

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

Physician: Doctor, Example

**Patient: Patient, Example**

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

**Acute Myeloid Leukemia Panel by FISH**

ARUP test code 3016654

FISH AML Panel	Normal	(Ref Interval: Normal)
	INTERPRETIVE INFORMATION: AML Panel by FISH	
	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.	

EER AML Panel by FISH	EERUnavailable	
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**PML-RARA Translocation by FISH**

ARUP test code 2002363

PML-RARA Translocation by FISH	Normal	(Ref Interval: Normal)
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H=High, L=Low, \*=Abnormal, C=Critical

Test Performed: Acute Myeloid Leukemia Panel by FISH (FISHAML)  
Specimen Type: Bone Marrow  
Indication for Testing: Concern for AML

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**RESULT**  
Normal FISH Result

inv(3) or t(3;3) GATA2::MECOM Fusion: not detected  
Deletion 5q: not detected  
Monosomy 7: not detected  
Deletion 7q: not detected  
t(8;21) RUNX1::RUNX1T1 Fusion: not detected  
11p15 (NUP98) Rearrangement: not detected  
11q23 (KMT2A) Rearrangement: not detected  
inv(16) or t(16;16) CBFβ::MYH11 Fusion: not detected

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**INTERPRETATION**  
There was no evidence of GATA2::MECOM (also known as RPN1-EVI1) fusion due to 3q21/3q26.2 inversion or translocation, deletion 5q31, monosomy 7, deletion 7q31, RUNX1::RUNX1T1 fusion due to translocation (8;21)(q21.3;q22), 11p15 (NUP98) rearrangement, 11q23 KMT2A (MLL) rearrangement, or CBFβ::MYH11 fusion due to either 16p13.1/16q22 inversion or translocation.

This analysis was performed with the AML panel probes RPN1/MECOM, D5S23/EGR1, D7Z1/D7S486, RUNX1/RUNX1T1 (Abbott Molecular), NUP98 and CBFβ-MYH11 (MetaSystems), and MLL (KMT2A) (CytoCell). A total of 200 cells were scored for each probe.

Cytogenomic Nomenclature (ISCN):  
nuc  
ish(RPN1,MECOM,D5S23,EGR1,D7Z1,D7S486,RUNX1T1,NUP98,KMT2A,MYH11,CBFB,RUNX1)x2[200]

This result has been reviewed and approved by [REDACTED]

**INTERPRETIVE INFORMATION: PML/RARA Translocation by FISH**

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EER PML-RARA Translocation by FISH

EERUnavailable

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

Unless otherwise indicated, testing performed at:

VERIFIED/REPORTED DATES				
Procedure	Accession	Collected	Received	Verified/Reported
FISH AML Panel	24-166-106097	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
PML-RARA Translocation by FISH	24-166-106097	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
EER PML-RARA Translocation by FISH	24-166-106097	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
EER AML Panel by FISH	24-166-106097	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: