



**Prima visita -
counseling
prenatale: screening
delle anomalie
cromosomiche, dei
disordini
ipertensivi, della
sindrome metabolica**

Enrico Ferrazzi
Daniela Di Martino



Fondazione IRCCS Ca' Granda
Ospedale Maggiore Policlinico

Sistema Socio Sanitario
 Regione
Lombardia



UNIVERSITÀ DEGLI STUDI
DI MILANO

*Direttore -MANGIAGALLI CENTER
IRCCS CA GRANDA POLICLINICO
University of Milan, School of Medicine,*

Amniocentesi e CVS per età



Invasive diagnostic testing

CVS / Amnio	Post-test risk
T21, 18 & 13	0
Other chromosomal	1 in 10,000
Uncertain results	1 in 100
Fetal loss	1 in 100
Maternal death	Very rare but not 0

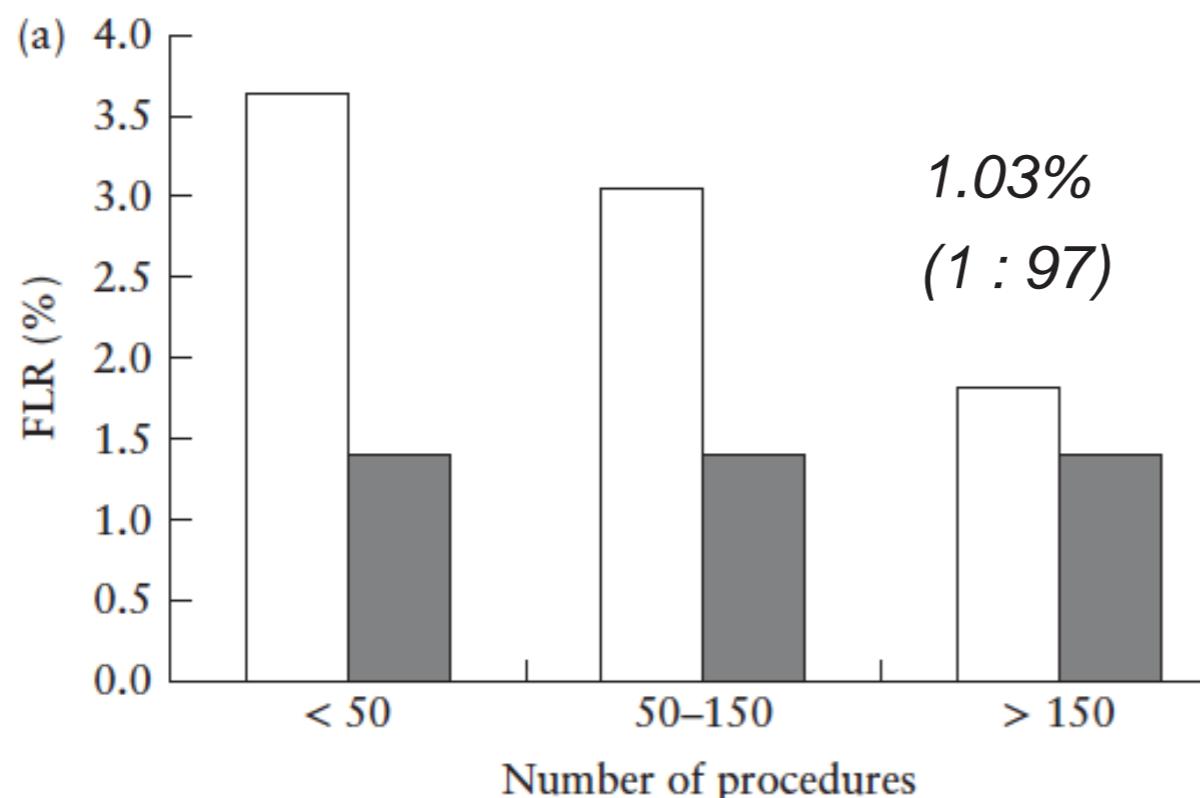
3 feti normali
persi per ogni
T21



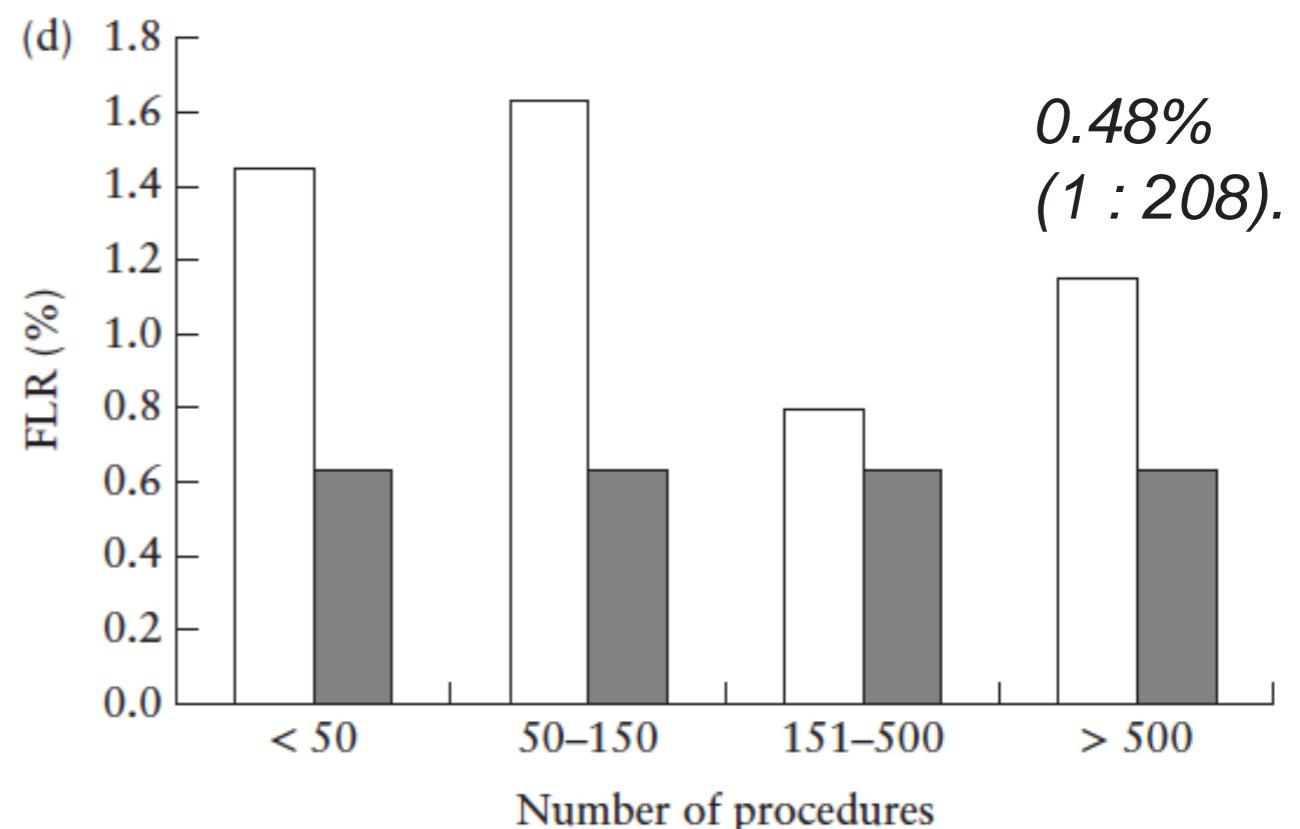
Total pregnancy loss after chorionic villus sampling and amniocentesis: a cohort study

M. BAKKER¹, E. BIRNIE^{1,2}, P. ROBLES DE MEDINA³, K. M. SOLLIE¹, E. PAJKRT³
and C. M. BILARDO¹

CVS



amniocentesis



2018 QUALE SCENARIO



EDITORS' CHOICE

Prospective first-trimester screening for trisomy 21 in 30,564 pregnancies

Kyriaki Avgidou, MD,^a Aris Papageorghiou, MD,^a Renu Bindra, MD,^a
Kevin Spencer, MD,^b Kypros H. Nicolaides, MD^{a,*}

