

## APPLICATION FORM AND INFORMED CONSENT INFORMATION ON THE POTENTIAL AND LIMITATIONS OF THE TEST

AFFIX LABEL

PLEASE COMPLETE ALL RELEVANT DATA IN CAPITAL LETTERS  PATIENT	TEST DECUMEN
	TEST REQUIRED
Surname	Basic
Name	BASIC PLUS DIGEORGE SYNDROME
Date of birth DD / MM / YYYY	Basic Plus + 90 Microdeletions
Place of birth	KARYOTYPE
	KARYOTYPE PLUS
Address - Post Code - City	KARYOTYPE PLUS + MONOGENIC DISEASES
Country	TOTAL SCREEN
C.F.	MONOGENIC DISEASES
Phone Number	MINI CGS 1  Basic Plus + 90 Microdeletions +  Maternal Carrier Screening
Email	MINI CGS 2  KARYOTYPE + MATERNAL CARRIER SCREENING
Date / / / / / / DD/MM/YYYY	MINI CGS 3  KARYOTYPE PLUS + MONOGENIC DISEASES + MATERNAL CARRIER SCREENING
PREGNANCY (ALL FIELDS ARE REQUIRED)	
I want to be informed of <b>fetal sex?</b> Yes NO	ETHNIC GROUP
	(required)
PREGNANCY  Single Monochorional Bicorial	Caucasian African North African
Single twin twin	Asian Other
Spontaneous Homologous Heterologous IVF Heterologous IVF (Age)	Weight (required) Height (required)
Date Last Period DD / MM / YYYY	Smoker Yes No
	Clinical History
Actual gestational age at the date of collection	
WEEKS DAYS	
DOCTOR (LARORATORY)	
DOCTOR / LABORATORY  Surname of the doctor (required)	Address
Name of the doctor (required)	Post Code City
Doctor's phone number	Email
Laboratory / Clinical Diagnostic Center of Belonging (required)	Date DD / MM / YYYY
	Doctor's signature that has collected the informed consent



## INFORMATION ON THE POTENTIAL OF THE TEST AND ACCEPTANCE OF ITS LIMITATIONS

TEST SELECTION

THE ANALYSES AVAILABLE WITHIN THIS TEST MUST BE SELECTED BY THE PREGNANT WOMAN UNDER THE GUIDANCE OF A SPECIALIST, WHO WILL ENSURE THAT THIS INFORMED CONSENT IS FULLY UNDERSTOOD IN ACCORDANCE WITH THE NEEDS AND REQUESTS OF THE PARENTAL COUPLE.

I hereby declare that I have received detailed information regarding the NIPT screening test that I have selected and requested.

I fully understand that, although the test I am undergoing demonstrates extremely high diagnostic accuracy, as stated by the current guidelines in use in our country, definitive diagnostic certainty is provided exclusively by invasive tests (Amniocentesis and Chorionic Villus Sampling).

I also understand that all fetal DNA tests (NIPT) do not provide definitive diagnostic certainty. While rare, cases of both false positives and false negatives have been reported, and I accept this rare eventuality.

BASIC exclusively investigates the most common forms of chromosomal anomalies, **Down syndrome** (trisomy of chromosome **21**), **Edwards syndrome** (trisomy of chromosome **18**), and **Patau** syndrome (trisomy of chromosome **13**), with a **sensitivity of 99,8%**, as outlined in the current guidelines. **If requested, fetal sex determination can also be provided; however, as previously stated, it does not include the detection of sex chromosome anomalies.** 

BASIC PLUS investigates the three main fetal chromosomal aneuploidies associated with chromosomes 21, 18, and 13 (as previously described), as well as the X and Y sex chromosomes, with a sensitivity of 99,8%. It also determines fetal sex, which, upon our request, can remain

Additionally, I understand that the likelihood of incorrect interpretation of fetal sex, as estimated in global literature, is 3%.

