Juvenile Polyposis Syndrome: **SMAD4** Gene Deletion/Duplication

**Test Code:** UU  
**Turnaround time:** 4 weeks  
**CPT Codes:** 81405 x1

### Condition Description

Juvenile polyposis syndrome (JPS) is an autosomal dominant condition characterized by predisposition to hamartomatous polyps in the gastrointestinal (GI) tract, specifically in the stomach, small intestine, colon, and rectum. The term “juvenile” refers to the type of polyp rather than to the age of onset of polyps. Most individuals with JPS have some polyps by age 20 years; some may have only four or five polyps over their lifetime, whereas others in the same family may have more than a hundred. If the polyps are left untreated, they may cause bleeding and anemia. Most juvenile polyps are benign; however, malignant transformation can occur. Most of this increased risk is attributed to colon cancer, but cancers of the stomach, upper GI tract, and pancreas have been reported. The incidence of colorectal cancer is 17%-22% by age 35 years; some individuals may have only a few polyps over their lifetime, whereas others in the same family may have more than a hundred. The incidence of gastric cancer is 21% in those with juvenile polyps.

JPS is clinically diagnosed if any one of the three following findings is present: more than five juvenile polyps of the colorectum; multiple juvenile polyps throughout the GI tract; any number of juvenile polyps and a family history of juvenile polyps. Juvenile polyps are hamartomas with a distinct histology that differs from that of adenomas. The genes known to be associated with JPS are **SMAD4** and **BMPR1A**. Approximately 20% of individuals with JPS have mutations in **SMAD4** (18q21.1); approximately 20% have mutations in **BMPR1A**. Recent studies suggest that deletion/duplication testing can identify an additional 9%-14% of mutations in **SMAD4**.

A combined syndrome of JPS and hereditary hemorrhagictelangiectasia (HHT) (termed JPS/HHT) may be present in 15%-22% of individuals with a **SMAD4** mutation. Some clinicians suggest that patients with juvenile polyposis who have a **SMAD4** mutation should be screened for the vascular lesions associated with hereditary hemorrhagictelangiectasia, especially occult arteriovenous malformations in visceral organs that may otherwise present suddenly with serious medical consequences.

Click here for the GeneTests summary on this condition.

### Genes

**SMAD4**

### Indications

This test is indicated for:

- Confirmation of a clinical diagnosis of juvenile polyposis syndrome in individuals who have tested negative for sequence analysis
- Individuals at-risk for juvenile polyposis syndrome due to family history who have tested negative for sequence analysis

### Methodology

DNA isolated from peripheral blood is hybridized to a CGH array to detect deletions and duplications. The targeted CGH array has overlapping probes which cover the entire genomic region.

### Detection

Detection is limited to duplications and deletions. The CGH array will not detect point or intronic mutations. Results of molecular analysis must be interpreted in the context of the patient's clinical and/or biochemical phenotype.

### Specimen Requirements

Submit only 1 of the following specimen types

**Type: Whole Blood (EDTA)**

**Specimen Requirements:**
- EDTA (Purple Top)
- Infants and Young Children (2 years of age to 10 years old: 3-5 ml)
- Older Children & Adults: 5-10 ml
- Autopsy: 2-3 ml unclotted cord or cardiac blood

**Specimen Collection and Shipping:**
Ship sample at room temperature for receipt at EGL within 24 hours of collection. Do not refrigerate or freeze.

**Type: DNA, Isolated**

**Specimen Requirements:**
- Microtainer
- 3µg
- Isolation using the Perkin Elmer™Chemagen™ Chemagen™ Automated Extraction method or Qiagen™ Puregene kit for DNA extraction is recommended.
**Specimen Collection and Shipping:**
Refrigerate until time of shipment in 100 ng/µL in TE buffer. Ship sample 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.
Sequence analysis is required before deletion/duplication analysis by targeted CGH array. If sequencing is performed outside of EGL Genetics, please submit a copy of the sequencing report with the test requisition.

**Related Tests**
- Sequencing analysis of the SMAD4 gene is available (UT) and is required before deletion/duplication analysis.
- Prenatal testing is available to individuals who are confirmed carriers of mutations. Please contact the laboratory genetic counselor to discuss appropriate testing prior to collecting a prenatal specimen.