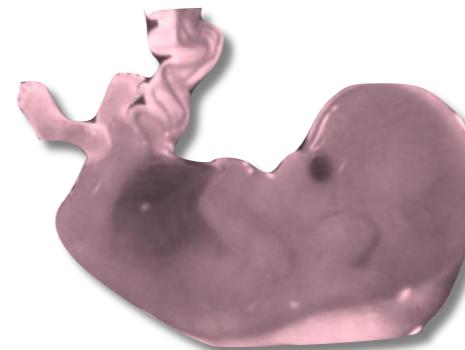
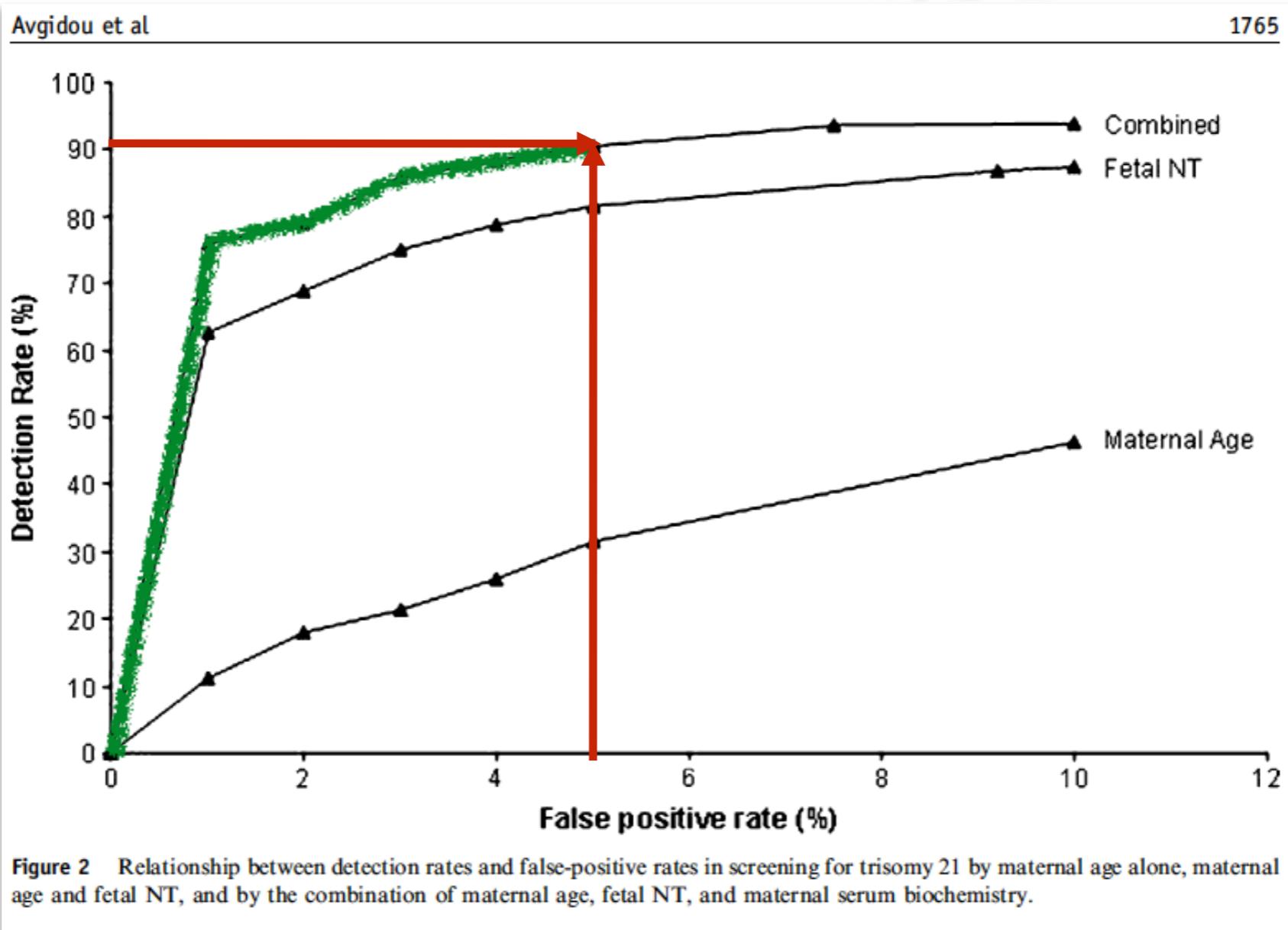


Figure 1 Fetus with nuchal translucency (NT) at the end of the first trimester kindly provided by Dr. Kypros Nicolaides, University of Southampton.

