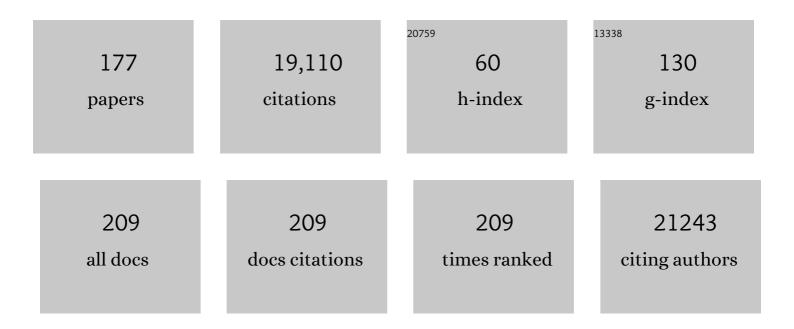
Heather C Mefford

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | De novo mutations in epileptic encephalopathies. Nature, 2013, 501, 217-221. | 13.7 | 1,351 |
| 2 | Exome sequencing identifies MLL2 mutations as a cause of Kabuki syndrome. Nature Genetics, 2010, 42, 790-793. | 9.4 | 1,238 |
| 3 | Multiplex Targeted Sequencing Identifies Recurrently Mutated Genes in Autism Spectrum Disorders. Science, 2012, 338, 1619-1622. | 6.0 | 1,133 |
| 4 | Recurrent Rearrangements of Chromosome 1q21.1 and Variable Pediatric Phenotypes. New England Journal of Medicine, 2008, 359, 1685-1699. | 13.9 | 663 |
| 5 | Disruptive CHD8 Mutations Define a Subtype of Autism Early in Development. Cell, 2014, 158, 263-276. | 13.5 | 637 |
| 6 | SFARI Gene 2.0: a community-driven knowledgebase for the autism spectrum disorders (ASDs). Molecular Autism, 2013, 4, 36. | 2.6 | 632 |
| 7 | Targeted resequencing in epileptic encephalopathies identifies de novo mutations in CHD2 and SYNGAP1. Nature Genetics, 2013, 45, 825-830. | 9.4 | 589 |
| 8 | Refining analyses of copy number variation identifies specific genes associated with developmental delay. Nature Genetics, 2014, 46, 1063-1071. | 9.4 | 583 |
| 9 | A recurrent 16p12.1 microdeletion supports a two-hit model for severe developmental delay. Nature Genetics, 2010, 42, 203-209. | 9.4 | 539 |
| 10 | Population Analysis of Large Copy Number Variants and Hotspots of Human Genetic Disease. American Journal of Human Genetics, 2009, 84, 148-161. | 2.6 | 530 |
| 11 | 15q13.3 microdeletions increase risk of idiopathic generalized epilepsy. Nature Genetics, 2009, 41, 160-162. | 9.4 | 511 |
| 12 | A recurrent 15q13.3 microdeletion syndrome associated with mental retardation and seizures. Nature Genetics, 2008, 40, 322-328. | 9.4 | 509 |
| 13 | Genome-Wide Copy Number Variation in Epilepsy: Novel Susceptibility Loci in Idiopathic Generalized and Focal Epilepsies. PLoS Genetics, 2010, 6, e1000962. | 1.5 | 414 |
| 14 | Recurrent microdeletions at 15q11.2 and 16p13.11 predispose to idiopathic generalized epilepsies. Brain, 2010, 133, 23-32. | 3.7 | 406 |
| 15 | High Rate of Recurrent De Novo Mutations in Developmental and Epileptic Encephalopathies. American Journal of Human Genetics, 2017, 101, 664-685. | 2.6 | 337 |
| 16 | GRIN2A mutations cause epilepsy-aphasia spectrum disorders. Nature Genetics, 2013, 45, 1073-1076. | 9.4 | 326 |
| 17 | Genomics, Intellectual Disability, and Autism. New England Journal of Medicine, 2012, 366, 733-743. | 13.9 | 276 |
| 18 | The phenotypic spectrum of <i>SCN8A</i> encephalopathy. Neurology, 2015, 84, 480-489. | 1.5 | 246 |

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| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | <i>GABRA1</i> and <i>STXBP1</i> : Novel genetic causes of Dravet syndrome. Neurology, 2014, 82, 1245-1253. | 1.5 | 229 |
| 20 | Recurrent Reciprocal Genomic Rearrangements of 17q12 Are Associated with Renal Disease, Diabetes, and Epilepsy. American Journal of Human Genetics, 2007, 81, 1057-1069. | 2.6 | 222 |
| 21 | Rare copy number variants are an important cause of epileptic encephalopathies. Annals of Neurology, 2011, 70, 974-985. | 2.8 | 222 |
