.	0.6	6
17	Alternative genomic diagnoses for individuals with a clinical diagnosis of Dubowitz syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 119-133.	0.7	17
18	A relatively common homozygous <i>TRAPPC4</i> splicing variant is associated with an early-infantile neurodegenerative syndrome. <i>European Journal of Human Genetics</i> , 2021, 29, 271-279.	1.4	8

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19	Homozygous missense <i>WIP12</i> variants cause a congenital disorder of autophagy with neurodevelopmental impairments of variable clinical severity and disease course. <i>Brain Communications</i> , 2021, 3, fcab183.	1.5	10
20	Expanding the phenotype of <i>PIGS</i> associated early onset epileptic developmental encephalopathy. <i>Epilepsia</i> , 2021, 62, e35-e41.	2.6	11
21	Neurodevelopmental disorder in an Egyptian family with a biallelic <i>ALKBH8</i> variant. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1288-1293.	0.7	13
22	Novel <i>NDUFA12</i> variants are associated with isolated complex I defect and variable clinical manifestation. <i>Human Mutation</i> , 2021, 42, 699-710.	1.1	12
23	Biallelic hypomorphic mutations in <i>HEATR5B</i> , encoding HEAT repeat-containing protein 5B, in a neurological syndrome with pontocerebellar hypoplasia. <i>European Journal of Human Genetics</i> , 2021, 29, 957-964.	1.4	7
24	Combining exome/genome sequencing with data repository analysis reveals novel gene-disease associations for a wide range of genetic disorders. <i>Genetics in Medicine</i> , 2021, 23, 1551-1568.	1.1	30
25	Loss of function mutations in <i>GEMIN5</i> cause a neurodevelopmental disorder. <i>Nature Communications</i> , 2021, 12, 2558.	5.8	28
26	Pathogenic variants in <i>PIDD1</i> lead to an autosomal recessive neurodevelopmental disorder with pachygyria and psychiatric features. <i>European Journal of Human Genetics</i> , 2021, 29, 1226-1234.	1.4	8
27	Biallelic variants in <i>KARS1</i> are associated with neurodevelopmental disorders and hearing loss recapitulated by the knockout zebrafish. <i>Genetics in Medicine</i> , 2021, 23, 1933-1943.	1.1	11
28	Loss of <i>C2orf69</i> defines a fatal autoinflammatory syndrome in humans and zebrafish that evokes a glycogen-storage-associated mitochondriopathy. <i>American Journal of Human Genetics</i> , 2021, 108, 1301-1317.	2.6	11
29	Samia Temtamy. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3613-3614.	0.7	0
30	Biallelic loss-of-function variants in the splicing regulator <i>NSRP1</i> cause a severe neurodevelopmental disorder with spastic cerebral palsy and epilepsy. <i>Genetics in Medicine</i> , 2021, 23, 2455-2460.	1.1	9
31	Undiagnosed Phenylketonuria Can Exist Everywhere: Results From an International Survey. <i>Journal of Pediatrics</i> , 2021, 239, 231-234.e2.	0.9	9
32	Implication of folate deficiency in <i>CYP2U1</i> loss of function. <i>Journal of Experimental Medicine</i> , 2021, 218, .	4.2	13
33	<i>ABHD16A</i> deficiency causes a complicated form of hereditary spastic paraplegia associated with intellectual disability and cerebral anomalies. <i>American Journal of Human Genetics</i> , 2021, 108, 2017-2023.	2.6	9
34	Bi-allelic <i>TTC5</i> variants cause delayed developmental milestones and intellectual disability. <i>Journal of Medical Genetics</i> , 2021, 58, 237-246.	1.5	4
35	Chromosome 9p terminal deletion in nine Egyptian patients and narrowing of the critical region for trigonocephaly. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1829.	0.6	4
36	The potential impact of <i>COMT</i> gene variants on dopamine regulation and phenotypic traits of ASD patients. <i>Behavioural Brain Research</i> , 2020, 378, 112272.	1.2	15

