Peroxisome Biogenesis Disorders, Zellweger Syndrome Spectrum,
PEX12-related:

PEX12 Gene

Test Code: SPEXX

Turnaround time: 4 weeks

CPT Codes: 81479 x1

Condition Description

Peroxisome biogenesis disorders, Zellweger syndrome spectrum (PBD, ZSS) is a continuum that encompasses three distinct phenotypes; Zellweger syndrome (ZS), neonatal adrenoleukodystrophy (NALD), and infantile Refsum disease (IRD). Individuals with PBD, ZSS usually present during the newborn period or later in childhood. Features during the newborn period include hypotonia, poor feeding, distinctive facial features, seizures, and liver cysts with hepatic dysfunction. Infants with ZS, the most severe phenotype on the spectrum, are significantly impaired and usually do not survive past the first year of life. Those that do survive past the first year have developmental delay, hypotonia, liver dysfunction, sensorineural hearing loss, and retinal dystrophy. Features of NALD and IRD (the least severe phenotype on the spectrum) are variable. They include developmental delays, liver dysfunction, episodes of intracranial bleeding, hearing loss, and vision impairments. PBD, ZSS can be slowly progressive.

PBD, ZSS is inherited in an autosomal recessive manner. Biochemical assays can definitively diagnose PBD, ZSS. Mutations in twelve different PEX genes have been identified to cause PBD, ZSS. They are the PEX1, PEX2, PEX3, PEX5, PEX6, PEX10, PEX12, PEX13, PEX14, PEX16, PEX19, and PEX26 genes. About 68% of individuals with PBD, ZSS have mutations in the PEX1 gene. Mutations in the PEX6, PEX10, PEX12, and PEX26 genes account for an additional 26% of mutations in individuals with PBD, ZSS.

Please note that this test is for the PEX12 gene (17q12) only.

For patients with suspected PBD, ZSS, 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 #601758: PEX12 gene
OMIM #214100: ZS

Indications

This test is indicated for:

- Confirmation of a clinical diagnosis of peroxisome biogenesis disorders, Zellweger syndrome spectrum.
- Carrier testing in adults with a family history of peroxisome biogenesis disorders, Zellweger syndrome spectrum.

Methodology

PCR amplification of 3 exons contained in the PEX12 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

Clinical Sensitivity: Mutations in the PEX12 gene account for ~4% of individuals with PBD, ZSS. Mutations in the PEX1 gene are the most common cause of PBD, ZSS.

Analytical Sensitivity: ~99%

Specimen Requirements

- Type: Whole Blood
- Preferred specimen type: Whole Blood

Specimen Requirements

- Anticoagulant: EDTA (purple top)
- Minimum volume: 5 ml whole blood
- Specimen stability: 7 days at room temperature, 14 days at 2-8°C, and 6 months at -20°C
- Collection equipment: EDTA (purple top) tube

Specimen Handling

- Whole blood is stable for 48 hours at room temperature. No preservative is necessary if whole blood is stored at room temperature for less than 48 hours. Whole blood should be refrigerated for longer storage.

Specimen Collection

- Whole blood is collected in EDTA (purple top) tubes.

Specimen Transport

- Whole blood is stable for 48 hours at room temperature. No preservative is necessary if whole blood is stored at room temperature for less than 48 hours. Whole blood should be refrigerated for longer storage.

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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.

**Special Instructions**

Submit copies of diagnostic biochemical test results with the sample, if appropriate. Contact the laboratory if further information is needed.

**Related Tests**

- Deletion/duplication analysis of the *PEX12* gene by CGH array is available for those individuals in whom sequence analysis is negative.
- Sequence and deletion/duplication analysis for the *PEX1*, *PEX2*, *PEX3*, *PEX5*, *PEX6*, *PEX14*, and *PEX26* genes are also available.
- 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.