| 22 | Clinical whole-genome sequencing in severe early-onset epilepsy reveals new genes and improves molecular diagnosis. Human Molecular Genetics, 2014, 23, 3200-3211. | 1.4 | 222 |
| 23 | Familial and sporadic 15q13.3 microdeletions in idiopathic generalized epilepsy: precedent for disorders with complex inheritance. Human Molecular Genetics, 2009, 18, 3626-3631. | 1.4 | 211 |
| 24 | <i>SCN2A</i> encephalopathy. Neurology, 2015, 85, 958-966. | 1.5 | 211 |
| 25 | Duplication hotspots, rare genomic disorders, and common disease. Current Opinion in Genetics and Development, 2009, 19, 196-204. | 1.5 | 191 |
| 26 | Ultra-rare genetic variation in common epilepsies: a case-control sequencing study. Lancet Neurology, The, 2017, 16, 135-143. | 4.9 | 190 |
| 27 | <i>GRIN2B</i> encephalopathy: novel findings on phenotype, variant clustering, functional consequences and treatment aspects. Journal of Medical Genetics, 2017, 54, 460-470. | 1.5 | 190 |
| 28 | Recurrent 200-kb deletions of 16p11.2 that include the SH2B1 gene are associated with developmental delay and obesity. Genetics in Medicine, 2010, 12, 641-647. | 1.1 | 178 |
| 29 | Advancing epilepsy genetics in the genomic era. Genome Medicine, 2015, 7, 91. | 3.6 | 173 |
| 30 | Mutations in the GABA Transporter SLC6A1 Cause Epilepsy with Myoclonic-Atonic Seizures. American Journal of Human Genetics, 2015, 96, 808-815. | 2.6 | 173 |
| 31 | The genetic landscape of infantile spasms. Human Molecular Genetics, 2014, 23, 4846-4858. | 1.4 | 156 |
| 32 | AMPA receptor GluA2 subunit defects are a cause of neurodevelopmental disorders. Nature Communications, 2019, 10, 3094. | 5.8 | 150 |
| 33 | The Genetics of Microdeletion and Microduplication Syndromes: An Update. Annual Review of Genomics and Human Genetics, 2014, 15, 215-244. | 2.5 | 145 |
| 34 | Copy number variants are frequent in genetic generalized epilepsy with intellectual disability. Neurology, 2013, 81, 1507-1514. | 1.5 | 140 |
| 35 | Genetic and neurodevelopmental spectrum of <i>SYNGAP1</i> -associated intellectual disability and epilepsy. Journal of Medical Genetics, 2016, 53, 511-522. | 1.5 | 135 |
| 36 | Clinical phenotype of the recurrent 1q21.1 copy-number variant. Genetics in Medicine, 2016, 18, 341-349. | 1.1 | 134 |

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 37 | <i>SYNGAP1</i> encephalopathy. Neurology, 2019, 92, e96-e107. | 1.5 | 131 |
| 38 | A method for rapid, targeted CNV genotyping identifies rare variants associated with neurocognitive disease. Genome Research, 2009, 19, 1579-1585. | 2.4 | 118 |
| 39 | Mutations in <i><scp>KCNT</scp>1</i> cause a spectrum of focal epilepsies. Epilepsia, 2015, 56, e114-20. | 2.6 | 117 |
| 40 | <i>CHD2</i> variants are a risk factor for photosensitivity in epilepsy. Brain, 2015, 138, 1198-1208. | 3.7 | 112 |
| 41 | Targeted long-read sequencing identifies missing disease-causing variation. American Journal of Human Genetics, 2021, 108, 1436-1449. | 2.6 | 105 |
| 42 | Parental Mosaicism in "De Novo―Epileptic Encephalopathies. New England Journal of Medicine, 2018, 378, 1646-1648. | 13.9 | 104 |
| 43 | Germline De Novo Mutations in GNB1 Cause Severe Neurodevelopmental Disability, Hypotonia, and Seizures. American Journal of Human Genetics, 2016, 98, 1001-1010. | 2.6 | 102 |
