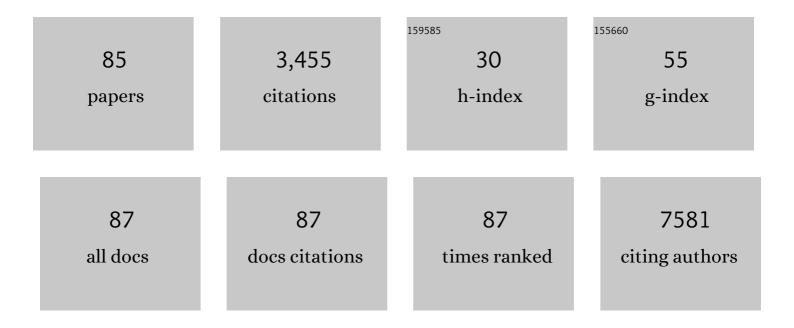
List of Publications by Year in descending order

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ΔΝΝΑΙΕΗΜΑΝ

#	Article	IF	CITATIONS
1	High Rate of Recurrent De Novo Mutations in Developmental and Epileptic Encephalopathies. American Journal of Human Genetics, 2017, 101, 664-685.	6.2	337
2	Exome Sequencing and the Management of Neurometabolic Disorders. New England Journal of Medicine, 2016, 374, 2246-2255.	27.0	254
3	Transcriptional Regulation of BACE1, the β-Amyloid Precursor Protein β-Secretase, by Sp1. Molecular and Cellular Biology, 2004, 24, 865-874.	2.3	207
4	Histone Lysine Methylases and Demethylases in the Landscape of Human Developmental Disorders. American Journal of Human Genetics, 2018, 102, 175-187.	6.2	204
5	Defects in the IFT-B Component IFT172 Cause Jeune and Mainzer-Saldino Syndromes in Humans. American Journal of Human Genetics, 2013, 93, 915-925.	6.2	196
6	Mutations in NOTCH1 Cause Adams-Oliver Syndrome. American Journal of Human Genetics, 2014, 95, 275-284.	6.2	150
7	Mutations in B4GALNT1 (GM2 synthase) underlie a new disorder of ganglioside biosynthesis. Brain, 2013, 136, 3618-3624.	7.6	115
8	Mutations of AKT3 are associated with a wide spectrum of developmental disorders including extreme megalencephaly. Brain, 2017, 140, 2610-2622.	7.6	102
9	Loss-of-Function and Gain-of-Function Mutations in KCNQ5 Cause Intellectual Disability or Epileptic Encephalopathy. American Journal of Human Genetics, 2017, 101, 65-74.	6.2	99
10	Mosaic Activating Mutations in FGFR1 Cause Encephalocraniocutaneous Lipomatosis. American Journal of Human Genetics, 2016, 98, 579-587.	6.2	88
11	Mitochondrial Carbonic Anhydrase VA Deficiency Resulting from CA5A Alterations Presents with Hyperammonemia in Early Childhood. American Journal of Human Genetics, 2014, 94, 453-461.	6.2	82
12	<i>OCRL1</i> Mutations in Dent 2 Patients Suggest a Mechanism for Phenotypic Variability. Nephron Physiology, 2009, 112, p27-p36.	1.2	79
13	The cost and diagnostic yield of exome sequencing for children with suspected genetic disorders: a benchmarking study. Genetics in Medicine, 2018, 20, 1-9.	2.4	79
14	Heterozygous Loss-of-Function Mutations in DLL4 Cause Adams-Oliver Syndrome. American Journal of Human Genetics, 2015, 97, 475-482.	6.2	73
15	Mutations in the Chromatin Regulator Gene BRPF1 Cause Syndromic Intellectual Disability and Deficient Histone Acetylation. American Journal of Human Genetics, 2017, 100, 91-104.	6.2	72
16	Mutations in the Spliceosome Component CWC27 Cause Retinal Degeneration with or without Additional Developmental Anomalies. American Journal of Human Genetics, 2017, 100, 592-604.	6.2	61
17	RAPIDOMICS: rapid genome-wide sequencing in a neonatal intensive care unit—successes and challenges. European Journal of Pediatrics, 2019, 178, 1207-1218.	2.7	59
18	De Novo Mutations in EBF3 Cause a Neurodevelopmental Syndrome. American Journal of Human Genetics, 2017, 100, 138-150.	6.2	52

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19	Rare deleterious mutations of HNRNP genes result in shared neurodevelopmental disorders. Genome Medicine, 2021, 13, 63.	8.2	50
20	A novel RYR2 loss-of-function mutation (I4855M) is associated with left ventricular non-compaction and atypical catecholaminergic polymorphic ventricular tachycardia. Journal of Electrocardiology, 2017, 50, 227-233.	0.9	47
21	De Novo Heterozygous POLR2A Variants Cause a Neurodevelopmental Syndrome with Profound Infantile-Onset Hypotonia. American Journal of Human Genetics, 2019, 105, 283-301.	6.2	46
22	Schinzel–Giedion syndrome: Report of splenopancreatic fusion and proposed diagnostic criteria. American Journal of Medical Genetics, Part A, 2008, 146A, 1299-1306.	1.2	45
23	Etiologies of uterine malformations. American Journal of Medical Genetics, Part A, 2016, 170, 2141-2172.	1.2	45
24	<i>NBEA</i> : Developmental disease gene with early generalized epilepsy phenotypes. Annals of Neurology, 2018, 84, 788-795.	5.3	44
25	Histone H3.3 beyond cancer: Germline mutations in <i>Histone 3 Family 3A and 3B</i> cause a previously unidentified neurodegenerative disorder in 46 patients. Science Advances, 2020, 6, .	10.3	43
26	Illness experience, depression, and anxiety in chronic fatigue syndrome. Journal of Psychosomatic Research, 2002, 52, 461-465.	2.6	41
27	DECIDE: a Decision Support Tool to Facilitate Parents' Choices Regarding Genomeâ€Wide Sequencing. Journal of Genetic Counseling, 2016, 25, 1298-1308.	1.6	36
28	Complex translocation disrupting TCF4 and altering TCF4 isoform expression segregates as mild autosomal dominant intellectual disability. Orphanet Journal of Rare Diseases, 2016, 11, 62.	2.7	35
29	Diffuse angiopathy in Adamsâ€Oliver syndrome associated with truncating <i>DOCK6</i> mutations. American Journal of Medical Genetics, Part A, 2014, 164, 2656-2662.	1.2	32
30	<scp> <i>GREB1L</i></scp> variants in familial and sporadic hereditary urogenital adysplasia and <scp>Mayerâ€Rokitanskyâ€Kusterâ€Hauser</scp> syndrome. Clinical Genetics, 2020, 98, 126-137.	2.0	32
31	Coâ€occurrence of Joubert syndrome and Jeune asphyxiating thoracic dystrophy. American Journal of Medical Genetics, Part A, 2010, 152A, 1411-1419.	1.2	30
32	Missense Variants in the Histone Acetyltransferase Complex Component Gene TRRAP Cause Autism and Syndromic Intellectual Disability. American Journal of Human Genetics, 2019, 104, 530-541.	6.2	30
33	The Canadian Rare Diseases Models and Mechanisms (RDMM) Network: Connecting Understudied Genes to Model Organisms. American Journal of Human Genetics, 2020, 106, 143-152.	6.2	30
34	A Clinical Classification Scheme for Tracheobronchomegaly (Mounier-Kuhn Syndrome). Lung, 2015, 193, 815-822.	3.3	27
35	De novo TBR1 variants cause a neurocognitive phenotype with ID and autistic traits: report of 25 new individuals and review of the literature. European Journal of Human Genetics, 2020, 28, 770-782.	2.8	27
36	KDM5A mutations identified in autism spectrum disorder using forward genetics. ELife, 2020, 9, .	6.0	27

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37	Novel findings and expansion of phenotype in a mosaic <scp>RASopathy</scp> caused by somatic <scp><i>KRAS</i></scp> variants. American Journal of Medical Genetics, Part A, 2021, 185, 2829-2845.	1.2	23
38	Phenotypic spectrum and transcriptomic profile associated with germline variants in TRAF7. Genetics in Medicine, 2020, 22, 1215-1226.	2.4	22
39	Identifying candidate genes for 2p15p16.1 microdeletion syndrome using clinical, genomic, and functional analysis. JCI Insight, 2016, 1, e85461.	5.0	22
40	A characteristic syndrome associated with microduplication of 8q12, inclusive of CHD7. European Journal of Medical Genetics, 2009, 52, 436-439.	1.3	21
41	New developmental syndromes: Understanding the family experience. Journal of Genetic Counseling, 2019, 28, 202-212.	1.6	21
42	Causal Attributions, Perceived Control, and Psychological Adjustment: A Study of Chronic Fatigue Syndrome ¹ . Journal of Applied Social Psychology, 2006, 36, 75-99.	2.0	20