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37	Mutation spectrum in the gene encoding methyl-CpG-binding protein 2 in Egyptian patients with Rett syndrome. <i>Meta Gene</i> , 2020, 24, 100620.	0.3	1
38	Defining the clinical, molecular and imaging spectrum of adaptor protein complex 4-associated hereditary spastic paraplegia. <i>Brain</i> , 2020, 143, 2929-2944.	3.7	29
39	Biallelic in-frame deletion in <i>TRAPPC4</i> in a family with developmental delay and cerebellar atrophy. <i>Brain</i> , 2020, 143, e83-e83.	3.7	8
40	MINPP1 prevents intracellular accumulation of the chelator inositol hexakisphosphate and is mutated in Pontocerebellar Hypoplasia. <i>Nature Communications</i> , 2020, 11, 6087.	5.8	28
41	Biallelic loss of function variants in <i>SYT2</i> cause a treatable congenital onset presynaptic myasthenic syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2272-2283.	0.7	20
42	Clinical, neuropathological, and genetic characterization of STUB1 variants in cerebellar ataxias: a frequent cause of predominant cognitive impairment. <i>Genetics in Medicine</i> , 2020, 22, 1851-1862.	1.1	30
43	Micro and Martsolf syndromes in 34 new patients: Refining the phenotypic spectrum and further molecular insights. <i>Clinical Genetics</i> , 2020, 98, 445-456.	1.0	12
44	<i>ASAHI</i> -related disorders: Description of 15 novel pediatric patients and expansion of the clinical phenotype. <i>Clinical Genetics</i> , 2020, 98, 598-605.	1.0	3
45	Blepharophimosis-epitosis-intellectual disability syndrome: A report of nine Egyptian patients with further expansion of phenotypic and mutational spectrum. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2857-2866.	0.7	4
46	International consensus recommendations on the diagnostic work-up for malformations of cortical development. <i>Nature Reviews Neurology</i> , 2020, 16, 618-635.	4.9	53
47	Prenatal delineation of a distinct lethal fetal syndrome caused by a homozygous truncating <i>KIDINS220</i> variant. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2867-2876.	0.7	10
48	A founder mutation in PEX12 among Egyptian patients in peroxisomal biogenesis disorder. <i>Neurological Sciences</i> , 2020, 42, 2737-2745.	0.9	1
49	Molecular diagnosis in recessive pediatric neurogenetic disease can help reduce disease recurrence in families. <i>BMC Medical Genomics</i> , 2020, 13, 68.	0.7	4
50	Recurrent homozygous damaging mutation in <i>TMX2</i> , encoding a protein disulfide isomerase, in four families with microlissencephaly. <i>Journal of Medical Genetics</i> , 2020, 57, 274-282.	1.5	6
51	Loss of PYCR2 Causes Neurodegeneration by Increasing Cerebral Glycine Levels via SHMT2. <i>Neuron</i> , 2020, 107, 82-94.e6.	3.8	30
52	Regulation of human cerebral cortical development by EXOC7 and EXOC8, components of the exocyst complex, and roles in neural progenitor cell proliferation and survival. <i>Genetics in Medicine</i> , 2020, 22, 1040-1050.	1.1	13
53	Microcephalic osteodysplastic primordial dwarfism type II: Additional nine patients with implications on phenotype and genotype correlation. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1407-1420.	0.7	11
54	Bi-allelic Variants in the GPI Transamidase Subunit PIGK Cause a Neurodevelopmental Syndrome with Hypotonia, Cerebellar Atrophy, and Epilepsy. <i>American Journal of Human Genetics</i> , 2020, 106, 484-495.	2.6	22

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55	RSRC1 loss-of-function variants cause mild to moderate autosomal recessive intellectual disability. <i>Brain</i> , 2020, 143, e31-e31.	3.7	6