| 44 | <i>SCN8A</i> encephalopathy: Research progress and prospects. Epilepsia, 2016, 57, 1027-1035. | 2.6 | 101 |
| 45 | Defining the phenotypic spectrum of <i>SLC6A1</i> mutations. Epilepsia, 2018, 59, 389-402. | 2.6 | 99 |
| 46 | Not all <i>SCN1A</i> epileptic encephalopathies are Dravet syndrome. Neurology, 2017, 89, 1035-1042. | 1.5 | 97 |
| 47 | The Genetic Landscape of Epilepsy of Infancy with Migrating Focal Seizures. Annals of Neurology, 2019, 86, 821-831. | 2.8 | 96 |
| 48 | <i>TRIO</i> loss of function is associated with mild intellectual disability and affects dendritic branching and synapse function. Human Molecular Genetics, 2016, 25, 892-902. | 1.4 | 94 |
| 49 | Complex Compound Inheritance of Lethal Lung Developmental Disorders Due to Disruption of the TBX-FGF Pathway. American Journal of Human Genetics, 2019, 104, 213-228. | 2.6 | 90 |
| 50 | Mutations in <i>GABRB3</i> . Neurology, 2017, 88, 483-492. | 1.5 | 87 |
| 51 | De Novo Pathogenic Variants in CACNA1E Cause Developmental and Epileptic Encephalopathy with Contractures, Macrocephaly, and Dyskinesias. American Journal of Human Genetics, 2018, 103, 666-678. | 2.6 | 87 |
| 52 | GGC Repeat Expansion and Exon 1 Methylation of XYLT1 Is a Common Pathogenic Variant in Baratela-Scott Syndrome. American Journal of Human Genetics, 2019, 104, 35-44. | 2.6 | 81 |
| 53 | <i>CHD2</i> myoclonic encephalopathy is frequently associated with self-induced seizures. Neurology, 2015, 84, 951-958. | 1.5 | 79 |
| 54 | Primer Part 1—The building blocks of epilepsy genetics. Epilepsia, 2016, 57, 861-868. | 2.6 | 77 |

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 55 | A populationâ€based costâ€effectiveness study of early genetic testing in severe epilepsies of infancy. Epilepsia, 2018, 59, 1177-1187. | 2.6 | 77 |
| 56 | Aberrant Inclusion of a Poison Exon Causes Dravet Syndrome and Related SCN1A-Associated Genetic Epilepsies. American Journal of Human Genetics, 2018, 103, 1022-1029. | 2.6 | 76 |
| 57 | Autism and developmental disability caused by <i>KCNQ3</i> gainâ€ofâ€function variants. Annals of Neurology, 2019, 86, 181-192. | 2.8 | 73 |
| 58 | De novo mutations in GRIN1 cause extensive bilateral polymicrogyria. Brain, 2018, 141, 698-712. | 3.7 | 72 |
| 59 | Dominant <i>KCNA2</i> mutation causes episodic ataxia and pharmacoresponsive epilepsy. Neurology, 2016, 87, 1975-1984. | 1.5 | 71 |
| 60 | Copy number variation analysis in singleâ€suture craniosynostosis: Multiple rare variants including <i>RUNX2</i> duplication in two cousins with metopic craniosynostosis. American Journal of Medical Genetics, Part A, 2010, 152A, 2203-2210. | 0.7 | 69 |
| 61 | Seizures Are Regulated by Ubiquitin-specific Peptidase 9 X-linked (USP9X), a De-Ubiquitinase. PLoS Genetics, 2015, 11, e1005022. | 1.5 | 66 |
| 62 | Epileptic spasms are a feature of <i>DEPDC5</i> mTORopathy. Neurology: Genetics, 2015, 1, e17. | 0.9 | 63 |
| 63 | 16p11.2 600 kb Duplications confer risk for typical and atypical Rolandic epilepsy. Human Molecular Genetics, 2014, 23, 6069-6080. | 1.4 | 61 |
| 64 | Diagnostic implications of genetic copy number variation in epilepsy plus. Epilepsia, 2019, 60, 689-706. | 2.6 | 61 |