undisclosed.			
			on caused by the deletion of a portion of
	ence of approximately 1 in 3,500 live birt		
I confirm my request for the so	creening of <b>DiGeorge Syndrome</b> ? YES	NO	
<b>21, 18, and 13,</b> as well as the chromosomal alterations caus is nonetheless correlated to th any NIPT test. The term micro in the loss of genetic inform	X and Y sex chromosomes (with a sense by structural rearrangements (referred e fetal fraction (sensitivity increases with deletions/microduplications refers to an	sitivity of 99,8%). The test also includes ed to as microdeletions/microduplication in a higher fetal fraction, up to 90%). Highe omalies characterized by the absence of in of extra genomic material (microdupli	euploidies associated with chromosomes the screening of a large number of small s) at an average resolution of 5Mb, which or sensitivity values cannot be achieved by a small chromosomal segment, resulting cations). Other molecular variants than
The complete list of investigat	ions is as follows:		
1p31, microduplication	5q12, microdeletion	11p13, <b>WAGR</b>	17q11.2, microduplication
1p36, microdeletion	5q35.3, <b>Sotos</b>	11p15-p14, microdeletion	17p11.2, <b>Potocki-Lupski</b>
1q21q32, monosomy	6p21, Cleidocranial Dysplasia	11q, <b>Jacobsen</b>	17p11.2, <b>Smith-Magenis</b>
1q21.1, microdeletion	6q24-q25, microdeletion	11q23.3-q25, microdeletion	17p13.3, Miller-Dieker
1q21.1, microduplication	7q11.23, microduplication	12q14, microdeletion	17q21, Koolen-de Vries
1q23qter, trisomy	7q11.23, Williams-Beuren	13q14, microdeletion	17q21.31, microduplication
1q41-q42, microdeletion	7q21.q31, trisomy	13q21qter, monosomy	18p, microdeletion
1q42qter, monosomy	7q32qter, monosomy	13q21qter, trisomy	18pterq12, trisomy
2p15-p16.1, microdeletion	7q32qter, partial trisomy	14q11-q22, microdeletion	18q, microdeletion
2q22.3, Mowat-Wilson	8p23.1, microdeletion	14q24-qter, trisomy	18q12qter, trisomy
2q33.1, microdeletion	8p23.1, microduplication	14q32.13, Wilms tipo 1	19p13, microduplication
2q33.1, microduplication	8q12.1-q21.2, microdeletion	15q11, <b>Angelman</b>	19q13.11, microdeletion
2q35, microduplication	8q13.3, Branchio-Oto-Renal	15q11-q13, <b>Prader-Willi</b>	20p, trisomy
2q37, microdeletion	Syndrome	15q14, microdeletion	20p12, <b>Alagille</b>
3p11-p21, monosomy	8q21qter, monosomy	15q22qter, trisomy	20q13.1-q13.3, microduplication
3q22, <b>Dandy-Walker</b>	8q21.11, microdeletion	15q26-qter, microdeletion	22q11.2, <b>DiGeorge</b>
3p25pter, monosomy	8q24.11, Langer-Giedion	15q26-qter, microduplication	22q11.2, microduplication
3q29, microdeletion	9p, microdeletion	15q26.1, <b>Congenital</b>	22q13, Phelan-mcdermid
3q29, microduplication	9q22.3-q33, microdeletion	Diaphragmatic Hernia Type 1	Xp11.3, microdeletion
4p16.3, Wolf-Hirschhorn	9q33.2-q34.3, microduplication	16p11.2-p12.2, microdeletion	Xp11.23-p11.22, microduplication
4q21q31, monosomy	9q34, <b>Kleefstra</b>	16p11.2-p12.2, microduplication	Xp21.3, Lissencefalia X-linked
4q31qter, monosomy	10q26, microdeletion	16p13.3, <b>Rubinstein-Taybi</b>	Xq27.3-q28, microduplication
5n <b>Cri-du-chat</b>	11n Potocki-Shaffer	17a11 2 microdeletion	Xq28, microdeletion

KARYOTYPE extends the investigation of numerical alterations to all chromosomes. This NIPT, In other words, detects the presence of an altered number in all 23 pairs of chromosomes related to the fetal karyotype, including the sex chromosomes X and Y (with a sensitivity of 99,8%).



