



VeriRef Gold®

NONINVASIVE PRENATAL SCREENING FOR THE DETECTION OF ANEUPLOIDIES AND DELETIONS/DUPLICATIONS (CNVs) IN MATERNAL BLOOD

High Risk of Sex Chromosome Trisomy (XYY)

| Patient Ref.: | |
|----------------|----------------|
| Gestional Age: | |
| | |
| Report Date: | |
| | Gestional Age: |

Number of fetuses: 1

RESULT: Sex chromosome trisomy is detected (XYY)

| Chromosomes | Result | Interpretation |
|---------------------|--------------------------|---|
| Chromosome 21 | No aneuploidy detected | Compatible with two copies of chromosome 21 |
| Chromosome 18 | No aneuploidy detected | Compatible with two copies of chromosome 18 |
| Chromosome 13 | No aneuploidy detected | Compatible with two copies of chromosome 13 |
| Chromosome X/Y | Aneuploidy detected | Compatible with one copy of chromosome X and two copies of chromosome Y (XYY) |
| Rest of chromosomes | No aneuploidy detected | Compatible with two copies of the rest of chromosomes |
| CNVs | No variants are detected | Compatible with normality |

Fetal sex: Male
Fetal fraction: 5 %

A high risk result should be confirmed from an amniotic fluid sample. In these cases we offer a free confirmation by QF-PCR or CGH array.



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Physician, technical specialist responsible for Clinical Analysis: Jaime Torrents Pont. The results relate to samples received and analyzed. This report may not be reproduced in part without permission. This document is addressed to the addressee and contains confidential information. Hereby, it is notified that any use, dissemination and/or unauthorized copying is prohibited by current legislation. Reference Laboratory has the certifications of its Quality System according to UNE-EN ISO 9001(ER-1087/1998) and its Environmental Management System according to EN ISO 14001 (GA-2001/0146) issued by AEN/NR

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Request No.:

Patient Name:

Patient Ref.:

METHODOLOGY

Whole genome analysis of fetal cfDNA by means of MPS (Massive Parallel sequencing). Illumina NextSeq® 500 Sequencer and last generation bioinformatic analysis used.

OBSERVATIONS

VeriRef Gold® is a laboratory test that detects, from the DNA released by the placenta in maternal blood, aneuploidies on all chromosomes and CNVs (deletions and duplications) on autosomal chromosomes

This test has the following yields in the case of simple pregnancies:

| Chromosome | Sensibility | Specificity |
|---------------------|------------------|-------------------|
| 21 | >99.9% (130/130) | 99.9% (1982/1984) |
| 18 | >99.9% (41/41) | 99.9% (1995/1997) |
| 13 | >99.9% (26/26) | 99.9% (2000/2002) |
| Monosomy X | 95.0% (19/20) | 99.9% (1961/1963) |
| XX | >99.9% (995/995) | 99.8% (995/997) |
| XY | >99.9% (966/966) | >99.9% (966/966) |
| Rest of chromosomes | 96.4% (136/141) | 99.8% (2001/2005) |

| Deletions and duplications | Sensibility | Specificity |
|----------------------------|---------------|-------------------|
| CNVs | 74.1% (20/27) | 99.8% (2000/2004) |

This test is valid in case of in vitro fertilization and in gestations from oocyte donation.

Noninvasive prenatal testing (NIPT) based on cell-free DNA analysis from maternal blood is a screening test; it is not diagnostic. Test results must not be used as the sole basis for diagnosis. Further confirmatory testing is necessary prior to making any irreversible pregnancy decision.

LIMITATIONS: Currently, in twin pregnancies it is only possible to detect aneuploidies on chromosomes 13, 18 and 21, and the presence of the Y chromosome. In simple pregnancies the reported sexual abnormalities are as follows: XO, XXX, XXY and XYY. This test does not detect: CNVs of size <7Mb, CNVs on sex chromosomes, polyploidies (such as triploidies), or balanced chromosomal rearrangements. A negative result does not eliminate the possibility of pregnancy presenting other genetic conditions or congenital defects. The presence of large duplications or deletions greater than 75% of the chromosome size are interpreted as aneuploidy of the complete chromosome. A false positive or negative may be due to the presence of maternal chromosomal alterations, a high index of maternal body mass, recent transfusion in the mother, mosaicism confined to placenta or the existence of an evanescent twin or fetal death.

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