| 65 | Exonâ€disrupting deletions of <scp><i>NRXN1</i></scp> in idiopathic generalized epilepsy. Epilepsia, 2013, 54, 256-264. | 2.6 | 59 |
| 66 | Mutations Affecting the SAND Domain of DEAF1 Cause Intellectual Disability with Severe Speech Impairment and Behavioral Problems. American Journal of Human Genetics, 2014, 94, 649-661. | 2.6 | 59 |
| 67 | Copy number variant analysis from exome data in 349 patients with epileptic encephalopathy. Annals of Neurology, 2015, 78, 323-328. | 2.8 | 59 |
| 68 | Epilepsy and the new cytogenetics. Epilepsia, 2011, 52, 423-432. | 2.6 | 56 |
| 69 | Clinical phenotype of ASD-associated DYRK1A haploinsufficiency. Molecular Autism, 2017, 8, 54. | 2.6 | 55 |
| 70 | Comparative sequencing of a multicopy subtelomeric region containing olfactory receptor genes reveals multiple interactions between non-homologous chromosomes. Human Molecular Genetics, 2001, 10, 2363-2372. | 1.4 | 51 |
| 71 | A targeted resequencing gene panel for focal epilepsy. Neurology, 2016, 86, 1605-1612. | 1.5 | 48 |
| 72 | De Novo Variants in the F-Box Protein FBXO11 in 20 Individuals with a Variable Neurodevelopmental Disorder. American Journal of Human Genetics, 2018, 103, 305-316. | 2.6 | 48 |

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 73 | De novo mutations of <i>KIAA2022</i> in females cause intellectual disability and intractable epilepsy. Journal of Medical Genetics, 2016, 53, 850-858. | 1.5 | 47 |
| 74 | Recent advances in epilepsy genomics and genetic testing. F1000Research, 2020, 9, 185. | 0.8 | 47 |
| 75 | Somatic mutation: The hidden genetics of brain malformations and focal epilepsies. Epilepsy Research, 2019, 155, 106161. | 0.8 | 45 |
| 76 | Bi-allelic Loss-of-Function CACNA1B Mutations in Progressive Epilepsy-Dyskinesia. American Journal of Human Genetics, 2019, 104, 948-956. | 2.6 | 45 |
| 77 | A homozygous <i>B3GAT3</i> mutation causes a severe syndrome with multiple fractures, expanding the phenotype of linkeropathy syndromes. American Journal of Medical Genetics, Part A, 2015, 167, 2691-2696. | 0.7 | 44 |
| 78 | Severe infantile onset developmental and epileptic encephalopathy caused by mutations in autophagy gene <i><scp>WDR</scp>45</i> . Epilepsia, 2018, 59, e5-e13. | 2.6 | 44 |
| 79 | <i>NBEA</i> : Developmental disease gene with early generalized epilepsy phenotypes. Annals of Neurology, 2018, 84, 788-795. | 2.8 | 44 |
| 80 | De Novo Mutations in PPP3CA Cause Severe Neurodevelopmental Disease with Seizures. American Journal of Human Genetics, 2017, 101, 516-524. | 2.6 | 43 |
| 81 | Disruptive mutations in TANC2 define a neurodevelopmental syndrome associated with psychiatric disorders. Nature Communications, 2019, 10, 4679. | 5.8 | 43 |
| 82 | Microdeletion syndromes. Current Opinion in Genetics and Development, 2013, 23, 232-239. | 1.5 | 42 |
| 83 | Genetic contribution to common epilepsies. Current Opinion in Neurology, 2011, 24, 140-145. | 1.8 | 41 |
| 84 | TANGO2: expanding the clinical phenotype and spectrum of pathogenic variants. Genetics in Medicine, 2019, 21, 601-607. | 1.1 | 41 |
| 85 | Further clinical and molecular delineation of the 15q24 microdeletion syndrome. Journal of Medical Genetics, 2012, 49, 110-118. | 1.5 | 40 |
