

Genetic Testing for Epilepsy: Sequence Analysis and Exon-Level Deletion/Duplication Testing of up to 127 Genes

Tests Included:

- Comprehensive Epilepsy Panel
- Infantile Epilepsy Panel
- Childhood Epilepsy Panel
- STAT Epilepsy Panel
- Progressive Myoclonic Epilepsy Panel
- Rett/Angelman Syndrome Epilepsy Panel

Clinical Features:

Epilepsy is defined by the occurrence of at least two unprovoked seizures occurring more than 24 hours apart. It is a common neurological disorder that affects at least 0.8% of the population. The International League against Epilepsy (ILAE) classifies seizures into two main categories.¹ **Generalized epileptic seizures** originate in and rapidly engage both cerebral hemispheres. Tonic-clonic, absence, myoclonic, clonic, tonic, and atonic seizures are all types of generalized seizures. **Focal seizures** originate from neuronal networks within a single hemisphere. Traditionally, focal seizures have been classified as “simple partial seizures,” which do not result in an alteration of consciousness, and “complex partial seizures,” which cause a change in behavior or consciousness.

Some types of seizures, such as infantile spasms, do not fit into either category and remain unclassified. Seizures can be self-limiting or controlled by standard therapeutic treatments in some cases; however, individuals with epileptic encephalopathy have severe seizures that are refractory to treatment, leading to cognitive and behavioral impairment secondary to the epileptic activity. Epilepsy may be an isolated neurological symptom, or it may occur in association with other neurological symptoms or medical problems.² Some individuals with epilepsy are diagnosed with an electroclinical syndrome such as West or Ohtahara syndrome based on the presence of characteristic EEG findings and the clinical and family history.¹

Inheritance Pattern/Genetics:

Epilepsy can be caused by genetic disorders, metabolic diseases, trauma, infection, and structural brain abnormalities, although the cause is not known in many cases. A genetic etiology underlies epilepsy in approximately 40% of individuals.³ Genes have been identified that cause both generalized seizures and focal seizures, as well as unclassified epilepsy types such as infantile spasms. The genetic etiology of idiopathic generalized epilepsy (IGE) is

frequently complex because it is due to a combination of multiple genetic factors that each confer a small risk for epilepsy and may be modified by environmental influences.³ Currently, approximately 2% of patients with IGE harbor an identifiable variant in a single gene associated with Mendelian inheritance of epilepsy.⁴ However, the percentage of patients with Mendelian epilepsy is higher for specific epilepsy types such as infantile spasms, benign familial neonatal and neonatal-infantile seizures (BFNS and BFNIS), and others.^{3,5,6,7} The inheritance pattern can be autosomal dominant, autosomal recessive, or X-linked. Pathogenic variants in a single gene may be associated with different types of seizures (clinical heterogeneity), and conversely, pathogenic variants in different genes can cause the same epilepsy phenotype (genetic heterogeneity).

The Epilepsy Panels at GeneDx includes sequencing and deletion/duplication analysis of genes causing Mendelian forms of epilepsy. Many of these genes encode subunits of ion channels involved in stabilizing or propagating neuronal activity, including components of the voltage-gated sodium, potassium, and calcium channels and the ligand-gated gamma-aminobutyric (GABA) channel.^{5,7,8,9} The panels also includes non-ion channel genes associated with a variety of neurotransmitter disorders, storage and other neurometabolic disorders, as well as genes causing syndromic forms of epilepsy, many of which are involved in transcriptional activation or repression.^{5,6,8,10,11,12}

Several different Epilepsy Panels are available at GeneDx, based on seizure age-of-onset and/or clinical presentation. The complete list of genes included on each panel is included in the table in the appendix. Available panels include:

- Comprehensive Epilepsy Panel (127 genes)
- Infantile Epilepsy Panel (111 genes)
- Childhood Epilepsy Panel (84 genes)
- STAT Epilepsy Panel* (26 genes, many with treatment implications)
- Progressive Myoclonic Epilepsy Panel* (18 genes)
- Rett/Angelman Syndrome Panel*

* Separate information sheets are available on our website (www.genedx.com) with more specific information about our STAT Epilepsy, Rett/Angelman Syndrome, and Progressive Myoclonic Epilepsy testing panels. In addition to the epilepsy panels mentioned above, GeneDx also offers panels or single gene testing for many other disorders causing epilepsy, such as tuberous sclerosis complex, mitochondrial disorders, and metabolic disorders.

