Glucose Transporter Type 1 Deficiency Syndrome: SLC2A1 Gene Sequencing

Test Code: SSLC2
Turnaround time: 2 weeks
CPT Codes: 81479 x1

Condition Description

Glucose transporter type 1 deficiency syndrome (Glut-1-DS) is characterized by infantile seizures that are resistant to treatment by anticonvulsants. This is followed by delays in mental and motor development, ataxia, dysarthria, spasticity, and uncontrolled eye movement before meals. The most important laboratory observation in Glut-1-DS is a reduced cerebrospinal fluid glucose concentration. Affected infants appear normal at birth. Seizures typically begin between one and four months of age. The frequency, type, and severity of seizures vary among individuals with Glut-1-DS. Additionally, cognitive impairment can range from learning difficulties to severe intellectual disability.

Mutation of the SLC2A1 (1p35-p31.3) gene cause Glut-1-DS. Glut-1-DS is inherited in an autosomal dominant manner with the majority of mutations occurring de novo. Affected parents have been reported with a mild or subclinical degree of impairment.

For patients with suspected Glut-1-DS, sequence analysis is recommended as the first step in mutation identification. For patients in whom mutations are not identified by full gene sequencing, deletion/duplication analysis is appropriate.

References:
- GeneReviews
- OMIM #138140: SLC2A1 gene
- OMIM #606777: Glut-1-DS

Genes
SLC2A1

Indications

This test is indicated for:
- Confirmation of a clinical diagnosis of Glucose Transporter Type 1 Deficiency syndrome.
- Carrier testing in adults with a family history of Glucose Transporter Type 1 Deficiency syndrome.

Methodology

PCR amplification of 10 exons contained in the SLC2A1 gene is performed on the patient's genomic DNA. Direct sequencing of amplification products is performed in both forward and reverse directions, using automated fluorescence dideoxy sequencing methods. The patient's gene sequences are then compared to a normal reference sequence. Sequence variations are classified as mutations, benign variants unrelated to disease, or variations of unknown clinical significance. Variants of unknown clinical significance may require further studies of the patient and/or family members. This assay does not interrogate the promoter region, deep intronic regions, or other regulatory elements, and does not detect large deletions.

Detection

Sequencing analysis detects 91% of mutations in individuals with Glut-1-DS. Mutations in the promoter region, some mutations in the introns and other regulatory element mutations cannot be detected by this analysis. Large deletions will not be detected by this analysis. Results of molecular analysis should be interpreted in the context of the patient's clinical and/or biochemical phenotype.

Analytical Sensitivity: ~99%

Specimen Requirements

Submit only 1 of the following specimen types
- Preferred specimen type: Whole Blood

Type: Whole Blood

Specimen Requirements:

In EDTA (purple top) or ACD (yellow top) tube:
- Infants (2 years): 3-5 ml
- Older Children & Adults: 5-10 ml

Specimen Collection and Shipping: Refrigerate until time of shipment. Ship sample within 5 days of collection at room temperature with overnight delivery.

Type: Saliva
Specimen Requirements:

Oragene™ Saliva Collection kit (available through EGL) used according to manufacturer instructions.

Specimen Collection and Shipping: Store sample at room temperature. Ship sample within 5 days of collection at room temperature with overnight delivery.

Related Tests

- Deletion/duplication analysis of the SLC2A1 gene by CGH array is available for those individuals in whom sequence analysis is negative.
- Custom diagnostic mutation analysis (KM) is available to family members if mutations are identified by targeted mutation testing or sequencing analysis.
- Prenatal testing is available only for known familial mutations to individuals who are confirmed carriers of mutations. Please contact the laboratory genetic counselor to discuss appropriate testing prior to collecting a prenatal specimen.