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mentioned so far: <b>numerical alterations</b> , known as an euploidies (at the <b>sex chromosomes</b> ( <b>X</b> and <b>Y</b> , along with their numerical alterations <b>duplications</b> , at an average reso <b>fetal fraction</b> , <b>up to 90%</b> ). Higher sensitivity values cannot be achiencluded in the results and is clearly outlined on the FetalDNA.it was	non-invasive test of circulating free fetal DNA (NIPT). It includes all the investigations alterations in chromosome number only), for <b>all chromosomes</b> , including <b>13</b> , <b>18</b> , <b>21</b> , and tions) with a <b>sensitivity of 99,8%</b> . Additionally, it includes the analysis of <b>90 syndromes</b> lution of 5Mb, which depends on the fetal fraction ( <b>sensitivity increases with a higher</b> eved by any test. The list of the 90 main syndromes investigated in this screening will be ebsite and on the previous page under the section 'Base Plus+90 Microdeletions.' <b>t, the screening for the most common maternal Cystic Fibrosis mutations</b> .	
I confirm that I request the screening for the most common mater	rnal Cystic Fibrosis mutations? YES NO	
This test is a screening test and not a diagnostic one. Although high	uested individually or in combination with the levels described above) ghly accurate, the results do not have diagnostic value and must be assessed within the s not a substitute for invasive prenatal diagnostic procedures (Chorionic Villus Sampling	
Hereditary Monogenic Diseases	Monogenic Diseases due to de novo mutations	
Autosomal Recessive Polycystic Kidney Disease ( <b>PKHD1</b> gene)	Achondroplasia ( <b>FGFR3</b> gene)	
Beta-Thalassemia ( <b>HBB</b> gene)	Apert Syndrome ( <b>FGFR2</b> gene)	
Congenital Adrenal Hyperplasia (CYP21A2 gene)	Crouzon Syndrome (FGFR2 gene)	
Congenital Deafness ( <b>GJB2</b> gene)	Hypochondroplasia ( <b>FGFR3</b> gene)	
Fetal Cystic Fibrosis ( <b>CFTR</b> gene)	LEOPARD Syndrome ( <b>PTPN11</b> gene)	
Hemochromatosis ( <b>HFE</b> gene)	Noonan Syndrome (PTPN11 gene), (RAF1 gene), (SOS1 gene)	
Phenylketonuria ( <b>PAH</b> gene)	Pfeiffer Syndrome ( <b>FGFR2</b> gene)	
Rett Syndrome ( <b>MECP2</b> gene)	Thanatophoric Dysplasia ( <b>FGFR3</b> gene)	
I have been fully informed that it is not possible to obtain definitive evidence of these anomalies in the fetus through maternal blood testing. The test I am undergoing has a <b>sensitivity of up to 90%</b> . Higher values are NEITHER realistic NOR documented and cannot be achieved by any NIPT test based on fetal DNA. Certainty (as reiterated by the Guidelines of our country) can only be provided by invasive tests (Amniocentesis or Chorionic Villus Sampling).		
TOTAL SCREEN represents a highly advanced and co	omprehensive non-invasive prenatal test (NIPT) performed on circulating free fetal DNA.	
It analyzes all <b>23 pairs of chromosomes</b> related to the fetal karyotype, including the investigation of <b>sex chromosome aneuploidies (X, Y)</b> as well as the three main fetal chromosomal aneuploidies associated with <b>chromosomes 21, 18, and 13: Down syndrome</b> (Trisomy 21), <b>Edwards syndrome</b> (Trisomy 18), and <b>Patau syndrome</b> (Trisomy 13), with a <b>sensitivity of 99,8%</b> .		
at an average resolution of 5Mb, which depends on the fetal fraction	caused by structural rearrangements (referred to as microdeletions/microduplications) on (sensitivity increases with a higher fetal fraction, reaching up to 90%). Additionally, it seases as listed in the previous section, with a sensitivity of up to 90%.	
The FetalDNA Total Screen also includes investigations related to t	he pregnant woman, specifically:	
Screening for mutations responsible for maternal Cystic Fibrosis		
	enes associated with Spinal Muscular Atrophy (this test excludes the vast majority of	
<ul> <li>Screening for infectious agents present in the maternal blood, pregnancy (while reliable and thorough, it does not rule out fetal da</li> </ul>	allowing early detection prior to the positivity of antibody tests routinely used during image caused by infections occurring before or after the test).	
Screening for mutations associated with predisposition to preter	rm birth (this test does not exclude preterm birth caused by other clinical factors).	
<ul> <li>Biochemical risk assessment for preeclampsia (provides a risk value that, although highly useful for the treating physician, does not offer diagnostic certainty).</li> </ul>		
• Screening for the most common mutations responsible for <b>hereditary thrombophilia</b> (these investigations, considered by much of the international literature to be useful for preventing maternal-fetal complications such as miscarriage, growth restriction, placental abruption, and thrombosis, must be evaluated in the clinical context and do not exclude the existence of other factors causing the same conditions).		
MINI CGS 1 includes the investigations indicated in the Base Plus + 90 Microdeletions combined with Maternal Carrier Screening, which analyzes:		
<ul> <li>Cystic Fibrosis, Congenital Deafness, and Spinal Muscular Atrophy (SMA) inherited from both parents (recessive transmission).</li> <li>Muscular Dystrophy and Fragile X Syndrome inherited from healthy carrier mothers (X-linked transmission).</li> <li>If any positive results are found in the Maternal Carrier Screening, I will be contacted for genetic counseling.</li> </ul>		
MINI CGS 2 includes the investigations indicated in the Karyotype Plus combined with Maternal Carrier Screening, which analyzes:  - Cystic Fibrosis, Congenital Deafness, and Spinal Muscular Atrophy (SMA) inherited from both parents (recessive transmission).  - Muscular Dystrophy and Fragile X Syndrome inherited from healthy carrier mothers (X-linked transmission).  If any positive results are found in the Maternal Carrier Screening, I will be contacted for genetic counseling.		