Test Methods:

Using genomic DNA from the submitted specimen, the complete coding regions and splice site junctions of the genes on this panel are enriched using a proprietary targeted capture system

developed by GeneDx for next-generation sequencing with CNV calling (NGS-CNV). The enriched targets are simultaneously sequenced with paired-end reads on an Illumina platform. Bi-directional sequence reads are assembled and aligned to reference sequences based on NCBI RefSeq transcripts and human genome build GRCh37/UCSC hg19. After gene specific filtering, data are analyzed to identify sequence variants and most deletions and duplications involving coding exons; however, technical limitations and inherent sequence properties effectively reduce this resolution for some genes. Alternative sequencing or copy number detection methods are used to analyze or confirm regions with inadequate sequence or copy number data by NGS. Reportable variants include pathogenic variants, likely pathogenic variants and variants of uncertain significance. Likely benign and benign variants, if present, are not routinely reported but are available upon request.

The technical sensitivity of sequencing is estimated to be > 99% at detecting single nucleotide events. It will not reliably detect deletions greater than 20 base pairs, insertions or rearrangements greater than 10 base pairs, or low-level mosaicism. The copy number assessment methods used with this test cannot reliably detect copy number variants of less than 500 base pairs or mosaicism and cannot identify balanced chromosome aberrations. Assessment of exon-level copy number events is dependent on the inherent sequence properties of the targeted regions, including shared homology and exon size. For the *CHRNA7*, *MAGI2*, and *PLCB1* gene(s), deletion/duplication analysis, but not sequencing was performed. For the *DNM1*, *FOXP1*, and *SLC6A8* gene(s), sequencing but not deletion/duplication analysis, was performed. If indicated based on the patient's specific clinical features, multiplex ligation-dependent probe amplification (MLPA) of the *FOXP1* and *SLC6A8* genes is available as a separate test (test code 906).

Test Sensitivity:

The clinical sensitivity of sequencing and deletion/duplication analysis of the genes included in the Comprehensive Epilepsy Panel depends in part on the patient's clinical phenotype. In a prior study, 31% of individuals with infantile spasms who were tested using an epilepsy gene panel were found to harbor definitive pathogenic variant(s) to explain the phenotype (Wirrell et al., 2015). Overall, 17-20% of epileptic encephalopathies have an identifiable genetic etiology (EpiPM Consortium, 2015). Specific information about the diagnostic yield for each gene in selected populations is summarized in the table below.