| 86 | Simultaneous impairment of neuronal and metabolic function of mutated gephyrin in a patient with epileptic encephalopathy. EMBO Molecular Medicine, 2015, 7, 1580-1594. | 3.3 | 39 |
| 87 | Genetic heterogeneity in infantile spasms. Epilepsy Research, 2019, 156, 106181. | 0.8 | 38 |
| 88 | Genotype to phenotype—discovery and characterization of novel genomic disorders in a "genotype-first―era. Genetics in Medicine, 2009, 11, 836-842. | 1.1 | 37 |
| 89 | Loss of function of the retinoid-related nuclear receptor (RORB) gene and epilepsy. European Journal of Human Genetics, 2016, 24, 1761-1770. | 1.4 | 36 |
| 90 | The Impact of Rapid Exome Sequencing on Medical Management of Critically Ill Children. Journal of Pediatrics, 2020, 226, 202-212.e1. | 0.9 | 35 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 91 | Intragenic deletions of <i>ALDH7A1</i> in pyridoxine-dependent epilepsy caused by <i>Alu</i> - <i>Alu</i> recombination. Neurology, 2015, 85, 756-762. | 1.5 | 34 |
| 92 | De novo loss-of-function mutations in WAC cause a recognizable intellectual disability syndrome and learning deficits in Drosophila. European Journal of Human Genetics, 2016, 24, 1145-1153. | 1.4 | 34 |
| 93 | The ClinGen Epilepsy Gene Curation Expert Panel—Bridging the divide between clinical domain knowledge and formal gene curation criteria. Human Mutation, 2018, 39, 1476-1484. | 1.1 | 33 |
| 94 | Expanding the genetic and phenotypic relevance of <i>KCNB1</i> variants in developmental and epileptic encephalopathies: 27 new patients and overview of the literature. Human Mutation, 2020, 41, 69-80. | 1.1 | 33 |
| 95 | Cenetically complex epilepsies, copy number variants and syndrome constellations. Genome Medicine, 2010, 2, 71. | 3.6 | 32 |
| 96 | A genotype-first approach identifies an intellectual disability-overweight syndrome caused by PHIP haploinsufficiency. European Journal of Human Genetics, 2018, 26, 54-63. | 1.4 | 32 |
| 97 | The severe epilepsy syndromes of infancy: A populationâ€based study. Epilepsia, 2021, 62, 358-370. | 2.6 | 31 |
| 98 | Next-Generation Sequencing in Intellectual Disability. Journal of Pediatric Genetics, 2015, 04, 128-135. | 0.3 | 30 |
| 99 | Expanding clinical phenotype in <i>CACNA1C</i> related disorders: From neonatal onset severe epileptic encephalopathy to lateâ€onset epilepsy. American Journal of Medical Genetics, Part A, 2018, 176, 2733-2739. | 0.7 | 30 |
| 100 | ZMIZ1 Variants Cause a Syndromic Neurodevelopmental Disorder. American Journal of Human Genetics, 2019, 104, 319-330. | 2.6 | 30 |
| 101 | Absence seizures with intellectual disability as a phenotype of the 15q13.3 microdeletion syndrome. Epilepsia, 2011, 52, e194-8. | 2.6 | 29 |
| 102 | CNVs in Epilepsy. Current Genetic Medicine Reports, 2014, 2, 162-167. | 1.9 | 28 |
| 103 | Severe neurocognitive and growth disorders due to variation in <i>THOC2</i> , an essential component of nuclear mRNA export machinery. Human Mutation, 2018, 39, 1126-1138. | 1.1 | 28 |
| 104 | NEXMIF encephalopathy: an X-linked disorder with male and female phenotypic patterns. Genetics in Medicine, 2021, 23, 363-373. | 1.1 | 28 |
| 105 | Loss of function in <i>ROBO1</i> is associated with tetralogy of Fallot and septal defects. Journal of Medical Genetics, 2017, 54, 825-829. | 1.5 | 27 |