56	Loss of the neural-specific BAF subunit ACTL6B relieves repression of early response genes and causes recessive autism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 10055-10066.	3.3	34
57	Novel congenital disorder of <i>O</i> -linked glycosylation caused by GALNT2 loss of function. <i>Brain</i> , 2020, 143, 1114-1126.	3.7	46
58	Genetic pattern of SMN1, SMN2, and NAIP genes in prognosis of SMA patients. <i>Egyptian Journal of Medical Human Genetics</i> , 2020, 21, .	0.5	10
59	Variants in the NPC1 Gene in Egyptian Patients with Niemann-Pick Type C. <i>Open Access Macedonian Journal of Medical Sciences</i> , 2020, 8, 134-145.	0.1	0
60	<i>ASAH1</i> pathogenic variants associated with acid ceramidase deficiency: Farber disease and spinal muscular atrophy with progressive myoclonic epilepsy. <i>Human Mutation</i> , 2020, 41, 1469-1487.	1.1	8
61	Bi-allelic GOT2 Mutations Cause a Treatable Malate-Aspartate Shuttle-Related Encephalopathy. <i>American Journal of Human Genetics</i> , 2019, 105, 534-548.	2.6	46
62	Phenotypic and mutational spectrum of thirty-five patients with Sjögren-Larsson syndrome: identification of eleven novel ALDH3A2 mutations and founder effects. <i>Journal of Human Genetics</i> , 2019, 64, 859-865.	1.1	8
63	Loss of SMPD4 Causes a Developmental Disorder Characterized by Microcephaly and Congenital Arthrogyriposis. <i>American Journal of Human Genetics</i> , 2019, 105, 689-705.	2.6	48
64	Bi-allelic Loss of Human APC2, Encoding Adenomatous Polyposis Coli Protein 2, Leads to Lissencephaly, Subcortical Heterotopia, and Global Developmental Delay. <i>American Journal of Human Genetics</i> , 2019, 105, 844-853.	2.6	17
65	Study of C677T variant of methylene tetrahydrofolate reductase gene in autistic spectrum disorder Egyptian children. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 305-309.	1.1	12
66	Recessive mutations in muscle-specific isoforms of FXR1 cause congenital multi-minicore myopathy. <i>Nature Communications</i> , 2019, 10, 797.	5.8	24
67	Biallelic mutations in valyl-tRNA synthetase gene VARS are associated with a progressive neurodevelopmental epileptic encephalopathy. <i>Nature Communications</i> , 2019, 10, 707.	5.8	28
68	Bilateral Calcification of Basal Ganglia in a Patient with Duplication of Both 11q13.1q22.1 and 4q35.2 with New Phenotypic Features. <i>Cytogenetic and Genome Research</i> , 2019, 159, 130-136.	0.6	3
69	Genomic and phenotypic delineation of congenital microcephaly. <i>Genetics in Medicine</i> , 2019, 21, 545-552.	1.1	85
70	GAP0 syndrome in seven new patients: Identification of five novel <i>ANTXR1</i> mutations including the first large intragenic deletion. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 237-242.	0.7	8
71	Clinical, biomarker and genetic spectrum of Niemann-Pick type C in Egypt: The detection of nine novel <i>NPC1</i> mutations. <i>Clinical Genetics</i> , 2019, 95, 537-539.	1.0	4
72	Loss of the sphingolipid desaturase DEGS1 causes hypomyelinating leukodystrophy. <i>Journal of Clinical Investigation</i> , 2019, 129, 1240-1256.	3.9	68

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73	Analysis of 17 genes detects mutations in 81% of 811 patients with lissencephaly. <i>Genetics in Medicine</i> , 2018, 20, 1354-1364.	1.1	92
74	Identification of a novel homozygous ALX4 mutation in two unrelated patients with frontonasal dysplasia type 2. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1190-1194.	0.7	7
75	Expanding the phenome and variome of skeletal dysplasia. <i>Genetics in Medicine</i> , 2018, 20, 1609-1616.	1.1	46
76	Biallelic variants in KIF14 cause intellectual disability with microcephaly. <i>European Journal of Human Genetics</i> , 2018, 26, 330-339.	1.4	52