Diagnostic Yield of Epilepsy Panel Genes in Selected Populations

| Epilepsy Type | Gene | Protein | Inh | Diagnostic Yield in Selected Population(s) |
|---|---|---|---|---|
| Benign familial neonatal seizures (BFNS) | <i>KCNQ2</i> | Potassium voltage-gated channel subfamily KQT member 2 | AD | >50% BFNS ⁶ |
| | <i>KCNQ3</i> | Potassium voltage-gated channel subfamily KQT member 3 | AD | ~7% BFNS ⁶ |
| Benign familial neonatal-infantile seizures (BFNIS) | <i>SCN2A</i> | Sodium channel protein type 2 subunit alpha | AD | Unknown ⁶ |
| Benign familial infantile seizures (BFIS) | <i>PRRT2</i> | Proline-rich transmembrane protein 2 | AD | 83% familial and 30% sporadic BFIS ^{14,15,16} ; 62-96% of familial PKD/IC; 36% sporadic PKD/IC ^{14,17,18} |
| Familial infantile myoclonic epilepsy (FIME) | <i>TBC1D24</i> | TBC1 domain family member 24 | AR | Unknown ^{19,20} |
| Early-onset epileptic encephalopathy and/or infantile spasms (includes West and Ohtahara syndromes) | <i>CDKL5</i> | Cyclin-dependent kinase-like 5 | XL | 10-17% infantile spasms ⁶ |
| | <i>ARX</i> | Aristaless related homeobox | XL | 5% males with infantile spasms ⁶ |
| | <i>TSC1</i> | Hamartin | AD | 2-4% infantile spasms ^{21,22,23} |
| | <i>TSC2</i> | Tuberin | AD | 10-16% infantile spasms ^{21,22,23} |
| | <i>SCN1A</i> | Sodium channel protein type 1 alpha | AD | 70-80% Dravet syndrome ⁶ ; 20-24% early-onset cryptic epilepsy ^{24,25} |
| | <i>PCDH19</i> | Protocadherin-19 | XL | 2-14% females with infantile/childhood epilepsy ^{26,27,28,29,30} |
| | <i>KCNQ2</i> | Potassium voltage-gated channel subfamily KQT member 2 | AD | 10% neonatal epileptic encephalopathy ³¹ |
| | <i>STXBP1</i> | Syntaxin binding protein 1 | AD | 35% Ohtahara syndrome ⁶ ; Unknown in Dravet syndrome ⁸⁸ |
| | <i>SLC2A1</i> | Solute carrier family 2, facilitated glucose transporter member 1 | AD | 91% GLUT1 deficiency ³ ; ~10% early-onset absence epilepsy ⁶ |
| | <i>SLC6A1</i> | Solute carrier family 6 member 1 | AD | 4% myoclonic-astatic epilepsy (MAE) ¹⁴⁹ |
| | <i>STX1B</i> | Syntaxin 1B | AD | Rare in MAE and fever-associated epilepsy syndromes ¹⁴⁹ |
| | <i>ALDH7A1</i> | Alpha-aminoacidic semialdehyde dehydrogenase (antiquitin) | AR | >90% pyridoxine-responsive epilepsy ³² |
| | <i>POLG</i> | DNA polymerase subunit gamma-1 | AR | 63-87% Alpers syndrome ^{33,34,35} ; 4-5% infantile/childhood epileptic encephalopathy ³⁴ |
| | <i>MEF2C</i> | Myocyte-specific enhancer factor 2C | AD | 2% epileptic encephalopathy ³⁶ |
| | <i>SCN2A</i> | Sodium channel protein type 2 alpha | AD | 1-2% early-onset epileptic encephalopathy ^{37,38} |
| | <i>SCN8A</i> | Sodium channel protein type 8 subunit alpha | AD | Unknown ³⁹ |
| | <i>CHD2</i> | Chromodomain helicase DNA binding protein 2 | AD | 1% of epileptic encephalopathy ⁹¹ |
| | <i>DNM1**</i> | Dynamin 1 | AD | 1% of epileptic encephalopathy ⁹² |
| | <i>GABRA1</i> | Gamma-aminobutyric acid receptor subunit alpha-1 | AD | Unknown in Dravet syndrome ⁸⁸ |
| | <i>KCNT1</i> | Potassium channel, sodium activated subfamily T, member 1 | AD | 35% MMPS ⁹⁹⁻¹⁰¹ ; Rare in other epileptic encephalopathies ⁹⁹ |
| <i>GRIN2B</i> | Glutamate receptor ionotropic, NMDA 2B | AD | Rare ⁹⁰ | |
| <i>GRIN1</i> | Glutamate receptor, ionotropic, NMDA 1 | AD | Rare ⁹⁴ | |
| <i>CASK</i> | Peripheral plasma membrane protein CASK | XL | ~96% females with MICPCH, Rare in epileptic encephalopathy ¹⁵¹ | |
| <i>SYNGAP1</i> | Ras/Rap GTPase-activating protein SynGAP | AD | Rare ¹⁵¹ | |
| <i>GABBR2</i> | Gamma-aminobutyric acid type B receptor subunit 2 | AD | Rare in epileptic encephalopathy and | |