| 106 | Poison exons in neurodevelopment and disease. Current Opinion in Genetics and Development, 2020, 65, 98-102. | 1.5 | 26 |
| 107 | Mutations of the Transcriptional Corepressor ZMYM2 Cause Syndromic Urinary Tract Malformations. American Journal of Human Genetics, 2020, 107, 727-742. | 2.6 | 25 |
| 108 | Severe speech impairment is a distinguishing feature of <i>FOXP1</i> â€related disorder. Developmental Medicine and Child Neurology, 2021, 63, 1417-1426. | 1.1 | 24 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 109 | Multiplex families with epilepsy. Neurology, 2016, 86, 713-722. | 1.5 | 23 |
| 110 | Recurrent duplications of 17q12 associated with variable phenotypes. American Journal of Medical Genetics, Part A, 2015, 167, 3038-3045. | 0.7 | 22 |
| 111 | Clinical Genetic Testing in Epilepsy. Epilepsy Currents, 2015, 15, 197-201. | 0.4 | 22 |
| 112 | Clinical Phenotypes of Carriers of Mutations in CHD8 or Its Conserved Target Genes. Biological Psychiatry, 2020, 87, 123-131. | 0.7 | 22 |
| 113 | lterative phenotyping of 15q11.2, 15q13.3 and 16p13.11 microdeletion carriers in pediatric epilepsies. Epilepsy Research, 2014, 108, 109-116. | 0.8 | 21 |
| 114 | Pathogenic Variants in CEP85L Cause Sporadic and Familial Posterior Predominant Lissencephaly. Neuron, 2020, 106, 237-245.e8. | 3.8 | 21 |
| 115 | The epilepsy phenotypic spectrum associated with a recurrent <i>CUX2</i> variant. Annals of Neurology, 2018, 83, 926-934. | 2.8 | 20 |
| 116 | De novo and biallelic DEAF1 variants cause a phenotypic spectrum. Genetics in Medicine, 2019, 21, 2059-2069. | 1.1 | 20 |
| 117 | De novo variants in SNAP25 cause an early-onset developmental and epileptic encephalopathy. Genetics in Medicine, 2021, 23, 653-660. | 1.1 | 20 |
| 118 | Genetic investigations of the epileptic encephalopathies. Progress in Brain Research, 2016, 226, 35-60. | 0.9 | 19 |
| 119 | Biallelic <i>PI4KA</i> variants cause a novel neurodevelopmental syndrome with hypomyelinating leukodystrophy. Brain, 2021, 144, 2659-2669. | 3.7 | 19 |
| 120 | Epilepsy due to 20q13.33 subtelomere deletion masquerading as pyridoxineâ€dependent epilepsy. American Journal of Medical Genetics, Part A, 2012, 158A, 3190-3195. | 0.7 | 18 |
| 121 | Deletions of 16p11.2 and 19p13.2 in a family with intellectual disability and generalized epilepsy. American Journal of Medical Genetics, Part A, 2013, 161, 1722-1725. | 0.7 | 18 |
| 122 | <i><scp>BRAT</scp>1</i> encephalopathy: a recessive cause of epilepsy of infancy with migrating focal seizures. Developmental Medicine and Child Neurology, 2020, 62, 1096-1099. | 1.1 | 18 |
| 123 | Bi-allelic Loss-of-Function Variants in NUP188 Cause a Recognizable Syndrome Characterized by Neurologic, Ocular, and Cardiac Abnormalities. American Journal of Human Genetics, 2020, 106, 623-631. | 2.6 | 18 |
| 124 | Genetic literacy series: Primer part 2—Paradigm shifts in epilepsy genetics. Epilepsia, 2018, 59, 1138-1147. | 2.6 | 17 |
| 125 | Developmental and epilepsy spectrum of <i>KCNB1</i> encephalopathy with longâ€ŧerm outcome. Epilepsia, 2020, 61, 2461-2473. | 2.6 | 17 |