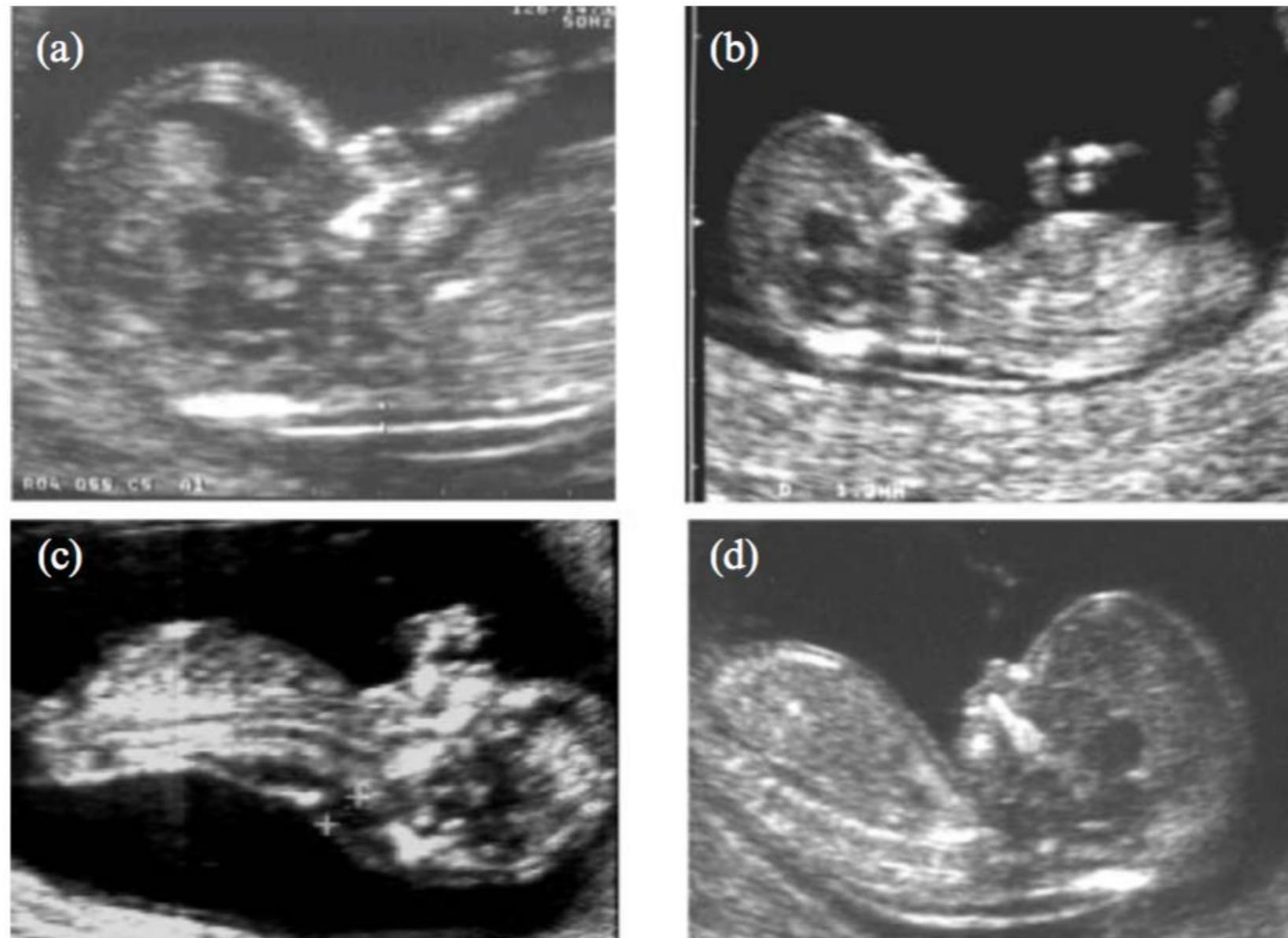




EDITORS' CHOICE

Prospective first-trimester screening for trisomy 21 in 30,564 pregnancies

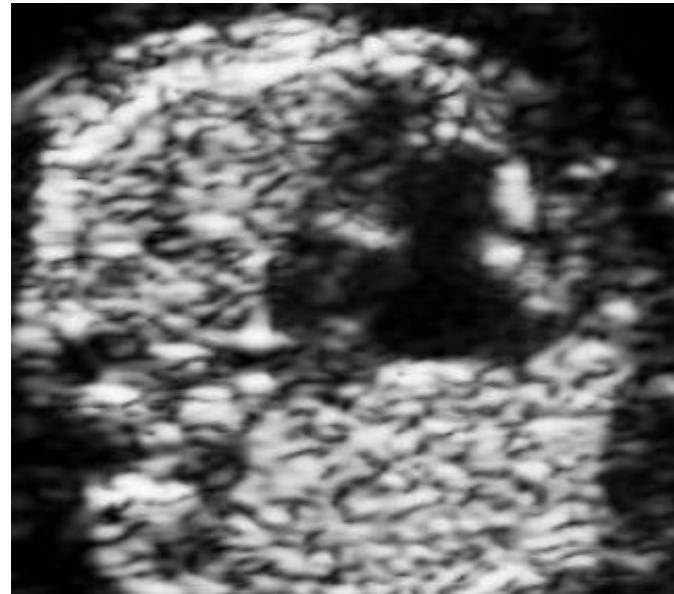
Kyriaki Avgidou, MD,^a Aris Papageorghiou, MD,^a Renu Bindra, MD,^a
Kevin Spencer, MD,^b Kypros H. Nicolaides, MD^{a,*}





Early detection of fetal defects

Atrioventricular septal defect



T 21/18/13 = 65%

Exomphalos



T 18/13 = 55%

Facial cleft



T 18/13 = 83%

Hydrops



T 21/18/13 = 80%



Clinical implementation of routine screening for fetal trisomies in the UK NHS: cell-free DNA test contingent on results from first-trimester combined test

M. M. GIL*, R. REVELLO*, L. C. POON*, R. AKOLEKAR*† and K. H. NICOLAIDES*

Table 3 Distribution of risk from the combined test according to trisomic outcome

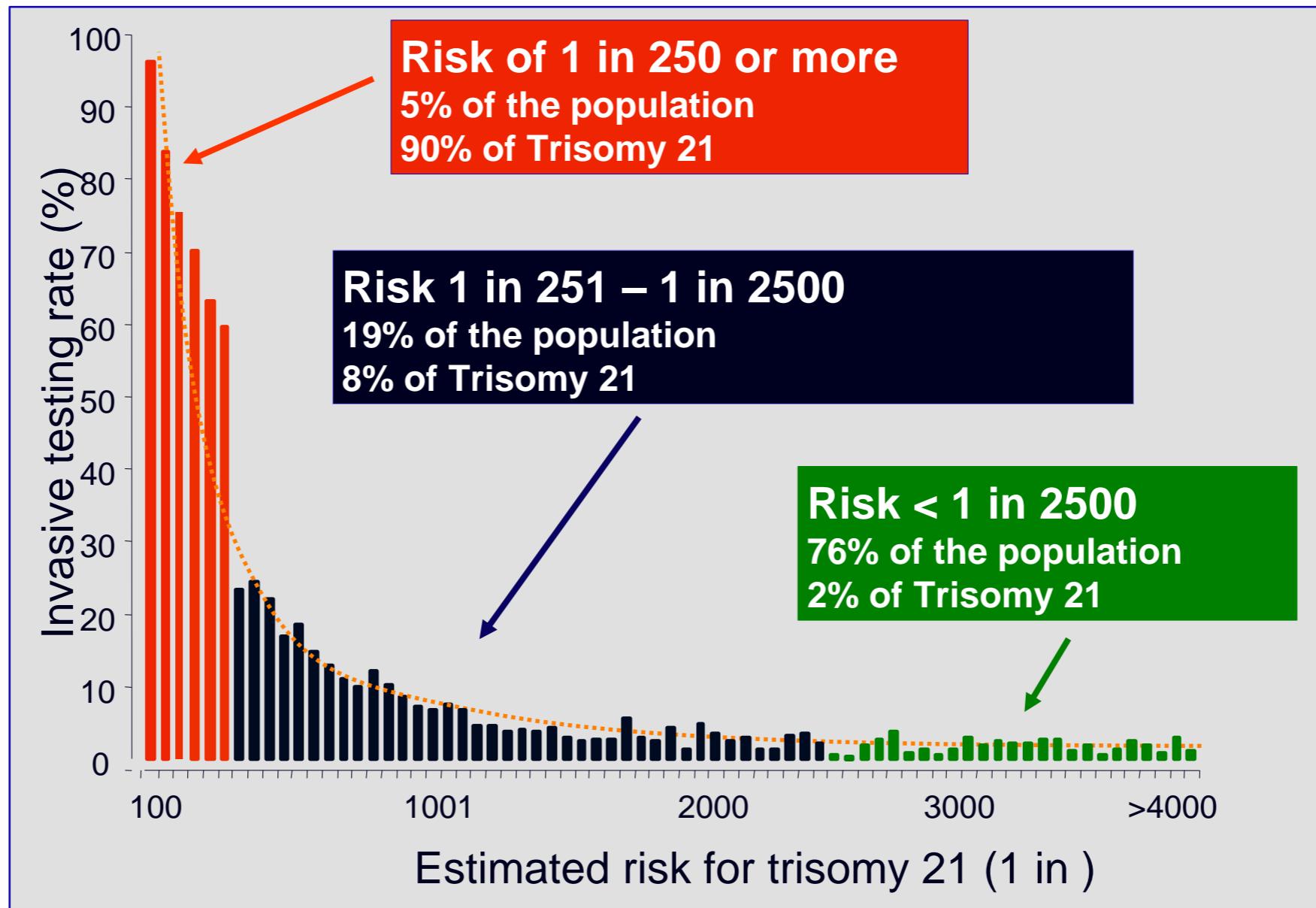
11.692

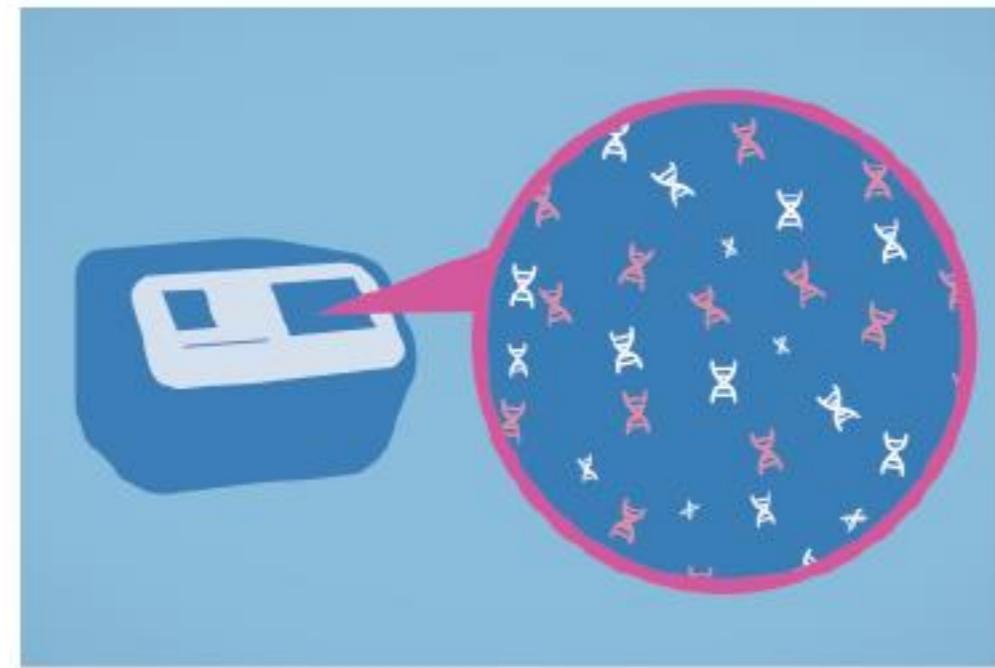
<i>Risk cut-off</i>		<i>Trisomy 21 (n = 47)</i>
≥ 1 in 10		30 (63.8)
≥ 1 in 20		36 (76.6)
≥ 1 in 50		38 (80.9)
≥ 1 in 100	+5	41 (87.2)
≥ 1 in 500		46 (97.9)
≥ 1 in 1000		46 (97.9)
≥ 1 in 1500		46 (97.9)
≥ 1 in 2000		46 (97.9)
≥ 1 in 2500	+1	46 (97.9)
≥ 1 in 3000		46 (97.9)
≥ 1 in 3500		47 (100)