| Epilepsy Type | Gene | Protein | Inh | Diagnostic Yield in Selected Population(s) |
|---------------|--------------------------|---|-----|---|
| | | | | intellectual disability with Rett-like features ¹¹⁸ |
| | <i>GABRB2</i> | Gamma-aminobutyric acid (GABA) A receptor, beta 2 | AD | Rare ¹¹² |
| | <i>GABRB3</i> | Gamma-aminobutyric acid (GABA) A receptor, beta 3 | AD | Rare ⁹⁵ |
| | <i>KCNA2</i> | Potassium voltage-gated channel subfamily A member 2 | AD | Rare ¹³⁸ |
| | <i>KCNB1</i> | Potassium channel, voltage gated shab related subfamily B, member 1 | AD | Rare ⁹⁸ |
| | <i>SMC1A</i> | Structural maintenance of chromosomes protein 1A | AD | Rare ^{151,152} |
| | <i>PNPO</i> | Pyridoxine-5'-phosphate oxidase | AR | Rare ⁴⁰ |
| | <i>SPTAN1</i> | Alpha-II spectrin | AD | Rare ⁴¹ |
| | <i>ALG13</i> | ALG13, UDP-N-acetylglucosaminyltransferase subunit | XL | Rare ⁹² |
| | <i>ARHGEF9</i> | Cdc42 guanine nucleotide exchange factor (GEF) 9 | XL | Rare ⁹⁴ |
| | <i>BRAT1</i> | BRCA1 associated ATM activator 1 | AR | Rare ¹²³ |
| | <i>CACNA1A</i> | Calcium channel, voltage-dependent, P/Q type, alpha 1A subunit | AD | Rare ⁹⁵ |
| | <i>CLCN4</i> | Chloride voltage-gated channel 4 | XL | Rare ¹³³ |
| | <i>EEF1A2</i> | Eukaryotic translation elongation factor 1 alpha 2 | AD | Rare ^{96, 97} |
| | <i>FRRS1L</i> | DOMON domain-containing protein FRRS1L | AR | Rare ¹³⁵ |
| | <i>GNAO1</i> | Guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O | AD | Rare in epileptic encephalopathy and early-onset hyperkinetic movement disorders ¹³⁶ |
| | <i>IQSEC2</i> | IQ motif and sec7 domain 7 | XL | Rare ⁹⁵ |
| | <i>NR2F1</i> | Nuclear receptor subfamily 2, group F, member 1 | AD | Rare ¹⁰³ |
| | <i>QARS</i> | Glutamyl-tRNA synthetase | AR | Rare ¹⁰⁴ |
| | <i>SLC13A5</i> | Solute carrier family 13(sodium-dependent citrate transporter), member 5 | AR | Rare ¹⁰⁸ |
| | <i>SZT2</i> | KICSTOR complex protein SZT2 | AR | Rare ¹⁴⁹ |
| | <i>TBL1XR1</i> | F-box-like/WD repeat-containing protein TBL1XR1 | AD | Rare ¹⁵⁰ |
| | <i>PIGA</i> | Phosphatidylinositol glycan anchor biosynthesis, class A | XL | Rare ¹⁰⁹ |
| | <i>PIGN</i> | Phosphatidylinositol glycan anchor biosynthesis, class N | AR | Rare ¹⁴⁵ |
| | <i>PIGO</i> | Phosphatidylinositol glycan anchor biosynthesis, class O | AR | Rare ¹¹⁰ |
| | <i>PIGV</i> | Phosphatidylinositol glycan anchor biosynthesis, class V | AR | Rare ¹¹¹ |
| | <i>LIAS</i> | Lipoyl synthase, mitochondrial | AR | Rare ⁴² |
| | <i>WWOX</i> | WW domain containing oxidoreductase | AR | Rare ¹⁰⁵ |
| | <i>MAGI2 (dels only)</i> | Membrane-associated guanylate kinase, WW and PDZ domain-containing protein 2 | AD | Rare ⁴³ |
| | <i>SLC25A22</i> | Mitochondrial glutamate carrier 1 | AR | Rare ⁴⁴ |
| | <i>PLCB1 (dels only)</i> | 1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase beta-1 | AR | Rare ¹⁵² |
| | <i>ATP6AP2</i> | Renin receptor | XL | Rare ⁴⁵ |
| Autosomal | <i>CHRNA4</i> | Neuronal acetylcholine receptor alpha-4 | AD | ~10% autosomal dominant nocturnal frontal lobe epilepsy (ADNFLE) ⁶ |