| 126 | The unexpected role of copy number variations in juvenile myoclonic epilepsy. Epilepsy and Behavior, 2013, 28, S66-S68. | 0.9 | 16 |

| # | Article | IF | CITATIONS |
|-----|---|-----|-----------|
| 127 | Investigating the genetic basis of feverâ€associated syndromic epilepsies using copy number variation analysis. Epilepsia, 2015, 56, e26-32. | 2.6 | 16 |
| 128 | Levetiracetam efficacy in PCDH19 Girls Clustering Epilepsy. European Journal of Paediatric Neurology, 2020, 24, 142-147. | 0.7 | 16 |
| 129 | Germline variants in tumor suppressor FBXW7 lead to impaired ubiquitination and a neurodevelopmental syndrome. American Journal of Human Genetics, 2022, 109, 601-617. | 2.6 | 16 |
| 130 | Five patients with a chromosome 1q21.1 triplication show macrocephaly, increased weight and facial similarities. European Journal of Medical Genetics, 2015, 58, 503-508. | 0.7 | 15 |
| 131 | Missense variants in <i>CTNNB1</i> can be associated with vitreoretinopathy—Seven new cases of <i>CTNNB1</i> â€associated neurodevelopmental disorder including a previously unreported retinal phenotype. Molecular Genetics & Genomic Medicine, 2021, 9, e1542. | 0.6 | 15 |
| 132 | Inherited <i>RORB</i> pathogenic variants: Overlap of photosensitive genetic generalized and occipital lobe epilepsy. Epilepsia, 2020, 61, e23-e29. | 2.6 | 14 |
| 133 | The epileptology of GNB5 encephalopathy. Epilepsia, 2019, 60, e121-e127. | 2.6 | 13 |
| 134 | Variants in GNAI1 cause a syndrome associated with variable features including developmental delay, seizures, and hypotonia. Genetics in Medicine, 2021, 23, 881-887. | 1.1 | 13 |
| 135 | Copy Number Matters in Epilepsy. Epilepsy Currents, 2015, 15, 180-182. | 0.4 | 12 |
| 136 | Dravet syndrome in South African infants: Tools for an early diagnosis. Seizure: the Journal of the British Epilepsy Association, 2018, 62, 99-105. | 0.9 | 12 |
| 137 | Beyond the single nucleotide variant in epilepsy genetics. Nature Reviews Neurology, 2014, 10, 490-491. | 4.9 | 11 |
| 138 | Description of a new oncogenic mechanism for atypical teratoid rhabdoid tumors in patients with ring chromosome 22. American Journal of Medical Genetics, Part A, 2017, 173, 245-249. | 0.7 | 11 |
| 139 | Developmental and epileptic encephalopathy: Personal utility of a genetic diagnosis for families. Epilepsia Open, 2021, 6, 149-159. | 1.3 | 11 |
| 140 | The Road to Diagnosis: Shortening the Diagnostic Odyssey in Epilepsy. Epilepsy Currents, 2019, 19, 307-309. | 0.4 | 10 |
| 141 | The phenotypic spectrum of Xâ€linked, infantile onset <i>ALG13</i> â€related developmental and epileptic encephalopathy. Epilepsia, 2021, 62, 325-334. | 2.6 | 10 |
| 142 | 2014 Epilepsy Benchmarks Area I: Understanding the Causes of the Epilepsies and Epilepsy-Related Neurologic, Psychiatric, and Somatic Conditions. Epilepsy Currents, 2016, 16, 182-186. | 0.4 | 9 |
| 143 | Return of individual results in epilepsy genomic research: A view from the field. Epilepsia, 2018, 59, 1635-1642. | 2.6 | 9 |
| 144 | Three novel patients with epileptic encephalopathy due to biallelic mutations in the <scp><i>PLCB1</i></scp> gene. Clinical Genetics, 2020, 97, 477-482. | 1.0 | 9 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 145 | Epilepsy Benchmarks Area III: Improved Treatment Options for Controlling Seizures and Epilepsy-Related Conditions Without Side Effects. Epilepsy Currents, 2020, 20, 23S-30S. | 0.4 | 9 |
