Result of the non-invasive prenatal examination



LifeCodexx AG | Line-Eid-Straße 3 | DE-78467 Konstanz

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Title, last name, first name of patient

Musterfrau, Martina

Singleton or multiple pregnancy

Date of birth

Singleton pregnancy

1974-06-28*

Test option

Test option 3

Sample received on	Examination material	Bar code no.	Lab ID	QC	cffDNA content
2013-01-23*	EDTA blood	00103541	LCD01673	approved	12 %

Chromosome	Result	Interpretation	
Chromosome 21	within the normal range	No evidence of fetal trisomy 21	
Chromosome 18	within the normal range	No evidence of fetal trisomy 18	
Chromosome 13	within the normal range	No evidence of fetal trisomy 13	
Sex chromosomes	within the normal range	No evidence of Turner, Triple-X, Klinefelter or XYY syndrome	

In the event of a high-risk pregnancy, international professional associations recommend further medical clarification, e.g. a second trimester ultrasound, despite a negative test result. We request a response in the event of discordant results.

Random massively parallel sequencing (rMPS) (or: next generation sequencing, NGS) was the method applied for this analysis.

Fetal sex

female

When informing the pregnant patient of the fetal sex, please ensure that the national regulations applicable in each case are complied with.

Test method and analysis result: The PrenaTest® for the determination of the chromosomal disorders tested is based on the latest next generation sequencing (NGS) and PCR technologies using CE-marked software and CE-marked in-vitro diagnostic test systems [according to the intended purpose and declaration of conformity]. During use of the PrenaTest® in clinical practice, 100% accuracy cannot be expected. In general, no statements regarding structural chromosomal changes, mosaics or polyploidy can be made with the PrenaTest®. More information on the appraisal of results (sensitivity/specificity) and accuracy of the PrenaTest®, the limits of the examination as well as fetal sex determination can be found at www.lifecodexx.com.

Konstanz, 2018-11-29*

Validation performed by

Dr. M. Mustermann (initial validation) and J. Mustermann, M.Sc. Bioinformatics (final validation)

This results report was electronically generated and is valid without a signature.