| Epilepsy Type | Gene | Protein | Inh | Diagnostic Yield in Selected Population(s) |
|---|----------------------|---|--------|---|
| Dominant Focal Epilepsies | <i>CHRNA2</i> | Neuronal acetylcholine receptor alpha-2 | AD | < 5% ADNLFLE ⁶ |
| | <i>CHRNA2</i> | Neuronal acetylcholine receptor alpha-2 | AD | Rare in ADNLFLE ⁶ |
| | <i>DEPDC5</i> | DEP domain containing 5 | AD | 5-37% autosomal dominant focal epilepsies, with or without cortical dysplasia ¹²⁶ |
| | <i>KCNT1</i> | Potassium channel, sodium activated subfamily T, member 1 | AD | < 5% ADNLFLE ^{99,102} |
| | <i>LGI1</i> | Leucine-rich glioma-inactivated protein 1 | AD | ~50% autosomal dominant partial epilepsy with auditory features (ADTLE) ⁶ |
| | <i>NPRL3</i> | GATOR complex protein NPRL3 | AD | ~2% autosomal dominant focal epilepsies, with or without cortical dysplasia ¹²⁷ |
| Generalized Epilepsy with Febrile Seizures Plus (GEFS+) | <i>SCN1A</i> | Sodium channel protein type 1 alpha | AD | 5-10% GEFS+ ⁵ |
| | <i>SCN1B</i> | Sodium channel subunit beta-1 | AD | <5% GEFS+ ⁶ |
| | <i>GABRG2</i> | Gamma-aminobutyric acid receptor subunit gamma-2 | AD | <1% GEFS+ ⁶ |
| | <i>SCN2A</i> | Sodium channel protein type 2 alpha | AD | Rare ⁵ |
| Epilepsy with paroxysmal dyskinesia | <i>PRRT2</i> | Proline-rich transmembrane protein 2 | AD | 62-96% of familial PKD/infantile convulsions (PKD/IC); 36% sporadic PKD/IC ^{14,17,18} |
| | <i>KCNMA1</i> | Potassium calcium-activated channel subfamily M alpha 1 | AD/A R | Rare ¹⁴⁰ |
| Progressive Myoclonic Epilepsy | <i>EPM2A</i> | Laforin | AR | 53% Lafora disease ⁴⁸ |
| | <i>NHLRC1</i> | NHL repeat-containing protein 1 (malin) | AR | 40% Lafora disease ⁴⁸ |
| | <i>CSTB*</i> (EPM1) | Cystatin-B | AR | >90% Unverricht-Lundborg disease ⁴⁹ |
| | <i>KCTD7</i> (EPM3) | BTB/POZ domain-containing protein KCTD7 | AR | Rare ⁵⁰ |
| | <i>SCARB2</i> (EPM4) | Lysosome membrane protein 2 | AR | ~7% progressive myoclonic epilepsy ⁵¹ ; Unknown in action myoclonus-renal failure syndrome ⁵² |
| | <i>GOSR2</i> | Golgi SNAP receptor complex member 2 | AR | Rare ^{53,54} |
| | <i>KCNC1</i> | Potassium voltage-gated channel subfamily C member 1 | AD | Rare KCNC1-associated myoclonus epilepsy and ataxia (MEAK) ¹¹⁵ |
| | <i>FOLR1</i> | Folate receptor alpha | AR | Rare ⁵⁵ |
| Rett/atypical Rett syndromes | <i>MECP2</i> | Methyl CpG binding protein 2 | XL | 88% females with Rett syndrome ⁵⁷ |
| | <i>CDKL5</i> | Cyclin-dependent kinase-like 5 | XL | 2-8% females with atypical Rett syndrome ^{58,59} |
| | <i>CTNNA1</i> | Catenin beta 1 | AD | Rare in Rett-like syndromes ¹¹⁶ |
| | <i>DDX3X</i> | DEAD-box helicase 3, X-linked | XL | Rare in Rett-like syndromes ¹¹⁷ |
| | <i>FOXP1**</i> | Forkhead box protein G1 | AD | ~1% Rett syndrome overall ⁶⁰ ; 25% congenital variant of Rett ⁶¹ |
| | <i>GABBR2</i> | Gamma-aminobutyric acid type B receptor subunit 2 | AD | Rare in epileptic encephalopathy and intellectual disability with Rett-like features ¹¹⁸ |
| | <i>KCNA2</i> | Potassium voltage-gated channel subfamily A member 2 | AD | Rare in Rett-like syndromes ¹¹⁹ |
| | <i>MBD5</i> | Methyl-CpG-binding domain protein 5 | AD | Rare in Rett-like syndromes ⁶² |
| | <i>MEF2C</i> | Myocyte-specific enhancer factor 2C | AD | Rare in Rett-like syndromes ⁶³ |
| | <i>SATB2</i> | SATB homeobox 2 | AD | Rare in Rett-like syndromes ¹²⁰ |
| | <i>STXBP1</i> | Syntaxin binding protein 1 | AD | Rare in Rett-like syndromes ¹²¹ |

