Neuro17
Hereditary Cancer Gene Panel

**including:** sequencing and high resolution deletion/duplication analysis

**PANEL DESCRIPTION:**
Using sequencing and high resolution deletion/duplication analysis, the Hereditary Neuro17 Cancer Panel analyzes 17 genes for genetic variants that confer an increased risk of developing a brain tumor and, in some cases, other cancers and noncancerous conditions. Identifying a causative genetic variant provides valuable information for the patient’s diagnosis, medical management, surveillance (early screening for other associated cancers), and familial screening.

**TEST DETAILS:**
- This panel includes both sequencing and high resolution deletion/duplication analysis of the genes specified when performed on a blood specimen.
  - **Sequencing** is performed using a customized next generation sequencing library. Analysis includes the coding exons of all genes in the panel plus ten bases into the introns and untranslated regions (5’ and 3’). Sanger sequencing is performed to confirm variants classified as pathogenic or suspected pathogenic.
  - **Deletion/duplication analysis** is performed using a high resolution, custom microarray platform designed to target the genes of interest at the exon level.
    - In addition, deletion/duplication studies of PMS2 will be performed using Multiplex Ligation-dependent Probe Amplification (MLPA) because of the presence of highly homologous pseudogenes. However, MLPA cannot be performed on a buccal specimen.
- Detection rates are limited to the genes specified; this test does not provide whole genome analysis.
- Gene panels are a more cost-effective approach than single gene testing to confirm or establish a diagnosis. However, if single gene testing is desired for the patient or family members of an individual with a known mutation, that must be ordered separately.

**INDICATIONS FOR TESTING:**
- Personal and/or family history of
  - Brain tumor(s)
  - Central nervous system (CNS) tumor(s)
  - Peripheral nervous system (PNS) tumor(s)
- Brain or CNS tumor associated with multiple genetic conditions, such as ependymoma, glioblastoma, meningioma, and medulloblastoma
- Personal and/or family history raising concern for the syndromes included in this panel
  - Nevoid basal cell carcinoma (Gorlin) syndrome GeneReviews® [http://www.ncbi.nlm.nih.gov/books/NBK1151/]
  - Li-Fraumeni syndrome GeneReviews® [http://www.ncbi.nlm.nih.gov/books/NBK1311/]
  - Neurofibromatosis type I (NF1) GeneReviews® [http://www.ncbi.nlm.nih.gov/books/NBK1109/]
  - PTEN hamartoma tumor (Cowden) syndrome GeneReviews® [http://www.ncbi.nlm.nih.gov/books/NBK1488/]
- Known familial variant in any of the genes included in this panel (targeted analysis)
SPECIMEN COLLECTION & TRANSPORT:
Complimentary test kits are available upon request, but are not required.

SAMPLE TYPE and REQUIREMENTS:
- blood: 3 - 5 ml whole blood in an EDTA tube (purple top)
- buccal mucosa swab: 2 swabs^  
- extracted DNA (from blood or buccal): 70 ng/µg with a total yield of 7-10 µg (in TE) in a DNA microcentrifuge tube

^ Deletion/duplication and rearrangement analyses cannot be performed for indicated genes on a buccal specimen.

SHIPPING:
- Maintain and ship samples at room temperature.
- Coordinate transport for sample to be received in our laboratory within 24-48 hours of collection.
  - LOCAL: Call 402-559-5070 (option 1)
  - OUT OF AREA: Prior to shipment, please fax the completed test request form to 402-559-7248, including the FedEx® airbill tracking number.
    - Saturday delivery MUST be checked when sending FedEx® on Friday.
    - Please include Internal Billing Reference # 3155070600 on the FedEx® airbill.
    - Ship To: Human Genetics Laboratory – Zip 5440
      UNMC Shipping & Receiving Dock
      601 S. Saddle Creek Road
      Omaha, NE 68106

FORMS FOR TESTING: The following forms can be downloaded via our website.
- Required: Hereditary Cancer Test Request Form
- Optional: Informed Consent for Hereditary Cancer Genetic Testing

POTENTIAL TEST RESULTS:
- Positive for a Pathogenic Variant – This designation is reported when a pathogenic change is identified in a gene and the change is known to increase the risk for certain cancers.
- Negative – This designation is reported when no harmful or uncertain changes are found in any of the genes tested.
- A Variant of Uncertain Clinical Significance (UCS) – This designation is reported when an inconclusive change in a gene is found; the change may increase the risk for cancer or be harmless. Variants of uncertain clinical significance may also be classified as “Likely Pathogenic” or “Likely Benign” based on the ACMG recommendations for variant classification.1

TURNAROUND TIME: For all sample types, results are typically available in 14 days.

BILLING: Our laboratory offers patient/self-pay, insurance (including Medicare/Medicaid), and client/institution billing options. Verifying coverage requirements or obtaining preauthorization PRIOR TO OR AT THE TIME OF SPECIMEN COLLECTION is often necessary. We provide preauthorization services upon request by calling 402-559-5070 (option 3); the following form is helpful for obtaining the information required by insurance providers and can be downloaded via our website.
- Insurance Preauthorization Request

In some circumstances, a test may be warranted even though insurance coverage is denied or not guaranteed. For these situations, we request the following form be signed by the patient and submitted with the sample. This helps inform patients of their potential financial responsibility, should the costs of genetic testing not be paid by their insurance provider.
- Advanced Beneficiary Notice of Noncoverage (ABN) – required when billing Medicare
CPT CODES: 81405, 81407

PRICING: For current costs contact the laboratory billing staff at 402-559-5070 (option 3).

GENE LIST:

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<thead>
<tr>
<th>ALK</th>
<th>MLH1</th>
<th>NBN</th>
<th>PHOX2B</th>
<th>SUFU</th>
<th>TSC2</th>
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<tr>
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<td>MSH2</td>
<td>NF1</td>
<td>PMS2^</td>
<td>TP53</td>
<td>VHL</td>
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<tr>
<td>MEN1</td>
<td>MSH6</td>
<td>NF2</td>
<td>PTCH1</td>
<td>TSC1</td>
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</tr>
</tbody>
</table>

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RESOURCES: The following resource can be downloaded via our website, or you may request brochures for your clinic by contacting our marketing specialist at 402-559-6935 | humangenetics@unmc.edu.

- PATIENT BROCHURE: Neuro17

REFERENCES:

1 Richards et al. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology. Genet Med advance online publication 5 March 2015; 1-20.

updated 3-12-2019