The combined test detection rate of 87% for T21 detection rate of 93% for T18/13 a FPR of 3.4%;



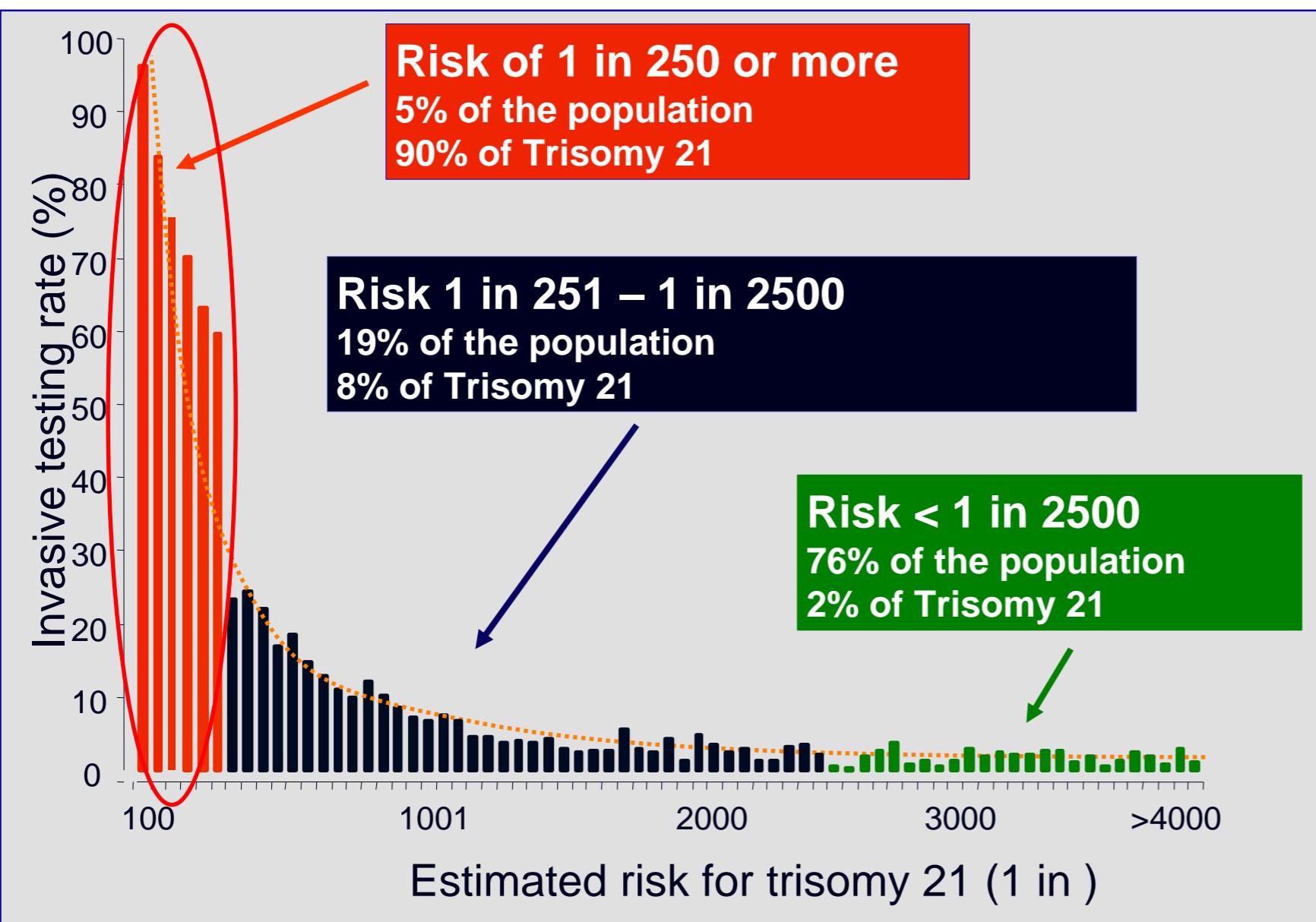
Risk distribution







Contingent screening



- Risk \geq 1 in ?**
- cfDNA to all?
 - Risk stratification?
 - Nuchal translucency?
 - Major fetal defects?
- Impact on invasive testing rate



Clinical implementation of routine screening for fetal trisomies in the UK NHS: cell-free DNA test contingent on results from first-trimester combined test

M. M. GIL*, R. REVELLO*, L. C. POON*, R. AKOLEKAR*† and K. H. NICOLAIDES*

<i>Risk cut-off</i>	<i>Trisomy 21 (n=47)</i>
≥ 1 in 10	30 (63.8)
≥ 1 in 20	36 (76.6)
≥ 1 in 50	38 (80.9)
≥ 1 in 100	41 (87.2)
≥ 1 in 500	46 (97.9)
≥ 1 in 1000	46 (97.9)
≥ 1 in 1500	46 (97.9)
≥ 1 in 2000	46 (97.9)
≥ 1 in 2500	46 (97.9)
≥ 1 in 3000	46 (97.9)
≥ 1 in 3500	47 (100)

Women with a risk **≥1 in 100 (high-risk group)**
were offered options of invasive testing, cfDNA
testing or no further testing,

Women with a between **1 in 101 and 1 in 2500
(intermediate-risk group)**
were offered cfDNA testing or no further
testing



Clinical implementation of routine screening for fetal trisomies in the UK NHS: cell-free DNA test contingent on results from first-trimester combined test

M. M. GIL*, R. REVELLO*, L. C. POON*, R. AKOLEKAR*† and K. H. NICOLAIDES*

Table 3 Distribution of risk from the combined test according to trisomic outcome

