

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

Physician: Doctor, Example

Patient: Patient, Example

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

RhE/e (RHCE) Antigen Genotyping, Fetal

ARUP test code 3016682

RHE GENO, Fetal Specimen Direct Amnio

RhEe Genotype Fetal, Interp

E/E

Indication for testing: Determine fetal RhEe genotype to assess risk for alloimmune hemolytic disease.

Fetal RhEe genotype: E/E

Interpretation: Two copies of the RHCE*3 (E) allele were detected in this prenatal sample; the RHCE*5 (e) allele was not identified. Homozygosity for the E allele predicts expression of E antigen and lack of expression of e antigen (also referred to as RhE+e- phenotype) in this fetus. If the pregnant patient is sensitized to e antigen, this fetus is predicted to not be at risk for anti-e-mediated alloimmune hemolytic disease.

This result has been reviewed and approved by [REDACTED]

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

Unless otherwise indicated, testing performed at:

BACKGROUND INFORMATION: RHe/e (RHCE) Antigen Genotyping, Fetal

CHARACTERISTICS: Erythrocyte alloimmunization may result in hemolytic transfusion reactions or hemolytic disease of the fetus and newborn (HDFN).

E ANTIGEN FREQUENCY: 0.29 white, 0.22 African American, 0.39 Asian.

e ANTIGEN FREQUENCY: 0.98 white, 0.98 African American, 0.96 Asian.

INHERITANCE: Co-dominant.

CAUSE: Antigen-antibody mediated red-cell hemolysis between donor/recipient or transferred maternal antibodies.

POLYMORPHISM TESTED: Rh blood group RHCE*3 (E), RHCE*5 (e): c.676G>C; p.Ala226Pro.

CLINICAL SENSITIVITY: 99 percent.

METHODOLOGY: Immucor PreciseType(TM) HEA Molecular BeadChip which is FDA-approved for clinical testing. Polymerase Chain Reaction (PCR)/Fragment Analysis.

ANALYTIC SENSITIVITY AND SPECIFICITY: 99 percent.

LIMITATIONS: Bloody amniotic fluid samples may give false-negative results because of maternal cell contamination. Rare nucleotide changes leading to altered or partial antigen expression and null phenotypes are not detected by this assay. This assay is occasionally limited in predicting genotype due to extreme variation in the Rh locus. False-negative RHe predictions may result due to RHCE-D-CE fusion genes. Patients who have had hematopoietic stem cell transplants may have inconclusive results on this test. Abnormal signal intensities may result in indeterminate genotyping results.

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.

Maternal Contamination Study Fetal Spec

Fetal Cells

Single fetal genotype present; no maternal cells present. Fetal and maternal samples were tested using STR markers to rule out maternal cell contamination.

Maternal Contam Study, Maternal Spec

whole blood

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

VERIFIED/REPORTED DATES				
Procedure	Accession	Collected	Received	Verified/Reported
RHE GENO, Fetal Specimen	23-235-113278	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
RhEe Genotype Fetal, Interp	23-235-113278	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
Maternal Contamination Study Fetal Spec	23-235-113278	00/00/0000 00:00	00/00/0000 00:00	00/00/0000 00:00
Maternal Contam Study, Maternal Spec	23-235-113278	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
Jonathan R. Genzen, MD, PhD, Laboratory Director

Patient: Patient, Example
ARUP Accession: 23-235-113278
Patient Identifiers: 01234567890ABCD, 012345
Visit Number (FIN): 01234567890ABCD
Page 3 of 3 | Printed: 8/28/2023 4:33:13 PM
4848