| 146 | 15q13.3 microdeletions in a prospectively recruited cohort of patients with idiopathic generalized epilepsy in Bulgaria. Epilepsy Research, 2013, 104, 241-245. | 0.8 | 8 |
| 147 | <i>FBXO28</i> causes developmental and epileptic encephalopathy with profound intellectual disability. Epilepsia, 2021, 62, e13-e21. | 2.6 | 8 |
| 148 | Phenotypic Spectrum of Seizure Disorders in MBD5-Associated Neurodevelopmental Disorder. Neurology: Genetics, 2021, 7, e579. | 0.9 | 8 |
| 149 | Association of ultraâ€rare coding variants with genetic generalized epilepsy: A case–control whole exome sequencing study. Epilepsia, 2022, 63, 723-735. | 2.6 | 8 |
| 150 | Biallelic <i>ADAM22</i> pathogenic variants cause progressive encephalopathy and infantile-onset refractory epilepsy. Brain, 2022, 145, 2301-2312. | 3.7 | 8 |
| 151 | A de novo inâ€frame deletion of <i>CASK</i> gene causes early onset infantile spasms and supratentorial cerebral malformation in a female patient. American Journal of Medical Genetics, Part A, 2018, 176, 2425-2429. | 0.7 | 7 |
| 152 | Double somatic mosaicism in a child with Dravet syndrome. Neurology: Genetics, 2019, 5, e333. | 0.9 | 7 |
| 153 | Reanalysis and optimisation of bioinformatic pipelines is critical for mutation detection. Human Mutation, 2019, 40, 374-379. | 1.1 | 7 |
| 154 | Rare variants in KDR, encoding VEGF Receptor 2, are associated with tetralogy of Fallot. Genetics in Medicine, 2021, 23, 1952-1960. | 1.1 | 7 |
| 155 | Pathogenic <scp><i>MAST3</i></scp> Variants in the <scp>STK</scp> Domain Are Associated with Epilepsy. Annals of Neurology, 2021, 90, 274-284. | 2.8 | 7 |
| 156 | Evaluation of multiple putative risk alleles within the 15q13.3 region for genetic generalized epilepsy. Epilepsy Research, 2015, 117, 70-73. | 0.8 | 6 |
| 157 | <i>De novo FZR1</i> loss-of-function variants cause developmental and epileptic encephalopathies. Brain, 2022, 145, 1684-1697. | 3.7 | 5 |
| 158 | <i>SCN1A</i> mutations in focal epilepsy with auditory features: widening the spectrum of GEFS <i>plus</i> . Epileptic Disorders, 2019, 21, 185-191. | 0.7 | 5 |
| 159 | Clarifying the role of the 22q11.2 microdeletion in juvenile myoclonic epilepsy. Epilepsy and Behavior, 2013, 29, 589-590. | 0.9 | 4 |
| 160 | Expanding role of GABAA receptors in generalised epilepsies. Lancet Neurology, The, 2018, 17, 657-658. | 4.9 | 4 |
| 161 | Genetic convergence of developmental and epileptic encephalopathies and intellectual disability. Developmental Medicine and Child Neurology, 2021, 63, 1441-1447. | 1.1 | 4 |
| 162 | Diagnostic Considerations in the Epilepsies—Testing Strategies, Test Type Advantages, and Limitations. Neurotherapeutics, 2021, 18, 1468-1477. | 2.1 | 4 |

| # | Article | IF | CITATIONS |
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| 163 | A recurrent, de novo pathogenic variant in ARPC4 disrupts actin filament formation and causes a neurodevelopmental disorder with microcephaly and speech delay. Human Genetics and Genomics Advances, 2021, 3, 100072. | 1.0 | 4 |
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