77	<i>PGAP3</i> -related hyperphosphatasia with mental retardation syndrome: Report of 10 new patients and a homozygous founder mutation. <i>Clinical Genetics</i> , 2018, 93, 84-91.	1.0	20
78	A homozygous founder mutation in <i>TRAPPC6B</i> associates with a neurodevelopmental disorder characterised by microcephaly, epilepsy and autistic features. <i>Journal of Medical Genetics</i> , 2018, 55, 48-54.	1.5	37
79	Hypermanganesemia with dystonia, polycythemia and cirrhosis in 10 patients: Six novel <i>SLC30A10</i> mutations and further phenotype delineation. <i>Clinical Genetics</i> , 2018, 93, 905-912.	1.0	30
80	Loss of tubulin deglutamylase <i>CCP1</i> causes infantile-onset neurodegeneration. <i>EMBO Journal</i> , 2018, 37, .	3.5	86
81	Unbalanced 14;X Translocation and Pattern of X Inactivation in a Female Patient with Multiple Congenital Anomalies. <i>Cytogenetic and Genome Research</i> , 2018, 156, 71-79.	0.6	1
82	Loss of <i>Protocadherin 12</i> leads to disjunction of the <i>D</i> domain of <i>PCDH12</i> in a patient with a form of <i>D</i> domain dysplasia syndrome. <i>Annals of Neurology</i> , 2018, 84, 638-647.	2.8	19
83	Genetic variants in components of the <i>NALCN</i> - <i>UNC80</i> - <i>UNC79</i> ion channel complex cause a broad clinical phenotype (<i>NALCN</i> channelopathies). <i>Human Genetics</i> , 2018, 137, 753-768.	1.8	38
84	Biallelic loss of human <i>CTNNA2</i> , encoding β -catenin, leads to <i>ARP2/3</i> complex overactivity and disordered cortical neuronal migration. <i>Nature Genetics</i> , 2018, 50, 1093-1101.	9.4	70
85	Mutations in <i>LNPK</i> , Encoding the Endoplasmic Reticulum Junction Stabilizer Lunapark, Cause a Recessive Neurodevelopmental Syndrome. <i>American Journal of Human Genetics</i> , 2018, 103, 296-304.	2.6	24
86	Biallelic mutations in the 3' exonuclease <i>TOE1</i> cause pontocerebellar hypoplasia and uncover a role in snRNA processing. <i>Nature Genetics</i> , 2017, 49, 457-464.	9.4	66
87	A novel frameshift mutation in the sterol 27-hydroxylase gene in an Egyptian family with cerebrotendinous xanthomatosis without cataract. <i>Metabolic Brain Disease</i> , 2017, 32, 311-315.	1.4	3
88	Band-like calcification with simplified gyration and polymicrogyria: report of 10 new families and identification of five novel <i>OCLN</i> mutations. <i>Journal of Human Genetics</i> , 2017, 62, 553-559.	1.1	18
89	Aicardi-Goutières syndrome: unusual neuro-radiological manifestations. <i>Metabolic Brain Disease</i> , 2017, 32, 679-683.	1.4	10
90	Biallelic Variants in <i>OTUD6B</i> Cause an Intellectual Disability Syndrome Associated with Seizures and Dysmorphic Features. <i>American Journal of Human Genetics</i> , 2017, 100, 676-688.	2.6	54

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91	Mutations in GPAA1 , Encoding a GPI Transamidase Complex Protein, Cause Developmental Delay, Epilepsy, Cerebellar Atrophy, and Osteopenia. American Journal of Human Genetics, 2017, 101, 856-865.	2.6	49
92	Hypomorphic Recessive Variants in SUFU Impair the Sonic Hedgehog Pathway and Cause Joubert Syndrome with Cranio-facial and Skeletal Defects. American Journal of Human Genetics, 2017, 101, 552-563.	2.6	45
93	Homozygous Mutations in TBC1D23 Lead to a Non-degenerative Form of Pontocerebellar Hypoplasia. American Journal of Human Genetics, 2017, 101, 441-450.	2.6	43
94	Extracellular miR-145, miR-223 and miR-326 expression signature allow for differential diagnosis of immune-mediated neuroinflammatory diseases. Journal of the Neurological Sciences, 2017, 383, 188-198.	0.3	36