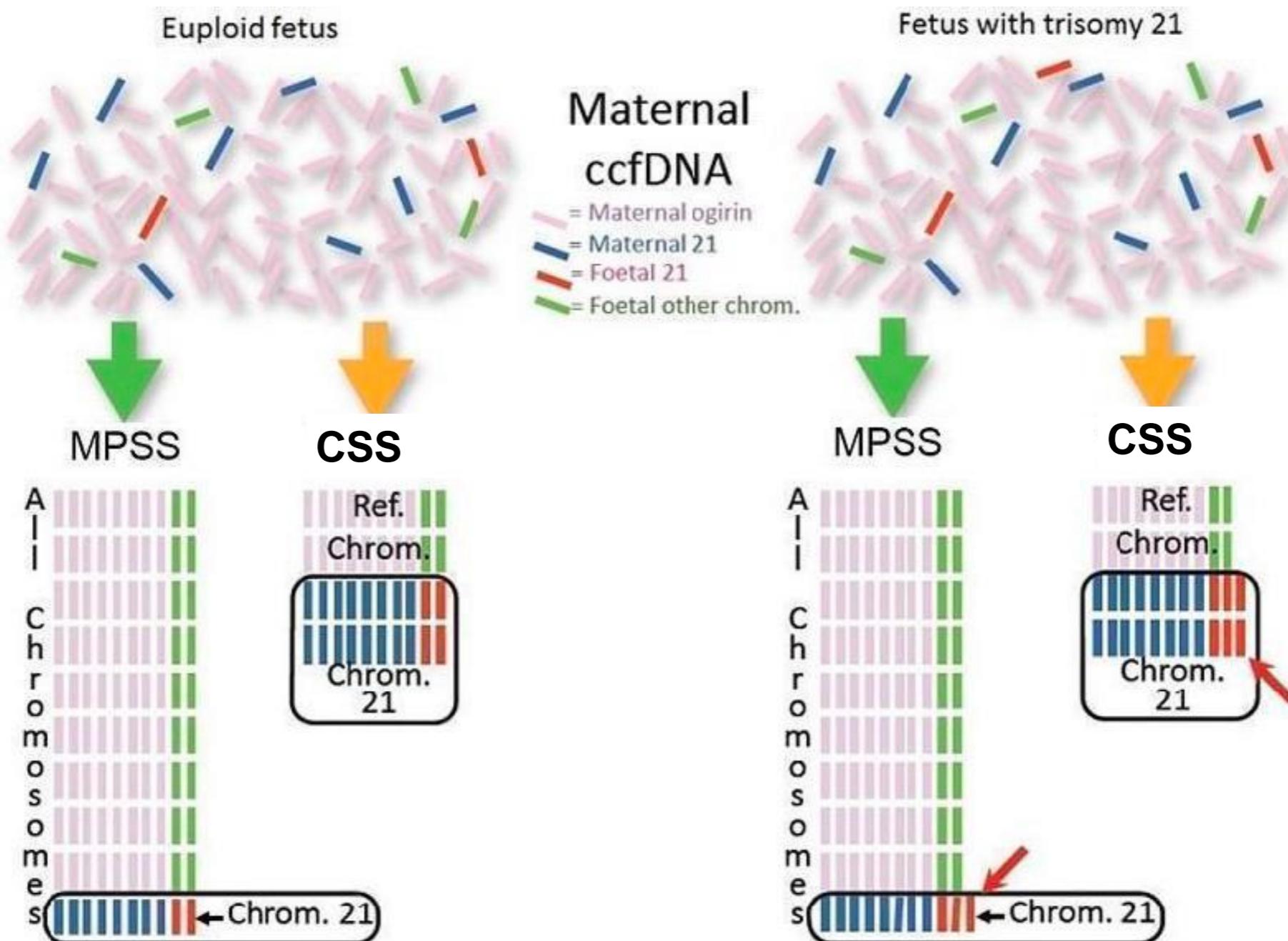
11.692

<i>Risk cut-off</i>		<i>Trisomy 21 (n = 47)</i>
≥ 1 in 10		30 (63.8)
≥ 1 in 20		36 (76.6)
≥ 1 in 50		38 (80.9)
≥ 1 in 100	+5	41 (87.2)
≥ 1 in 500		46 (97.9)
≥ 1 in 1000		46 (97.9)
≥ 1 in 1500		46 (97.9)
≥ 1 in 2000		46 (97.9)
≥ 1 in 2500	+1	46 (97.9)
≥ 1 in 3000		46 (97.9)
≥ 1 in 3500		47 (100)

The combined test detection rate of 87% for T21 detection rate of 93% for T18/13 a FPR of 3.4%; cfDNA test **detection rate of 98% for T21** **detection rate of 82% for T18/13 a FPR of 0.25%.**

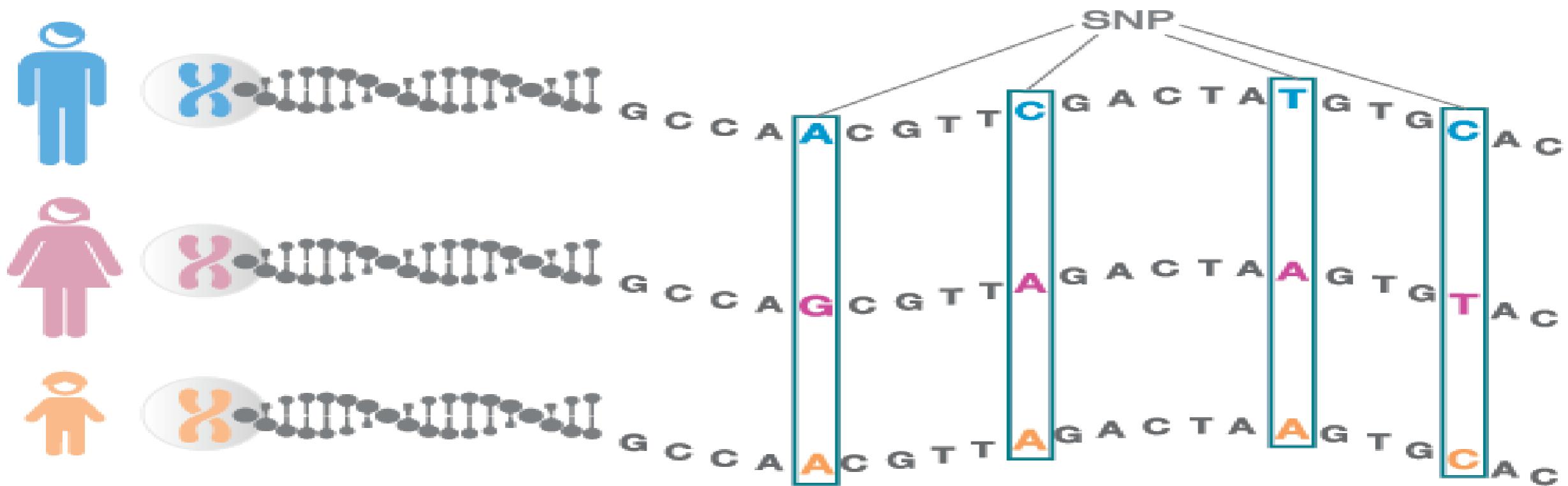


Badeau M, Lindsay C, Blais J, Takwoingi Y, Langlois S, Légaré F, Giguère Y, Turgeon AF, Witteman W, Rousseau F





SNP = Single Nucleotide Polymorphism



- Polymorphisms occur when a single base pair (nucleotide) is changed: A, T, C, or G
 - These are **normal** genetic changes that occur in every person and mark where people differ from one another

metanalisi delle casistiche pubblicate con le tre metodologie:

- 1.CSS, chromosome-specific sequencing
- 2.MPSS, massively parallel shotgun sequencing
- 3.SNP, single nucleotide polymorphism



Analysis of cell-free DNA in maternal blood in screening for aneuploidies: updated meta-analysis

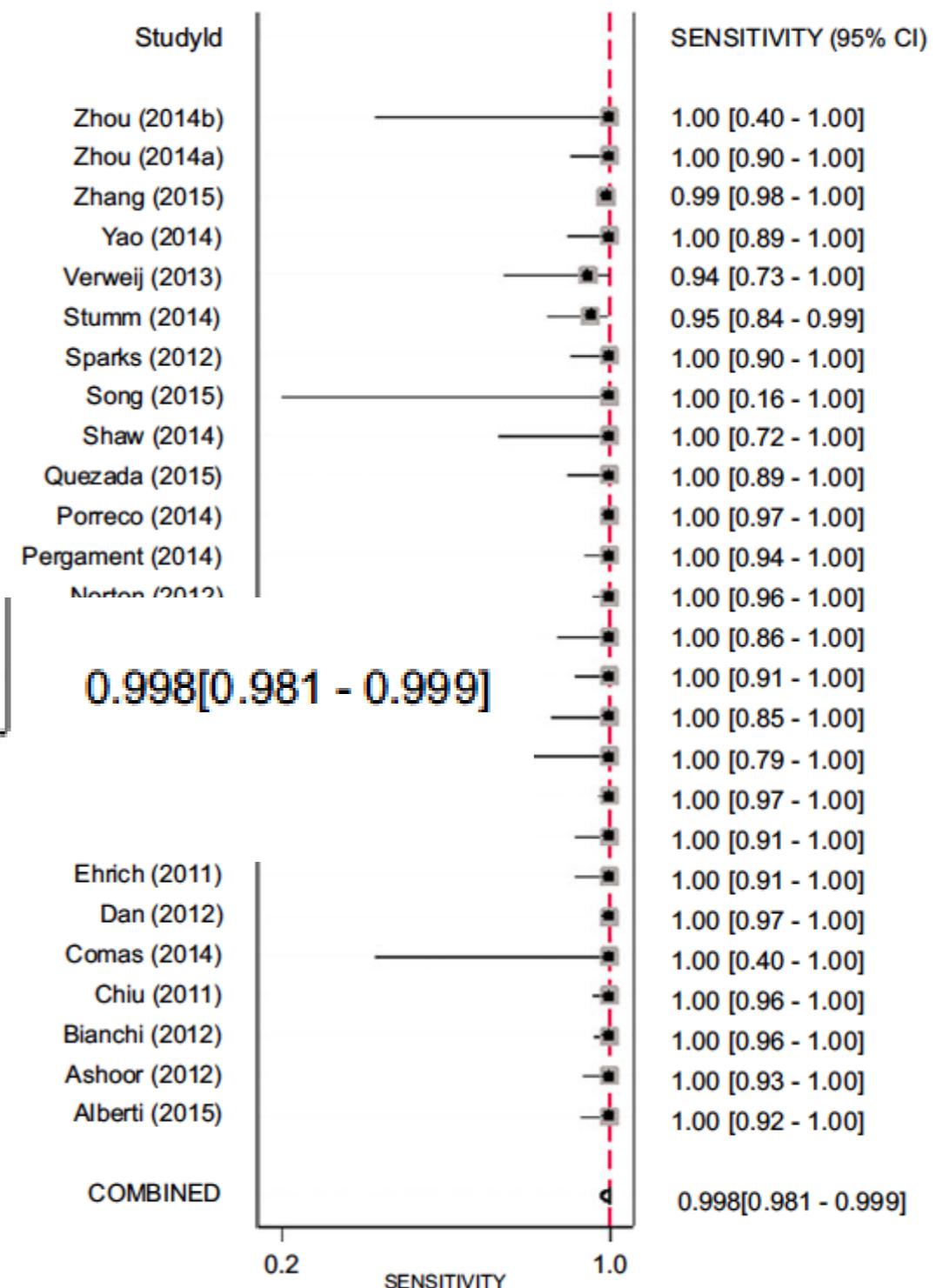
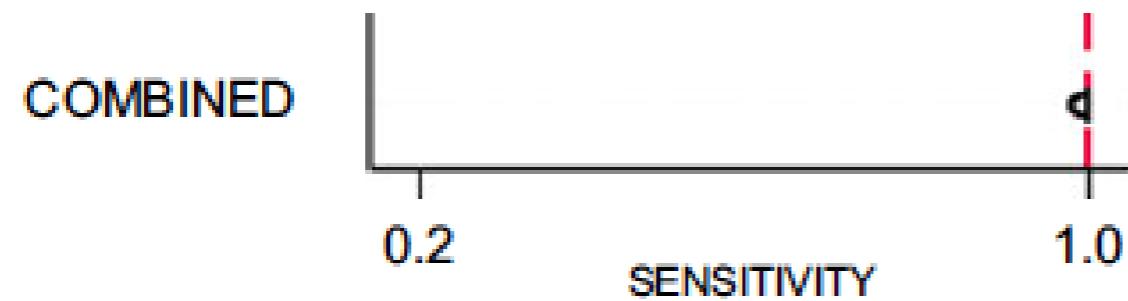
M. M. GIL^{1,2,3} V. ACCURTI¹, B. SANTACRUZ², M. N. PLAN⁴ and K. H. NICOLAIDES¹

¹Fetal Medicine Research Institute, King's College Hospital, London, UK; ²Obstetrics and Gynecology Department, Torrejon University Hospital, Torrejon de Ardoz, Madrid, Spain; ³Obstetrics and Gynecology Department, Universidad Francisco de Vitoria, Pozuelo de Alarcón, Madrid, Spain; ⁴Clinical Biostatistics Unit, Ramón y Cajal Hospital (IRYCIS), CIBER Epidemiology and Public Health (CIBERESP), Madrid, Spain

Quezada (2015) ²⁶	CSS	10 (10–11)	T21,T18,T13	53/2905 (1.8)
Gil (2016) ³⁰	CSS	12 (11–13)	T21	1/54 (1.9)
Mnyani (2016) ³³	SNP	14 (13–21)	T13	2/82 (2.4)
Qi (2016) ³⁸	MPSS	19 (11–30)	T21,T18,T13	4/2828 (0.1)
Ma (2017) ⁴⁰	MPSS	19 (—)	T21,T18,T13	15/10 594 (0.1)

Analysis of cell-free fetal DNA in maternal blood for detection of trisomy 21, 18 and 13 in a general pregnant population and in a high risk population – a systematic review and meta-analysis

ERIK IWARSSON^{1,*}, BO JACOBSSON^{2,3,*}, JESSICA DAGERHAMN⁴, THOMAS DAVIDSON^{4,5},
EDUARDO BERNABÉ⁶ & MARIANNE HEIBERT ARNLIND^{4,7}





Analysis of cell-free DNA in maternal blood in screening for fetal aneuploidies: updated meta-analysis