| Epilepsy Type | Gene | Protein | Inh | Diagnostic Yield in Selected Population(s) |
|--|-----------------------|--|-----|---|
| | <i>TBL1XR1</i> | Transducin (beta)-like 1 X-linked receptor 1 | AD | Rare in Rett-like syndromes ¹²² |
| | <i>WDR45</i> | WD repeat domain 45 | XL | Rare in Rett-like syndromes ^{106, 107} |
| Angelman/ Angelman-like/ Pitt-Hopkins syndromes | <i>UBE3A</i> | Ubiquitin protein ligase E3A | AD | 68% maternally inherited 15q11.2 deletion ⁶⁴ ; 11% UBE3A sequencing variant Angelman syndrome ⁶⁴ |
| | <i>SLC9A6</i> | Sodium/hydrogen exchanger 6 | XL | 6% Angelman-like syndrome ⁶⁵ |
| | <i>MBD5</i> | Methyl-CpG-binding domain protein 5 | AD | Rare in Angelman-like syndrome ⁶⁶ |
| | <i>TCF4</i> | Transcription factor 4 | AD | 36% Pitt-Hopkins syndrome (PHS) ⁶⁷ ; 2% Angelman syndrome ⁶⁷ |
| | <i>ATRX</i> | Transcriptional regulator ATRX | XL | 95% males with alpha-thalassemia X-linked intellectual disability syndrome ¹⁵³ |
| | <i>NRXN1</i> | Neurexin-1 | AR | Rare in PHS ⁶⁸ |
| | <i>CNTNAP2</i> | Contactin-associated protein-like 2 | AR | Rare in PHS ⁶⁸ |
| Mowat-Wilson syndrome | <i>ZEB2</i> | Zinc finger E-box-binding homeobox 2 | AD | 95% Mowat Wilson syndrome ^{69,70} |
| Creatine deficiency syndromes | <i>SLC6A8</i> ** | Solute carrier family 6 (neurotransmitter transporter), member 8 | XL | 2% males with epilepsy and intellectual disability ⁴⁶ ; 65% males with biochemical creatine deficiency ⁴⁷ |
| | <i>GAMT</i> ** | Guanidinoacetate N-methyltransferase | AR | Rare ⁷¹ |
| | <i>GATM</i> | Glycine amidinotransferase, mitochondrial | AR | Rare ⁷¹ |
| Neuronal Ceroid Lipofuscinoses (NCL) | <i>PPT1 (CLN1)</i> | Palmitoyl-protein thioesterase 1 | AR | 98% PPT1 deficiency ⁷² |
| | <i>TPP1 (CLN2)</i> | Tripeptidyl-peptidase 1 | AR | 95% TPP1 deficiency ⁷³ |
| | <i>CLN3</i> | Battenin | AR | 92% Juvenile NCL ⁷⁴ |
| | <i>DNAJC5 (CLN4B)</i> | DnaJ homolog subfamily C member 5 | AD | 25% Kufs disease ^{75,76} |
| | <i>CLN5</i> | Ceroid-lipofuscinosis neuronal protein 5 | AR | 94% Finnish late-infantile NCL ⁷⁷ ; Otherwise rare |
| | <i>CLN6</i> | Ceroid-lipofuscinosis neuronal protein 6 | AR | Rare ⁷⁷ |
| | <i>MFSD8 (CLN7)</i> | Major facilitator superfamily domain-containing protein 8 | AR | Rare ⁷⁷ |
| | <i>CLN8</i> | Ceroid-lipofuscinosis neuronal protein 8 | AR | 100% Finnish Northern epilepsy ⁷⁷ ; Otherwise rare |
| | <i>CTSD (CLN10)</i> | Cathepsin D | AR | Rare ⁷⁷ |
| | <i>CTSF (CLN13)</i> | Cathepsin F | AR | Rare ⁷⁷ |
| | <i>KCTD7 (CLN14)</i> | BTB/POZ domain-containing protein KCTD7 | AR | Rare ⁷⁸ |
| Metabolic disorders | <i>ALDH5A1</i> | Aldehyde dehydrogenase 5 family member A1 | AR | 96% SSADH deficiency ¹²⁹ |
| | <i>ADSL</i> | Adenylosuccinate lyase | AR | Rare adenylosuccinate lyase deficiency ⁷⁹ |
| | <i>ASNS</i> | Asparagine synthetase (glutamine-hydrolyzing) | AR | Unknown (asparagine synthetase deficiency) ¹³⁰ |
| | <i>FOLR1</i> | Folate receptor alpha | AR | Rare cerebral folate deficiency ⁵⁶ |
| | <i>GLDC</i> | Glycine decarboxylase | AR | 70-75% glycine encephalopathy (nonketotic hyperglycinemia) ¹²⁴ |
| | <i>NGLY1</i> | N-glycanase 1 | AR | Rare in congenital disorder of deglycosylation ¹⁴³ |
| | <i>SLC19A3</i> | Solute carrier family 19 member 3 | AR | Rare thiamine metabolism dysfunction disorders ¹²⁵ |
| Periventricular | <i>FLNA</i> | Filamin A | XL | 93% classic bilateral periventricular nodular |