43	FOXP1 haploinsufficiency: Phenotypes beyond behavior and intellectual disability?. American Journal of Medical Genetics, Part A, 2017, 173, 3172-3181.	1.2	18
44	The Genomic Consultation Service: A clinical service designed to improve patient selection for genomeâ€wide sequencing in British Columbia. Molecular Genetics & Genomic Medicine, 2018, 6, 592-600.	1.2	18
45	Heterozygous ANKRD17 loss-of-function variants cause a syndrome with intellectual disability, speech delay, and dysmorphism. American Journal of Human Genetics, 2021, 108, 1138-1150.	6.2	17
46	Treatable inborn errors of metabolism causing neurological symptoms in adults. Molecular Genetics and Metabolism, 2013, 110, 431-438.	1.1	16
47	Evidence of ancillary trigeminal innervation of levator palpebrae in the general population. Journal of Clinical Neuroscience, 2014, 21, 301-304.	1.5	16
48	Unique variants in CLCN3, encoding an endosomal anion/proton exchanger, underlie a spectrum of neurodevelopmental disorders. American Journal of Human Genetics, 2021, 108, 1450-1465.	6.2	16
49	Emphysema in an adult with galactosialidosis linked to a defect in primary elastic fiber assembly. Molecular Genetics and Metabolism, 2012, 106, 99-103.	1.1	15
50	Prenatal ultrasound and MRI findings of temporal and occipital lobe dysplasia in a twin with achondroplasia. Ultrasound in Obstetrics and Gynecology, 2014, 44, 365-368.	1.7	15
51	BMPER variants associated with a novel, attenuated subtype of diaphanospondylodysostosis. Journal of Human Genetics, 2015, 60, 743-747.	2.3	15
52	19p13.2 microduplication causes a Sotos syndromeâ€like phenotype and alters gene expression. Clinical Genetics, 2012, 81, 56-63.	2.0	14
53	Additional post-natal diagnoses following antenatal diagnosis of isolated cleft lip +/â^' palate. Archives of Disease in Childhood: Fetal and Neonatal Edition, 2014, 99, F286-F290.	2.8	14
54	Patient Recall, Interpretation, and Perspective of an Inconclusive Long QT Syndrome Genetic Test Result. Journal of Genetic Counseling, 2017, 26, 150-158.	1.6	13

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55	JARID2 haploinsufficiency is associated with a clinically distinct neurodevelopmental syndrome. Genetics in Medicine, 2021, 23, 374-383.	2.4	13
56	PSEN1 p.Met233Val in a Complex Neurodegenerative Movement and Neuropsychiatric Disorder. Journal of Movement Disorders, 2018, 11, 45-48.	1.3	12
57	Optic atrophy, cataracts, lipodystrophy/lipoatrophy, and peripheral neuropathy caused by a de novo <i>OPA3</i> mutation. Journal of Physical Education and Sports Management, 2017, 3, a001156.	1.2	11
58	Variant Reinterpretation in Survivors of Cardiac Arrest With Preserved Ejection Fraction (the Cardiac) Tj ETQq0 0 Laboratories. Circulation Genomic and Precision Medicine, 2021, 14, e003235.	0 rgBT /C 3.6	overlock 10 Tf 10
59	Bi-allelic variants in the ER quality-control mannosidase gene EDEM3 cause a congenital disorder of glycosylation. American Journal of Human Genetics, 2021, 108, 1342-1349.	6.2	9
60	A Novel Germline Heterozygous BCL11B Variant Causing Severe Atopic Disease and Immune Dysregulation. Frontiers in Immunology, 2021, 12, 788278.	4.8	9
61	How do Physicians Decide to Refer Their Patients for Psychiatric Genetic Counseling? A Qualitative Study of Physicians' Practice. Journal of Genetic Counseling, 2016, 25, 1235-1242.	1.6	8
62	Hypogonadotropic Hypogonadism in Males with Glycogen Storage Disease Type 1. JIMD Reports, 2017, 36, 79-84.	1.5	8
