



## Molecular Genetics Laboratory

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Additional information and test requisitions are available at:  
[www.cincinnatichildrens.org/molecular-genetics/](http://www.cincinnatichildrens.org/molecular-genetics/)

<b>TEST:</b>	<b><i>JAG1</i> sequence analysis for Alagille Syndrome</b>
<b>DISORDER:</b>	Alagille syndrome is a complex disorder characterized by cholestasis and bile duct paucity, cardiac defects, skeletal abnormalities, ophthalmologic abnormalities and characteristic facial features. Less common features of Alagille syndrome include kidney abnormalities, neurovascular accidents and pancreatic insufficiency. Developmental delay and mild mental retardation are seen in a minority of individuals with Alagille syndrome. Mutations in <i>JAG1</i> result in broad phenotypic variability, even within families.
<b>INDICATIONS:</b>	<ul style="list-style-type: none"><li>▪ Diagnostic testing in patients with suspected Alagille syndrome</li><li>▪ Targeted mutation analysis of at-risk relatives of patient with <i>JAG1</i> mutation</li><li>▪ Prenatal diagnosis of an at-risk fetus</li></ul>
<b>SPECIMEN:</b>	At least 3 mls of whole blood in purple top (EDTA) tube. Label tube with patient's name, birth date, and date of collection. Phlebotomist must initial tube to verify patient's identity.
<b>TESTING METHODOLOGY:</b>	PCR-based sequencing of the coding regions and their exon/intron boundaries of the <i>JAG1</i> gene.
<b>CLINICAL SENSITIVITY:</b>	PCR-based sequencing of the coding regions and their exon/intron boundaries of the <i>JAG1</i> gene detects ~ 88% patients with Alagille syndrome. 20p12 microdeletion syndrome, which includes deletion of the entire <i>JAG1</i> gene, is identified in approximately 7% of patients with Alagille syndrome and can be detected by fluorescence in-situ hybridization (FISH) analysis. Mutations in <i>NOTCH2</i> are associated with Alagille syndrome in approximately 1% of affected individuals.
<b>ANALYTICAL SENSITIVITY:</b>	The sensitivity of PCR-based DNA sequencing is over 99% for the detection of nucleotide base changes, small deletions and insertions in the regions analyzed. Mutations in regulatory regions or other untranslated regions are not detected by this test. Multiple exon deletions, large insertions, genetic recombinational events and rare, primer site mutations may not be identified using this methodology.
<b>COSTS:</b>	Please call 1-866-450-4198 for current costs, any billing questions, or for assistance with insurance preauthorization.
<b>CPT CODES:</b>	<ul style="list-style-type: none"><li>▪ <i>JAG1</i> sequence analysis: 83890, 83898(x3), 83894(x3), 83891(x2), 83904(x49), 83909(x49), 83912</li><li>▪ Family specific analysis: 83890, 83898, 83894, 83891, 83904, 83912</li></ul>
<b>SHIPPING INSTRUCTIONS:</b>	Please enclose <b>test requisition</b> with sample. <b>All information must be completed before sample can be processed.</b> Place samples in Styrofoam mailer and ship at room temperature by overnight Federal Express to arrive Monday through Friday. <b>Ship to:</b> Cytogenetics and Molecular Genetics Laboratories 3333 Burnet Avenue NRB 1042 Cincinnati, OH 45229 513-636-4474
<b>RESULTS:</b>	Each test report includes a detailed interpretation of the genetic findings, the clinical significance of the result, and recommendations for clinical management and additional testing, if warranted. Results will be reported to the referring physician or health care provider as specified on the test requisition form.