| Epilepsy Type | Gene | Protein | Inh | Diagnostic Yield in Selected Population(s) |
|---|--|---|---------------------|---|
| nodular heterotopia | | | | heterotopia ¹³⁴ |
| Kabuki syndrome | <i>KDM6A</i> | Lysine demethylase 6A | XL | Rare Kabuki syndrome ¹²⁸ |
| EAST/SeSAME syndrome | <i>KCNJ10</i> | ATP-sensitive inward rectifier potassium channel 10 | AR | Rare ^{81,82} |
| CLIFAHDD and infantile hypotonia with psychomotor retardation and characteristic facies (IHPRF) | <i>NALCN</i> | Sodium leak channel, non-selective | AD/AR | Rare ¹⁴² |
| Epilepsy and hemiplegic migraine | <i>ATP1A2</i> | Sodium/potassium-transporting ATPase subunit alpha-2 | AD | Unknown ⁶ |
| Epilepsy and alternating hemiplegia of childhood | <i>ATP1A3</i> | Sodium/potassium-transporting ATPase subunit alpha-3 | AD | Rare in early life epilepsy ¹³¹ , 78% alternating hemiplegia of childhood ¹³² |
| Idiopathic generalized epilepsy | <i>CHRNA7 (dels only)</i> | Neuronal acetylcholine receptor subunit alpha-7 | AD | ~1% patients with IGE ⁵⁶ |
| Epilepsy with learning, speech, and/or behavioral disorders | <i>GRIN2A</i> | Glutamate receptor ionotropic, NMDA 2A | AD | 9-20% atypical Rolandic epilepsy and epilepsy-aphasia ⁸⁹ ; ~2% patients with intellectual disability and epilepsy ^{83,84} |
| | <i>GRIN2B</i> | Glutamate receptor ionotropic, NMDA 2B | AD | Unknown ⁸³ |
| | <i>KANSL1</i> | KAT8 regulatory NSL complex subunit 1 | AD | Unknown ^{85,86} |
| | <i>DYRK1A</i> | Dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A | AD | Rare ⁹³ |
| | <i>EEF1A2</i> | Eukaryotic translation elongation factor 1 alpha 2 | AD | Rare ⁹⁶ |
| | <i>HNRNPU</i> | HNRNPU antisense RNA 1 | AD | Rare ¹³⁷ |
| | <i>KCNH1</i> | Potassium voltage-gated channel subfamily H member 1 | AD | Rare in Temple-Baraitser and Zimmermann-Laband syndromes ¹³⁹ |
| | <i>KIAA2022</i> | Neurite extension and migration factor | XL | Rare ¹⁴¹ |
| | <i>PACS1</i> | Phosphofurin acidic cluster sorting protein 1 | AD | Rare ¹⁴⁴ |
| | <i>PNKP</i> | Bifunctional polynucleotide phosphatase/kinase | AR | Rare ⁸⁶ |
| <i>PPP2R5D</i> | protein phosphatase 2 regulatory subunit B', delta | AD | Rare ¹⁴⁶ | |

* The dodecamer repeat expansion that accounts for ~90% of all CSTB mutations is not detectable by this test

** Does not include deletion/duplication testing of DNM1, FOXG1, and SLC6A8; duplications of GAMT may not be detectable

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Appendix: Genes Included on Comprehensive Epilepsy Panel and Subpanels