M. M. GIL*, M. S. QUEZADA*, R. REVELLO*, R. AKOLEKAR*† and K. H. NICOLAIDES*†

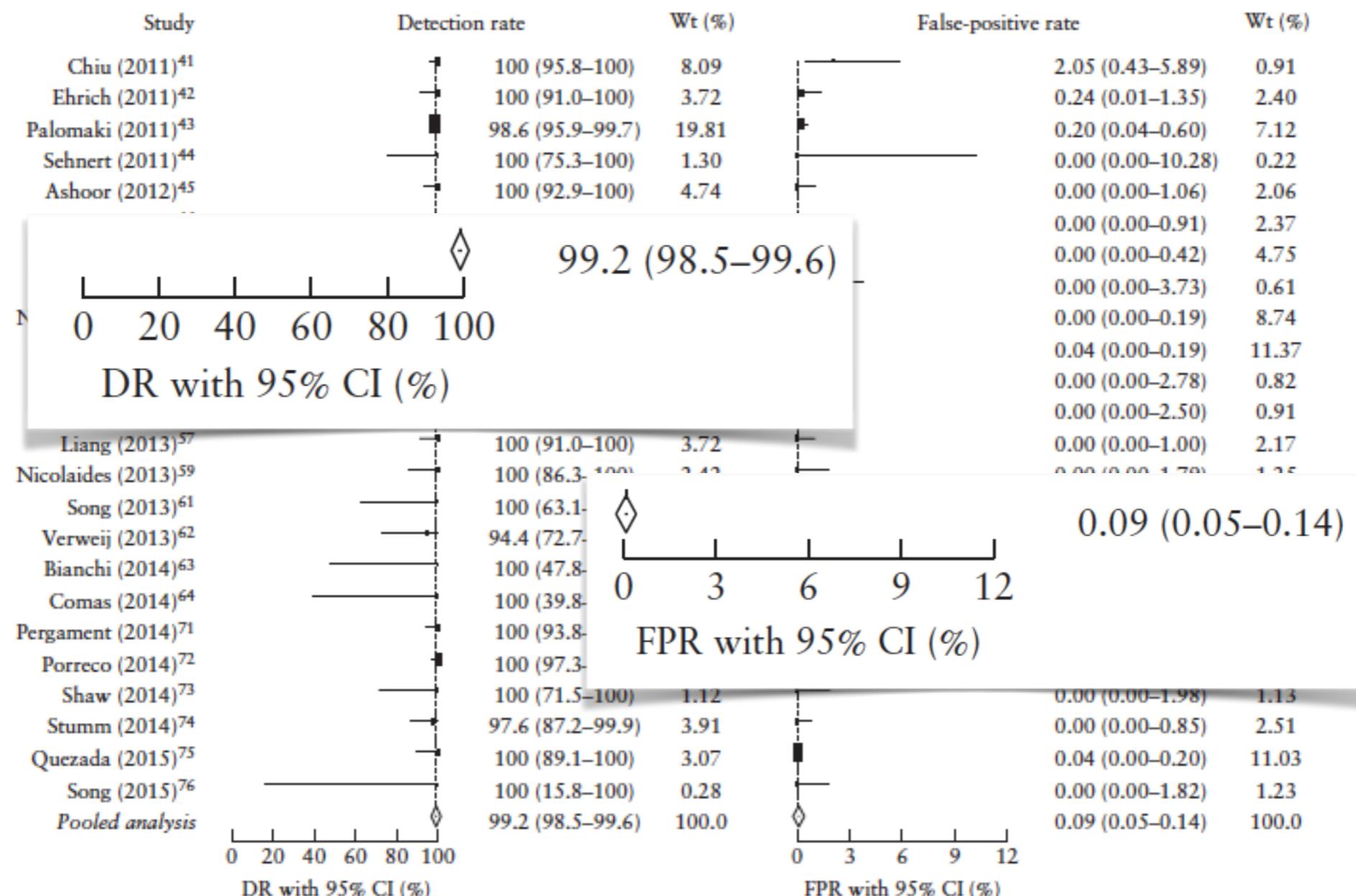


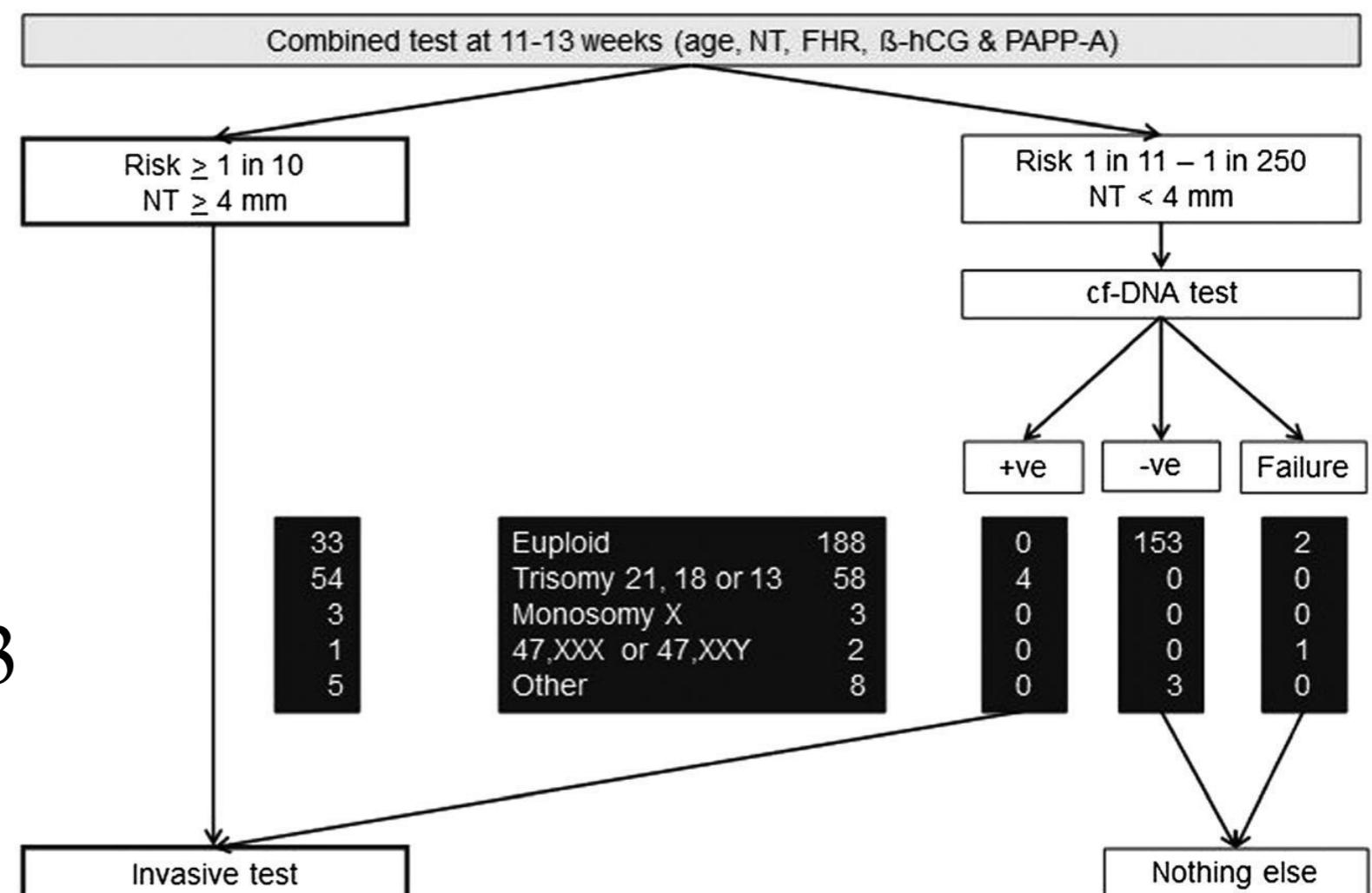
Figure 4 Forest plots of detection rates (DR) and false-positive rates (FPR) with 95% CIs and weighted pooled summary statistics using the random-effects model in assessing cell-free DNA analysis in screening for trisomy 21 in singleton pregnancy. Only the first author of each study is given.

Cell-free DNA testing in the maternal blood in high-risk pregnancies after first-trimester combined screening

Nicola Persico^{1*}, Simona Boito¹, Benedetta Ischia¹, Adalgisa Cordisco², Valentina De Robertis³, Isabella Fabietti¹, Enrico Periti², Paolo Volpe³, Luigi Fedele¹ and Georgios Rembouskos³

259 singleton pregnancies

estimated risk for trisomies 21, 18 or 13
 ≥ 1 in 250



Cell-free DNA testing in the maternal blood in high-risk pregnancies after first-trimester combined screening

Nicola Persico^{1*}, Simona Boito¹, Benedetta Ischia¹, Adalgisa Cordisco², Valentina De Robertis³, Isabella Fabietti¹, Enrico Periti², Paolo Volpe³, Luigi Fedele¹ and Georgios Rembouskos³

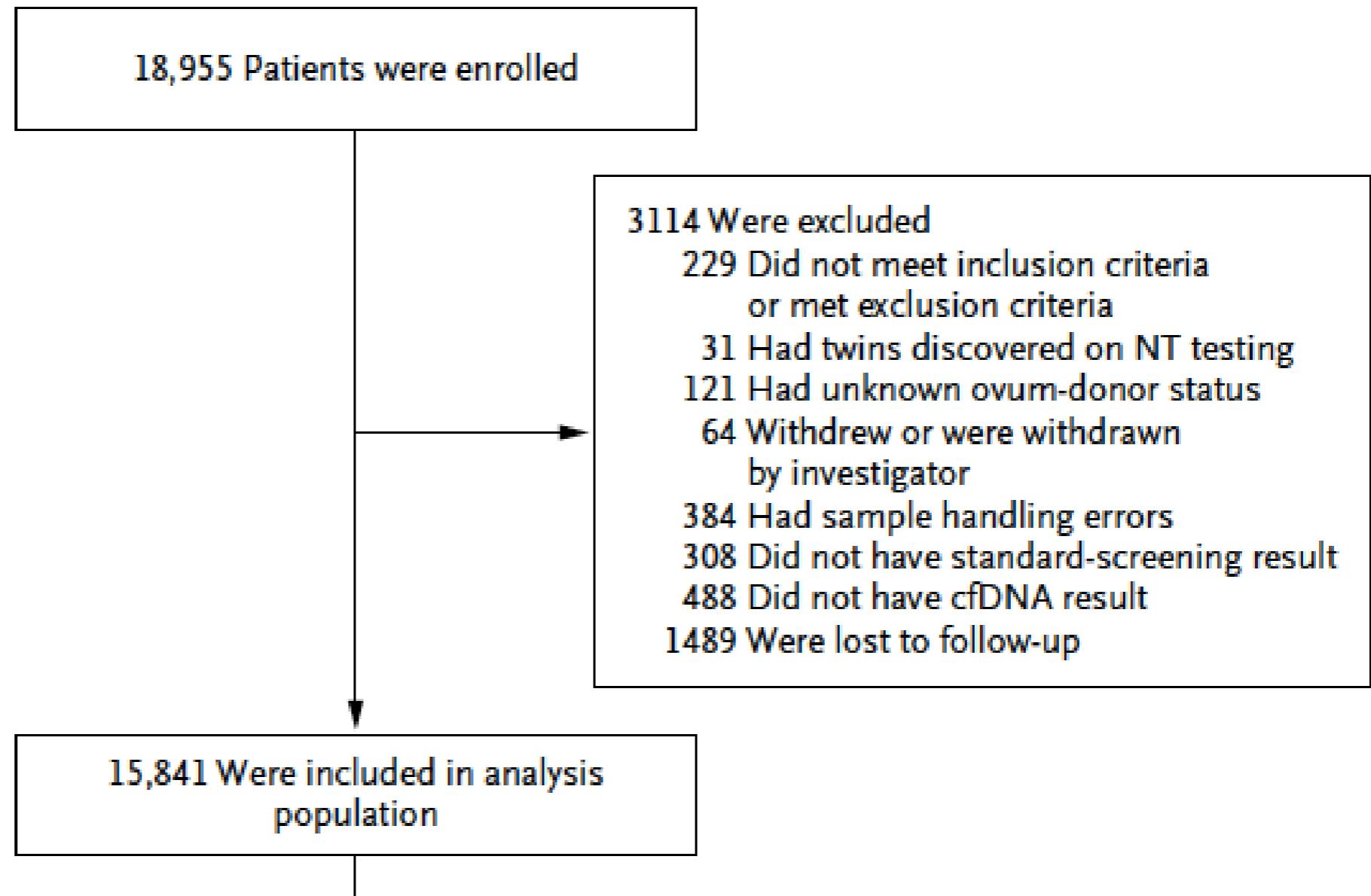
Table 1 Results from cfDNA testing according to fetal karyotype