- which tests for:Cystic Fibrosis, Congenital Deafness, and Spinal Muscular Atrophy (SMA) inherited from both parents (recessive transmission).
  - Muscular Dystrophy and Fragile X Syndrome inherited from healthy carrier mothers (X-linked transmission).

MINI CGS 3 includes the investigations indicated in the Karyotype Plus and monogenic diseases combined with Maternal Carrier Screening,

If any positive results are found in the Maternal Carrier Screening, I will be contacted for genetic counseling.



## INFORMATION ON THE POTENTIAL OF THE TEST AND ACCEPTANCE OF ITS LIMITATIONS

- Regarding the detection of fetal anomalies in maternal blood (NIPT), I fully understand that the test I am undergoing, as stated by the current Guidelines in use in our country, does not provide diagnostic certainty, which is exclusively offered by invasive tests (Amniocentesis and Chorionic Villus Sampling). I also understand that all fetal DNA tests (NIPT) do not provide definitive diagnostic results. Although rare, cases of false positives and false negatives have been reported. I accept this rare eventuality. Furthermore, the possibility of incorrect fetal sex determination is 3%. While this occurrence has no clinical relevance, it must be acknowledged due to its emotional impact.
- Additionally, NIPT does not detect balanced chromosomal rearrangements. It may fail to detect fetal and/or placental chromosomal mosaicisms (two cell
  lines with different chromosomal configurations). It does not analyze all point mutations associated with the investigated genes, and its sensitivity does not
  exceed 90%. It does not detect methylation defects, triploidy, polyploidy, or any chromosomal and molecular rearrangements that cannot be identified using
  NIPT techniques.
- If the screening test yield a positive result, current guidelines require confirmation through invasive prenatal diagnosis Chorionic Villus Sampling or Amniocentesis). These procedures will be scheduled at our Rome center completely **free of charge**, covering both the sampling technique and genetic testing.
- Result reporting times vary depending on the type of test requested and may be delayed due to technical issues or the need for additional analytical checks.
- I undertand that this NIPT, although performed using the most advanced molecular technologies, may not provide a result and might need to be repeated (approximately 1% of cases in the literature). This can also occur when a low percentage of fetal DNA is detected (generally below 4%). In such cases, an invasive diagnostic procedure is recommended, as a low level of fetal DNA in maternal blood may indicate an increased risk of chromosomal abnormalities. FetalDNA (like all NIPT tests) is performed by quantitatively comparing the DNA of selected chromosomes in maternal blood with that of fetal origin. Most of this DNA is of maternal origin, with only a small proportion being fetal. The test determines if the amount of DNA from a specific chromosome deviates from the expected amount. For example, an excess of DNA from chromosome 21 could indicate that the fetus has three copies of this chromosome (causing Down syndrome) instead of the usual two. The minimum threshold of 4% required to obtain a sufficiently reliable diagnosis has been defined through statistical models based on the minimum number of readings of aneuploid chromosome fragments sufficient to detect fetal aneuploidy at various levels of fetal fraction (FF). According to this model, at low FF levels, differences in circulating cfDNA between pregnancies with fetal trisomies and those with euploid fetuses may not be detectable, leading to false negatives. A factor associated with low fetal cfDNA percentages and the potential failure of the test is increased maternal body weight. The increased amount of maternal cfDNA in obese women may mask the fetal fraction, complicating the screening for aneuploidies and increasing the risk of test failure due to a high body mass index (BMI >30 in obesity and between 25 and 30 in overweight cases).
- It is emphasized and reiterated that any mutations other than those specifically targeted and reported in the test results will not be investigated, and the test has no capability to verify their presence.
- When the test needs to be repeated, a new blood sample is collected at no additional cost.