| Gene | Comprehensive | Infantile | Childhood | STAT | Progressive | Rett/Angelman |
|----------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| ADSL | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| ALDH5A1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| ALDH7A1 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| ALG13 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| ARHGEF9 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| ARX | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| ASNS | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| ATP1A2 | <input type="checkbox"/> | | | | | |
| ATP1A3 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| ATP6AP2 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| ATRX | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| BRAT1 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| CACNA1A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| CASK | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| CDKL5 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> |
| CHD2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| CHRNA2 | <input type="checkbox"/> | | <input type="checkbox"/> | | | |
| CHRNA4 | <input type="checkbox"/> | | <input type="checkbox"/> | | | |
| CHRNA7** | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| CHRN2 | <input type="checkbox"/> | | <input type="checkbox"/> | | | |
| CLCN4 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| CLN3 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| CLN5 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| CLN6 | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> | |
| CLN8 | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> | |
| CNTNAP2 | <input type="checkbox"/> | <input type="checkbox"/> | | | | <input type="checkbox"/> |
| CSTB*** | <input type="checkbox"/> | | | | <input type="checkbox"/> | |
| CTNNA1 | <input type="checkbox"/> | | | | | <input type="checkbox"/> |
| CTSD (CLN10) | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| CTSF | <input type="checkbox"/> | | | | <input type="checkbox"/> | |
| DDX3X | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| DEPDC5 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| DNAJC5 (CLN4B) | <input type="checkbox"/> | | | | <input type="checkbox"/> | |
| DNM1* | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| DYRK1A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| EF1A2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| EHMT1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| EPM2A | <input type="checkbox"/> | | <input type="checkbox"/> | | <input type="checkbox"/> | |
| FLNA | <input type="checkbox"/> | | | | | |
| FOLR1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | |
| FOXG1* | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| FRRS1L | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| GABBR2 | <input type="checkbox"/> | <input type="checkbox"/> | | | | <input type="checkbox"/> |
| GABRA1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GABRB2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GABRB3 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GABRG2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GAMT* | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GATM | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GLDC | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| GNAO1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GOSR2 | <input type="checkbox"/> | | <input type="checkbox"/> | | | |
| GRIN1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| GRIN2A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| GRIN2B | <input type="checkbox"/> | <input type="checkbox"/> | | | | |

| Gene | Comprehensive | Infantile | Childhood | STAT | Progressive | Rett/Angelman |
|----------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| HNRNPU | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| IQSEC2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| KANSL1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| KCNA2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| KCNB1 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| KCNC1 | <input type="checkbox"/> | | <input type="checkbox"/> | | <input type="checkbox"/> | |
| KCNH1 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| KCNJ10 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| KCNMA1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| KCNQ2 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| KCNQ3 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| KCNT1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| KCTD7 (CLN14) | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| KDM6A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| KIAA2022 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| LGII | <input type="checkbox"/> | | <input type="checkbox"/> | | | |
| MAGI2** | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| MBD5 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| MECP2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> |
| MEF2C | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> |
| MFSD8 (CLN7) | <input type="checkbox"/> | | <input type="checkbox"/> | | <input type="checkbox"/> | |
| NALCN | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| NGLY1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| NHLRC1 (EPM2B) | <input type="checkbox"/> | | <input type="checkbox"/> | | <input type="checkbox"/> | |
| NPRL3 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| NR2F1 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| NRXN1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| PACS1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| PCDH19 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| PIGA | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| PIGN | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| PIGO | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| PIGV | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| PLCB1** | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| PNKP | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| PNPO | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| POLG | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| PPP2R5D | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| PPT1 (CLN1) | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | |
| PRRT2 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| PURA | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| QARS | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| SATB2 | <input type="checkbox"/> | | | | | <input type="checkbox"/> |
| SCARB2 | <input type="checkbox"/> | | | | <input type="checkbox"/> | |
| SCN1A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SCN1B | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SCN2A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SCN8A | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| SLC13A5 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| SLC19A3 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SLC25A22 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| SLC2A1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SLC6A1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| SLC6A8* | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | |
| SLC9A6 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| SMC1A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| SPATA5 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |

| Gene | Comprehensive | Infantile | Childhood | STAT | Progressive | Rett/Angelman |
|-------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| SPTAN1 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| STX1B | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| STXBP1 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | <input type="checkbox"/> |
| SYNGAP1 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| SZT2 | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| TBC1D24 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | |
| TBL1XR1 | <input type="checkbox"/> | <input type="checkbox"/> | | | | <input type="checkbox"/> |
| TCF4 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| TPP1 (CLN2) | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | |
| TSC1 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| TSC2 | <input type="checkbox"/> | <input type="checkbox"/> | | <input type="checkbox"/> | | |
| UBE3A | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| WDR45 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |
| WWOX | <input type="checkbox"/> | <input type="checkbox"/> | | | | |
| ZEB2 | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | | | <input type="checkbox"/> |

* Sequencing only (no deletion/duplication analysis) for DNMI, FOXP1, or SLC6A8. This test can detect only complete gene deletions/duplications of GAMT. No sequence coverage of exon 22 of DNMI (NM_004408.3).

** Deletion/duplication testing only (no sequence analysis) for CHRNA7, MAGI2, and PLCB1.

*** The dodecamer repeat expansion that accounts for ~90% of all CSTB mutations is not be detectable by this test.