



Non-Invasive Prenatal Test Report

Name	Jane Doe	Institution	Hospital A
Sample ID	20230510-971-0000	Ordering Physician	N/A
Date of Birth	1983-01-01	Collection Date Report	2023/05/04
MRN		Date	2023/05/12

PREGNANCY INFORMATION						QUALITY CONTROL		
Gest.Age/Weight	Ultrasound Feature	Nuchal Translucency (NT)	Multiple Marker Screening Test	In-vitro Fertilization	No. of Fetus	DNA Quality	NGS Data Quality	QC Quality
13w+4d/51kg	N/A	N/A	N/A	None	Twin	Pass	Pass	Pass

FETAL FRACTION	8.4%	FETAL SEX	Y chromosome detected *
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* The Y chromosome was detected, However, this test cannot determine if one or two of the fetuses are male.

TEST RESULT	LOW RISK		
Chromosomal Abnormality	Result	Risk after NIPT ¹⁾	Maternal age-specific risk ²⁾
Trisomy 21	LOW RISK	< 1/10000 (0.01%)	1/57
Trisomy 18	LOW RISK	< 1/10000 (0.01%)	1/157
Trisomy 13	LOW RISK	< 1/10000 (0.01%)	1/495
XO	N/A	-	1/1000
XXX	N/A	-	1/1100
XXY	N/A	-	1/900
XYY	N/A	-	1/2750
Trisomy 9	LOW RISK	N/A	
Trisomy 16	LOW RISK		
Trisomy 22	LOW RISK		

Deletion Syndrome	Result
1p36	LOW RISK
2q33.1	LOW RISK
5p15 (Cri-du-chat)	LOW RISK
11qter (Jacobsen)	LOW RISK
Other Microdeletions³⁾	Not Detected

1) Risk after NIPT: Maternal age-specific risk * Relative risk for the corresponding chromosomal results of G-NIPT
 2) Maternal age-specific risk: Average risk for the chromosomal aneuploidy in the same age group of pregnant women.
 3) >7Mb deletion tested

Tested by : Myeong-Keun Lee M.T(20058) *MKlee* Confirmed by : Sae-Mi, Lee M.D(1067) *smlee* Ju Sun, Song M.D(997) *Song Ju Sun* [1/3]





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ADDITIONAL TEST RESULT			
Chromosomal Abnormality	Result	Chromosomal Abnormality	Result
Trisomy 1	LOW RISK	Trisomy 10	LOW RISK
Trisomy 2	LOW RISK	Trisomy 11	LOW RISK
Trisomy 3	LOW RISK	Trisomy 12	LOW RISK
Trisomy 4	LOW RISK	Trisomy 14	LOW RISK
Trisomy 5	LOW RISK	Trisomy 15	LOW RISK
Trisomy 6	LOW RISK	Trisomy 17	LOW RISK
Trisomy 7	LOW RISK	Trisomy 19	LOW RISK
Trisomy 8	LOW RISK	Trisomy 20	LOW RISK

* The clinical sensitivity was not determined due to low incidence.

INTERPRETATION

No fetal chromosomal abnormalities in autosomes were found. However, we cannot completely rule out the possibility of false negative results that may be caused by factors such as maternal chromosomal microdeletion/duplication/aneuploidy, blood transfusion, confined placental mosaicism (CPM) and fetal mosaicism. If any fetal abnormalities are found by ultrasonography, it is recommended to perform high-resolution cytogenetic testing regardless of the result of G-NIPT.

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TEST INFORMATION

- Test Method: Next Generation Sequencing (NGS)
- Test Subject: Fetal Trisomy (Chromosome 21, 18, 13, 9, 16, 22), Sex Chromosome Aneuploidy, Microdeletion Syndrome (>7Mb)
- Specimen Type: cfDNA tube WB 10mL
- Bioinformatics Pipeline: NIPT.v1.2

TEST PERFORMANCE

Test Item	Sensitivity	Specificity	NPV	PPV
Trisomy 21	99.79%	99.98%	>99.99%	98.73%
Trisomy 18	99.33%	99.97%	>99.99%	93.67%
Trisomy 13	>99.99%	99.98%	>99.99%	81.40%
Sex Chromosome Aneuploidies (XO, XXX, XXY)	>99.99%	99.86%	>99.99%	50.00%
Other Chromosomes	The clinical sensitivity was not determined due to low incidence.			
Microdeletion Syndrome	The clinical sensitivity was not determined due to low incidence, and the sensitivity may be significantly affected by factors such as fetal DNA fraction and microdeletion size.			

* Test performance is based on the G-NIPT test result conducted between 2015.12 ~ 2021.12 and may be changed in the future.

METHOD and LIMITATIONS

- The purpose of this test is for risk assessment of common fetal trisomies 21, 18, 13 and sex chromosome aneuploidies. This test is performed by massively parallel sequencing for whole-genome using circulating cell-free fetal DNA in maternal plasma and it is possible to detect abnormalities in all chromosomes as well as chromosome 21, 18 and 13. NIPT performance is superior to the existing prenatal multiple marker screening tests.
- This test cannot identify neural tube defects and polyploidy such as triploidy and tetraploidy.
- This test does not report monosomy.
- In case of trisomy 9, 16, 22 and microdeletion syndrome, the clinical sensitivity was not determined due to low incidence, and the sensitivity may be significantly affected by factors such as fetal DNA fraction and microdeletion size.
- This test is not to verify fetal karyotypes but is to determine the risk of fetal aneuploidies. If the result is positive, confirmatory test such as fetal karyotyping should be performed. Moreover, this is not a diagnostic test which does not rule out probability of false positive or false negative results.
- The factors affecting accuracy of this test are as follows: low fetal DNA fraction (early gestational weeks and high maternal BMI), undetermined maternal chromosomal abnormalities, confined placental mosaicism, fetal chromosomal mosaicism, multiple gestation, arithmetic error of calculating fetal DNA fraction, and maternal status (cancer, blood transfusion, transplantation, chemotherapy, stem cell treatment, or autoimmune disease), etc.

REFERENCE

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- JAMA. 2015 Jul 14;314(2):162-9. Noninvasive Prenatal Testing and Incidental Detection of Occult Maternal Malignancies
- N Engl J Med. 2015 Apr 23;372(17):1639-45. Copy-number variation and false positive prenatal aneuploidy screening results
- Clin Genet. 2016 May;89(5):523-30. Clinical implementation of NIPT - technical and biological challenges
- Fetal Diagn Ther. 1995 Nov-Dec;10(6):356-67

※ This test was developed and its performance characteristics determined by GC Genome. It has not been cleared or approved by the Korean Ministry of Food and Drug Safety (MFDS).

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