

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

Physician: Doctor, Example

**Patient: Patient, Example**

**DOB:** 4/3/1982  
**Gender:** Female  
**Patient Identifiers:** 01234567890ABCD, 012345  
**Visit Number (FIN):** 01234567890ABCD  
**Collection Date:** 01/01/2017 12:34

**Non-Invasive Prenatal Testing for Fetal Aneuploidy**

ARUP test code 2007537

**Result Summary**

**HIGH RISK T21**

This pregnancy is classified as HIGH RISK for trisomy 21 (Down syndrome) by this screen. This result should be confirmed by a diagnostic test. On average, 91% of pregnancies classified as HIGH RISK are found to have trisomy 21 based on a published study of 17,885 women (PMID 25111587).

**TEST INFORMATION:** Non-Invasive Prenatal Testing for Fetal Aneuploidy (Powered by Constellation) with or without Microdeletions

**METHODOLOGY:** DNA isolated from the maternal blood, which contains placental DNA, is amplified at 13,300+ loci using a targeted PCR assay and sequenced using a high-throughput sequencer. Sequence data are analyzed using Natera's Constellation software to estimate the fetal copy number and identify whole chromosome abnormalities for chromosomes 13, 18, 21, X, and Y as well as fetal sex. Barring QC failures and fetal fractions below the performance limits of the algorithm, the minimum confidence threshold is 0.98 for a high risk call. For both low risk and high risk calls, the majority of specimens will have a confidence of >0.99 across all regions tested. If a sample fails to meet the quality threshold, no result will be reported for one or more chromosomes.

**Microdeletions:** An additional 6,600+ loci are amplified to estimate the fetal copy numbers of chromosomal regions attributed to 22q11.2, Prader-willi, Angelman, Cri-du-chat, and 1p36 deletion syndromes. For any of the microdeletions, the minimum confidence threshold is 0.95 for a High Risk call. Under specific circumstances, the algorithm may return a result of "unchanged" which is equivalent to the population frequency of that microdeletion.

**SENSITIVITY AND SPECIFICITY:** For combined autosomal aneuploidies and Turner's syndrome, sensitivity and specificity are >99 percent. For combined microdeletions, sensitivity is >99 percent, specificity is >94 percent. Fetal sex has a sensitivity and specificity of >99 percent. Sex chromosome trisomies, if identified, will be reported with a specificity of 98 percent.

**DISCLAIMER:** Risks for aneuploidy are calculated based on maternal age, gestational age and test results. Risks for microdeletions are based on population frequency and test results. This test will not identify all deletions associated with each disorder. Ability to detect deletions will be based on size and location. Findings of unknown significance will not be reported. Cases with evidence of fetal and/or placental mosaicism will not be reported. As this assay is a screening test and not diagnostic, false positive and false negatives can

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

occur. Positive test results need diagnostic confirmation by alternative testing methods. Negative results do not fully exclude the diagnosis of any of the above syndromes. False positive and false negative results may be due to placental, fetal or maternal mosaicism, small imbalances, point mutations, gene inactivation, haploblocks, or other genetic/epigenetic mechanisms. Other potential sources of error include, but are not limited to, DNA sample contamination or degradation, limitations of current diagnostic techniques, misidentification of samples, or other factors that may interfere with correct interpretation of the analysis. This test has the potential to uncover consanguinity in the family. This test is not intended to identify pregnancies at risk for open neural tube defects. This test was developed and its performance characteristics determined by ARUP Laboratories. The U.S. Food and Drug Administration has not approved or cleared this test; however, FDA clearance or approval is not currently required for clinical use. The results are not intended to be used as the sole means for clinical diagnosis or patient management decisions.

SOFTWARE VERSION: Constellation 2.2

ABBREVIATIONS: T21 (trisomy 21); T18 (trisomy 18); T13 (trisomy 13); Trip (triploidy or vanishing twin); 45,X (monosomy X); XXX (trisomy X); XXY (Klinefelter syndrome); XYY (XYY syndrome); 22q (22q11.2 deletion syndrome); 1p36 (1p36 deletion syndrome); AS (Angelman syndrome); 5p- (Cri-du-chat syndrome); PWS (Prader-Willi syndrome)

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Trisomy 21

HIGH RISK

This is a screening test, and is NOT diagnostic for the conditions listed in this report. Both false positive and false negative results may occur. Genetic counseling and confirmatory fetal diagnostic testing is recommended. Irrevocable action such as pregnancy termination should not be taken based on the results of this test alone.

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Trisomy 18

Low risk

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Trisomy 13

Low risk

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Monosomy X

Low risk

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Triploidy/Vanishing Twin

Low risk

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Fetal Fraction

8.0

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Report Fetal Sex?

Yes

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Fetal Sex

Female

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Gestational Age at draw (Weeks) 20

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Gestational Age at draw (Days) 0

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Maternal Weight (Pounds) 150

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EER Non-Invasive Prenatal, Aneuploidy

**See Note**

Access ARUP Enhanced Report using either link below:

-Direct access:

-Enter Username, Password: <https://erpt.aruplab.com>

Username:

Password:

This result has been reviewed and approved by Yuan Ji, Ph.D.  
Electronic Signature

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| VERIFIED/REPORTED DATES               |               |                      |                      |                       |
|---------------------------------------|---------------|----------------------|----------------------|-----------------------|
| Procedure                             | Accession     | Collected            | Received             | Verified/Reported     |
| Result Summary                        | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Trisomy 21                            | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Trisomy 18                            | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Trisomy 13                            | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Monosomy X                            | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Triploidy/Vanishing Twin              | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Fetal Fraction                        | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Report Fetal Sex?                     | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 8:54:00 AM  |
| Fetal Sex                             | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |
| Gestational Age at draw (Weeks)       | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 8:54:00 AM  |
| Gestational Age at draw (Days)        | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 8:54:00 AM  |
| Maternal Weight (Pounds)              | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 8:54:00 AM  |
| EER Non-Invasive Prenatal, Aneuploidy | 18-030-101887 | 1/30/2018 8:14:00 AM | 1/30/2018 8:19:46 AM | 1/30/2018 10:17:00 AM |

END OF CHART

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