95	Recurrent and Prolonged Infections in a Child with a Homozygous IFIH1 Nonsense Mutation. Frontiers in Genetics, 2017, 8, 130.	1.1	35
96	Megalencephalic leukoencephalopathy with cysts in twelve Egyptian patients: novel mutations in MLC1 and HEPACAM and a founder effect. Metabolic Brain Disease, 2016, 31, 1171-1179.	1.4	7
97	Molecular and phenotypic spectrum of <i>ASPM</i>-related primary microcephaly: Identification of eight novel mutations. American Journal of Medical Genetics, Part A, 2016, 170, 2133-2140.	0.7	30
98	PEX6 is Expressed in Photoreceptor Cilia and Mutated in Deafblindness with Enamel Dysplasia and Microcephaly. Human Mutation, 2016, 37, 170-174.	1.1	36
99	Extending the mutation spectrum for Gallowayâ€™Mowat syndrome to include homozygous missense mutations in the WDR73 gene. American Journal of Medical Genetics, Part A, 2016, 170, 992-998.	0.7	26
100	<i>PYCR2</i> Mutations cause a lethal syndrome of microcephaly and failure to thrive. Annals of Neurology, 2016, 80, 59-70.	2.8	35
101	Genotype/phenotype correlation in a female patient with 21q22.3 and 12p13.33 duplications. American Journal of Medical Genetics, Part A, 2016, 170, 1050-1058.	0.7	6
102	Mutations in MBOAT7 , Encoding Lysophosphatidylinositol Acyltransferase I, Lead to Intellectual Disability Accompanied by Epilepsy and Autistic Features. American Journal of Human Genetics, 2016, 99, 912-916.	2.6	69
103	Biallelic Mutations in Citron Kinase Link Mitotic Cytokinesis to Human Primary Microcephaly. American Journal of Human Genetics, 2016, 99, 501-510.	2.6	70
104	Biallelic Mutations in TMTC3, Encoding a Transmembrane and TPR-Containing Protein, Lead to Cobblestone Lissencephaly. American Journal of Human Genetics, 2016, 99, 1181-1189.	2.6	30
105	Exome sequencing discloses KALRN homozygous variant as likely cause of intellectual disability and short stature in a consanguineous pedigree. Human Genomics, 2016, 10, 26.	1.4	13
106	Molybdenum cofactor and isolated sulphite oxidase deficiencies: Clinical and molecular spectrum among Egyptian patients. European Journal of Paediatric Neurology, 2016, 20, 714-722.	0.7	33
107	Pathogenic Variants in PIGG Cause Intellectual Disability with Seizures and Hypotonia. American Journal of Human Genetics, 2016, 98, 615-626.	2.6	71
108	Mutations in UNC80, Encoding Part of the UNC79-UNC80-NALCN Channel Complex, Cause Autosomal-Recessive Severe Infantile Encephalopathy. American Journal of Human Genetics, 2016, 98, 210-215.	2.6	37

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109	Dandyâ€“Walker malformation, genitourinary abnormalities, and intellectual disability in two families. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2503-2507.	0.7	6
110	Functional genome-wide siRNA screen identifies KIAA0586 as mutated in Joubert syndrome. <i>ELife</i> , 2015, 4, e06602.	2.8	64
111	Inactivating mutations in MFSD2A, required for omega-3 fatty acid transport in brain, cause a lethal microcephaly syndrome. <i>Nature Genetics</i> , 2015, 47, 809-813.	9.4	180
112	Non-manifesting AHI1 truncations indicate localized loss-of-function tolerance in a severe Mendelian disease gene. <i>Human Molecular Genetics</i> , 2015, 24, 2594-2603.	1.4	32
113	Characterization of human disease phenotypes associated with mutations in <i>TREX1</i> , <i>RNASEH2A</i> , <i>RNASEH2B</i> , <i>RNASEH2C</i> , <i>SAMHD1</i> , <i>ADAR</i> , and <i>IFIH1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 296-312.	0.7	447
114	Biallelic mutations in SNX14 cause a syndromic form of cerebellar atrophy and lysosome-autophagosome dysfunction. <i>Nature Genetics</i> , 2015, 47, 528-534.	9.4	111
115	Overexpression of <i>KLC2</i> due to a homozygous deletion in the non-coding region causes SPOAN syndrome. <i>Human Molecular Genetics</i> , 2015, 24, ddv388.	1.4	34
116	Autosomal dominant SCA5 and autosomal recessive infantile SCA are allelic conditions resulting from SPTBN2 mutations. <i>European Journal of Human Genetics</i> , 2014, 22, 286-288.	1.4	37
117	Mutations in ADAR1, IFIH1, and RNASEH2B Presenting As Spastic Paraplegia. <i>Neuropediatrics</i> , 2014, 45, 386-391.	0.3	72
118	Mutations in KATNB1 Cause Complex Cerebral Malformations by Disrupting Asymmetrically Dividing Neural Progenitors. <i>Neuron</i> , 2014, 84, 1226-1239.	3.8	95
119	Biallelic Truncating Mutations in FMN2, Encoding the Actin-Regulatory Protein Formin 2, Cause Nonsyndromic Autosomal-Recessive Intellectual Disability. <i>American Journal of Human Genetics</i> , 2014, 95, 721-728.	2.6	62
120	Mutations in CSPP1 Lead to Classical Joubert Syndrome. <i>American Journal of Human Genetics</i> , 2014, 94, 80-86.	2.6	75
121	Exome Sequencing Links Corticospinal Motor Neuron Disease to Common Neurodegenerative Disorders. <i>Science</i> , 2014, 343, 506-511.	6.0	466
122	Diagnostic Exome Sequencing to Elucidate the Genetic Basis of Likely Recessive Disorders in Consanguineous Families. <i>Human Mutation</i> , 2014, 35, 1203-1210.	1.1	75
123	Novel mutation in the fukutin gene in an Egyptian family with Fukuyama congenital muscular dystrophy and microcephaly. <i>Gene</i> , 2014, 539, 279-282.	1.0	5
124	AMPD2 Regulates GTP Synthesis and Is Mutated in a Potentially Treatable Neurodegenerative Brainstem Disorder. <i>Cell</i> , 2013, 154, 505-517.	13.5	94
125	Mutations in EOGT Confirm the Genetic Heterogeneity of Autosomal-Recessive Adams-Oliver Syndrome. <i>American Journal of Human Genetics</i> , 2013, 92, 598-604.	2.6	114
126	Mutations in LAMB1 Cause Cobblestone Brain Malformation without Muscular or Ocular Abnormalities. <i>American Journal of Human Genetics</i> , 2013, 92, 468-474.	2.6	96

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127	Whole-Exome Sequencing Identifies Mutated C12orf57 in Recessive Corpus Callosum Hypoplasia. American Journal of Human Genetics, 2013, 92, 392-400.	2.6	28
128	Assessment of interferon-related biomarkers in Aicardi-GoutiÃ¨res syndrome associated with mutations in TREX1, RNASEH2A, RNASEH2B, RNASEH2C, SAMHD1, and ADAR: a case-control study. Lancet Neurology, The, 2013, 12, 1159-1169.	4.9	473
129	Mutation Spectrum in <i>RAB3</i> , <i>GAP1</i> , <i>RAB3</i> , <i>GAP2</i> , and <i>RAB18</i> and Genotype-Phenotype Correlations in Warburg Micro Syndrome and Martsolf Syndrome. Human Mutation, 2013, 34, 686-696.	1.1	114
130	Mutations in ANTXR1 Cause GAPO Syndrome. American Journal of Human Genetics, 2013, 92, 792-799.	2.6	73
131	Clinical and molecular findings in eight Egyptian patients with suspected mitochondrial disorders and optic atrophy. Egyptian Journal of Medical Human Genetics, 2013, 14, 37-47.	0.5	2
132	Phenotypic spectrum and prevalence of INPP5E mutations in Joubert Syndrome and related disorders. European Journal of Human Genetics, 2013, 21, 1074-1078.	1.4	64
133	Further delineation of the clinical spectrum in <i>RNU4</i> related microcephalic osteodysplastic primordial dwarfism type I. American Journal of Medical Genetics, Part A, 2013, 161, 1875-1881.	0.7	20
134	Evolutionarily Assembled cis-Regulatory Module at a Human Ciliopathy Locus. Science, 2012, 335, 966-969.	6.0	84
135	Exome Sequencing Can Improve Diagnosis and Alter Patient Management. Science Translational Medicine, 2012, 4, 138ra78.	5.8	226
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