

Client: Example Client ABC123  
123 Test Drive  
Salt Lake City, UT 84108  
UNITED STATES

Physician: Doctor, Example

**Patient: Patient, Example**

**DOB:** 1/14/2021  
**Gender:** Female  
**Patient Identifiers:** 01234567890ABCD, 012345  
**Visit Number (FIN):** 01234567890ABCD  
**Collection Date:** 00/00/0000 00:00

**Tuberous Sclerosis Complex Panel, Sequencing and Deletion/Duplication**

ARUP test code 3002100

Tuberous Sclerosis Specimen

whole blood

Tuberous Sclerosis Interp

Negative

INDICATION FOR TESTING  
Patient with cardiac rhabdomyoma.

RESULT  
No pathogenic variants were detected in any of the genes tested.

INTERPRETATION  
No pathogenic variants were identified in the TSC1 and TSC2 genes by massively parallel sequencing of the coding regions and exon-intron boundaries or by deletion/duplication analysis. This result decreases the likelihood of, but does not exclude, a diagnosis of tuberous sclerosis complex (TSC). Please refer to the background information included in this report for the limitations of this test.

RECOMMENDATIONS  
Medical screening and management of this individual should rely on clinical findings and family history. Genetic consultation is recommended.

COMMENTS  
Likely benign and benign variants are not included in this report.

This result has been reviewed and approved by [REDACTED]

H=High, L=Low, \*=Abnormal, C=Critical

**BACKGROUND INFORMATION:** Tuberous Sclerosis Complex Panel, Sequencing and Deletion/Duplication

**CHARACTERISTICS:** Tuberous sclerosis complex (TSC) is a multisystem, genetic disorder causing numerous benign tumors, as well as intellectual and developmental disabilities. Tumors can occur in the skin, brain, kidneys, and other organs, and can lead to significant health complications and may be life threatening.

**PREVALENCE:** 1 in 6,000 individuals

**CAUSE:** Pathogenic germline variants in TSC1 and TSC2

**INHERITANCE:** Autosomal dominant; approximately 66 percent are de novo

**PENETRANCE:** Complete penetrance with variable expressivity

**CLINICAL SENSITIVITY:** 95 percent

**GENES TESTED:** TSC1, TSC2

**METHODOLOGY:** Targeted capture of all coding exons and exon-intron junctions of the targeted genes, followed by massively parallel sequencing. Sanger sequencing was performed as necessary to fill in regions of low coverage and confirm reported variants. A custom tiled comparative genomic hybridization array (aCGH) was used to detect large deletions or duplications in the targeted genes. Human genome build 19 (Hg 19) was used for data analysis.

**ANALYTICAL SENSITIVITY/SPECIFICITY:** The analytical sensitivity of this test is approximately 99 percent for single nucleotide variants (SNVs) and greater than 93 percent for insertions / duplications / deletions from 1-10 base pairs in size. Variants greater than 10 base pairs may be detected, but the analytical sensitivity may be reduced.

**LIMITATIONS:** A negative result does not exclude a diagnosis of TSC. This test only detects variants within the coding regions and intron-exon boundaries of the targeted genes. Regulatory region variants and deep intronic variants will not be identified and breakpoints of large deletions / duplications will not be determined. Single exon deletions / duplications or deletions / duplications less than 1 kb may not be detected. Deletions / duplications / insertions of any size may not be detected by massive parallel sequencing. Diagnostic errors can occur due to rare sequence variations. In some cases, variants may not be identified due to technical limitations in the presence of pseudogenes, repetitive, or homologous regions. This assay may not detect low-level mosaic or somatic variants associated with disease. Interpretation of this test result may be impacted if this patient has had an allogeneic stem cell transplantation. Non-coding transcripts were not analyzed. Single exon deletions / duplications will not be called for the following exons: TSC2 (NM\_000548) 17,29,41

This test was developed and its performance characteristics determined by ARUP Laboratories. It has not been cleared or approved by the US Food and Drug Administration. This test was performed in a CLIA certified laboratory and is intended for clinical purposes.

Counseling and informed consent are recommended for genetic testing. Consent forms are available online.

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VERIFIED/REPORTED DATES				
Procedure	Accession	Collected	Received	Verified/Reported
Tuberous Sclerosis Specimen	21-015-122320	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
Tuberous Sclerosis Interp	21-015-122320	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00

END OF CHART

H=High, L=Low, \*=Abnormal, C=Critical

Unless otherwise indicated, testing performed at:

ARUP LABORATORIES | 800-522-2787 | aruplab.com  
500 Chipeta Way, Salt Lake City, UT 84108-1221  
Tracy I. George, MD, Laboratory Director

Patient: Patient, Example  
ARUP Accession: 21-015-122320  
Patient Identifiers: 01234567890ABCD, 012345  
Visit Number (FIN): 01234567890ABCD  
Page 3 of 3 | Printed: 2/23/2021 8:23:53 AM  
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