63	Intracranial Calcification after Cord Blood Neonatal Transplantation for Krabbe Disease. Neuropediatrics, 2009, 40, 189-191.	0.6	7
64	Beyond the Electrocardiogram: Mutations in Cardiac Ion Channel Genes Underlie Nonarrhythmic Phenotypes. Clinical Medicine Insights: Cardiology, 2017, 11, 117954681769813.	1.8	7
65	High rate of hypertension in patients with m.3243A>G MELAS mutations and POLG variants. Mitochondrion, 2020, 53, 194-202.	3.4	7
66	Childhood-onset hemiatrophy caused by unilateral morphea. Clinical Dysmorphology, 2009, 18, 213-214.	0.3	6
67	Child Neurology: Krabbe disease. Neurology, 2012, 79, e170-2.	1.1	6
68	Rare disorders have many faces: in silico characterization of rare disorder spectrum. Orphanet Journal of Rare Diseases, 2022, 17, 76.	2.7	6
69	Corneal findings in Parry–Romberg syndrome. Canadian Journal of Ophthalmology, 2014, 49, e2-e5.	0.7	5
70	PIGG variant pathogenicity assessment reveals characteristic features within 19 families. Genetics in Medicine, 2021, 23, 1873-1881.	2.4	5
71	Utilization of telehealth in paediatric genome-wide sequencing: Health services implementation issues in the CAUSES Study. Journal of Telemedicine and Telecare, 2023, 29, 318-327.	2.7	5
72	Strabismus in Children With Intellectual Disability: Part of a Broader Motor Control Phenotype?. Pediatric Neurology, 2019, 100, 87-91.	2.1	4

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73	Renpenning syndrome in a female. American Journal of Medical Genetics, Part A, 2020, 182, 498-503.	1.2	4
74	Cost Analysis of Patients Referred for Inherited Heart Rhythm Disorder Evaluation. Canadian Journal of Cardiology, 2017, 33, 814-821.	1.7	3
75	Novel Exonic Deletions in TTC7A in a Newborn with Multiple Intestinal Atresia and Combined Immunodeficiency. Journal of Clinical Immunology, 2019, 39, 616-619.	3.8	3
76	Return of Results Policies for Genomic Research: Current Practices and the Hearts in Rhythm Organization (HiRO) Approach. Canadian Journal of Cardiology, 2022, 38, 526-535.	1.7	3
77	Can leaky splicing and evasion of premature termination codon surveillance contribute to the phenotypic variability in Alkuraya-Kucinskas syndrome?. European Journal of Medical Genetics, 2022, 65, 104427.	1.3	3
78	The effect of rapid exome sequencing on downstream health care utilization for infants with suspected genetic disorders in an intensive care unit. Genetics in Medicine, 2022, 24, 1675-1683.	2.4	3
79	Anterolateral diaphragmatic hernia with body wall defect understood in relation to the abaxial domain. American Journal of Medical Genetics, Part A, 2014, 164, 1860-1862.	1.2	1
80	Genomic and Cytogenetic Characterization of a Balanced Translocation Disrupting <i>NUP98</i> . Cytogenetic and Genome Research, 2017, 152, 117-121.	1.1	1
81	Secondary biogenic amine deficiencies: genetic etiology, therapeutic interventions, and clinical effects. Neurogenetics, 2021, 22, 251-262.	1.4	1
82	MG-141â€A further report of paediatric cancer and cleidocranial dysplasia raises the possibility of a causative association of weak effect. Journal of Medical Genetics, 2015, 52, A12.1-A12.	3.2	0
83	Familial impairment of vocal cord mobility in childhood with clubfoot. Clinical Dysmorphology, 2018, 27, 116-121.	0.3	0
84	Integration of genetic counsellors in genomic testing triage: Outcomes of a genomic consultation service in British Columbia, Canada. European Journal of Medical Genetics, 2021, 64, 104024.	1.3	0
85	Fetal Progeria: Prenatal Sonographic Findings in Petty Syndrome. Journal of Ultrasound in Medicine, 2013, 32, 881-883.	1.7	0