I hereby declare that I have FULLY UNDERSTOOD

the limitations of the selected screening test

- In dizygotic twin pregnancies, it is not possible to distinguish the condition of each fetus or accurately assess sex chromosome aneuploidies. However, the presence or absence of the Y chromosome can be detected. If the Y chromosome is identified, it cannot be determined whether one or both fetuses are male. In pregnancies that began as twin or multiple gestations followed by spontaneous miscarriage of one or more fetuses with reabsorption of the gestational sac (vanishing twin), the maternal blood may contain free fetal DNA from the miscarried fetus. This could interfere with the quality of results, leading to false positives if the miscarriage was caused by chromosomal aneuploidies in the lost fetus. Similarly, there may be discrepancies in sex determination (e.g., male sex identified due to the presence of the Y chromosome originating from the miscarried fetus).
- In cases of chromosomal mosaicism (with a frequency of approximately 1-2%), result discrepancies (false positives or false negatives) may occur. Specifically, the test may produce a positive result (aneuploidy detected), but the chromosomal anomaly is confined to the placenta due to chromosomal mosaicism. In such cases, the fetus may present a normal karyotype during invasive prenatal diagnosis (false positive). Conversely, the test may produce a negative result (aneuploidy not detected), but the fetal DNA without aneuploidy may be confined to the placenta due to chromosomal mosaicism, resulting in a fetus with an aneuploid karyotype during invasive prenatal diagnosis (false negative).

**FIELD REQUIRED** 

	January 1980	, and the second se		
Signature / Signatures pregnant woman		Doctor's signature that has collected the informed consent		
Your privacy is a priority for ALTAMEDICA. Artemisia SpA, whose registered office is located in Rome at Viale Liegi, 41 as the data controller, informs you that your data will be handled in compliance with the applicable laws and EU Regulation No. 2016/679. Your identity and all data related to your personal information will remain confidential, and only authorized personnel will have access to this information, along with competent authorities when required by local jurisdiction laws. We wish to inform you that your personal data will be processed solely for the following purposes: (1) To fulfill obligations arising from the provision of services you have subscribed to; (2) For research purposes, scientific publications, and presentations, provided that your data remains anonymous and cannot be identified during data analysis, and any identifiable data will be removed from any publication.				
In accordance with persona for no longer than permitted		s consent to perform the requested diagnostic tests and process their data, which will be stored		
You may exercise your rights at any time, including access, rectification, objection, deletion, withdrawal, automated decision-making, restriction, and portability, by contacting the company at Artemisia spa con sede in Viale Liegi 41 - Roma via registered letter with acknowledgment of receipt, or at the following certified email address: artemisiaspa@pec.it. Alternatively, you may contact the company's designated Data Protection Officer (DPO) at dpo@artemisia.it.				
Authorize	Do not authorize			
I authorize/do not authorize	the sending of the report to my email address	Signature / Signatures pregnant woman		

I hereby declare that I have NOT FULLY UNDERSTOOD

the limitations of the selected screening test