Fetal karyotype	<i>n</i>	Cell free DNA test			Screen -ve
		No result	Result	Screen +ve	
Euploid	188	4 (2.1)	184	—	184 (100)
Trisomy 21	37	1 (2.7)	36	35 (97.2)	1 (2.8)
Trisomy 18	15	2 (13.3)	13	13 (100)	—
Trisomy 13	6	1 (16.7)	5	5 (100)	—
Monosomy X	3	0 (0.0)	3	2 (66.7)	1 (33.3)
47,XXX or 47, XXY	2	1 (50.0)	1	1 (100)	—
Other	8	1 (12.5)	7	—	7 (100)
Total	259	10 (3.9)	249	56 (22.5)	193 (77.5)

Cell-free DNA Analysis for Noninvasive Examination of Trisomy

Mary E. Norton, M.D., Bo Jacobsson, M.D., Ph.D., Geeta K. Swamy, M.D., Louise C. Laurent, M.D., Ph.D.,
Angela C. Ranzini, M.D., Herb Brar, M.D., Mark W. Tomlinson, M.D., Leonardo Pereira, M.D., M.C.R.,
Jean L. Spitz, M.P.H., Desiree Hollomon, M.S.N., M.P.H., Howard Cuckle, D.Phil., M.B.A.,
Thomas J. Musci, M.D., and Ronald J. Wapner, M.D.

CSS, chromosome-specific sequencing



Cell-free DNA Analysis for Noninvasive Examination of Trisomy

Mary E. Norton, M.D., Bo Jacobsson, M.D., Ph.D., Geeta K. Swamy, M.D., Louise C. Laurent, M.D., Ph.D.,
Angela C. Ranzini, M.D., Herb Brar, M.D., Mark W. Tomlinson, M.D., Leonardo Pereira, M.D., M.C.R.,
Jean L. Spitz, M.P.H., Desree Hollomon, M.S.N., M.P.H., Howard Cuckle, D.Phil., M.B.A.,
Thomas J. Musci, M.D., and Ronald J. Wapner, M.D.

CSS, chromosome-specific sequencing

Trisomy 21 =

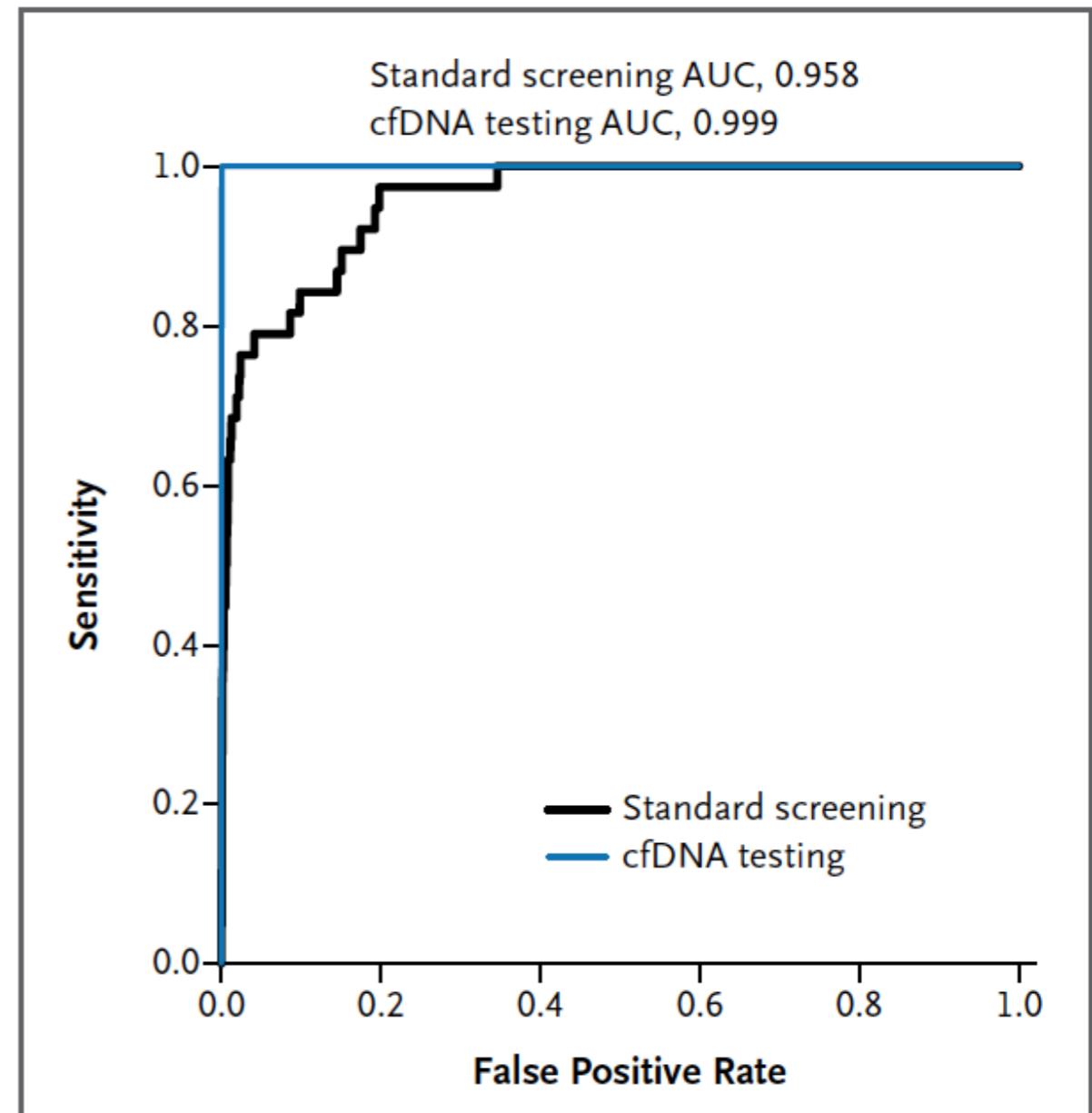
CSS: 38 of 38 women (100%; 95%CI, 90.7 to 100)

Standard: 30 of 38 women (78.9%; 95%CI, 62.7 to 90.4)

False positive rates

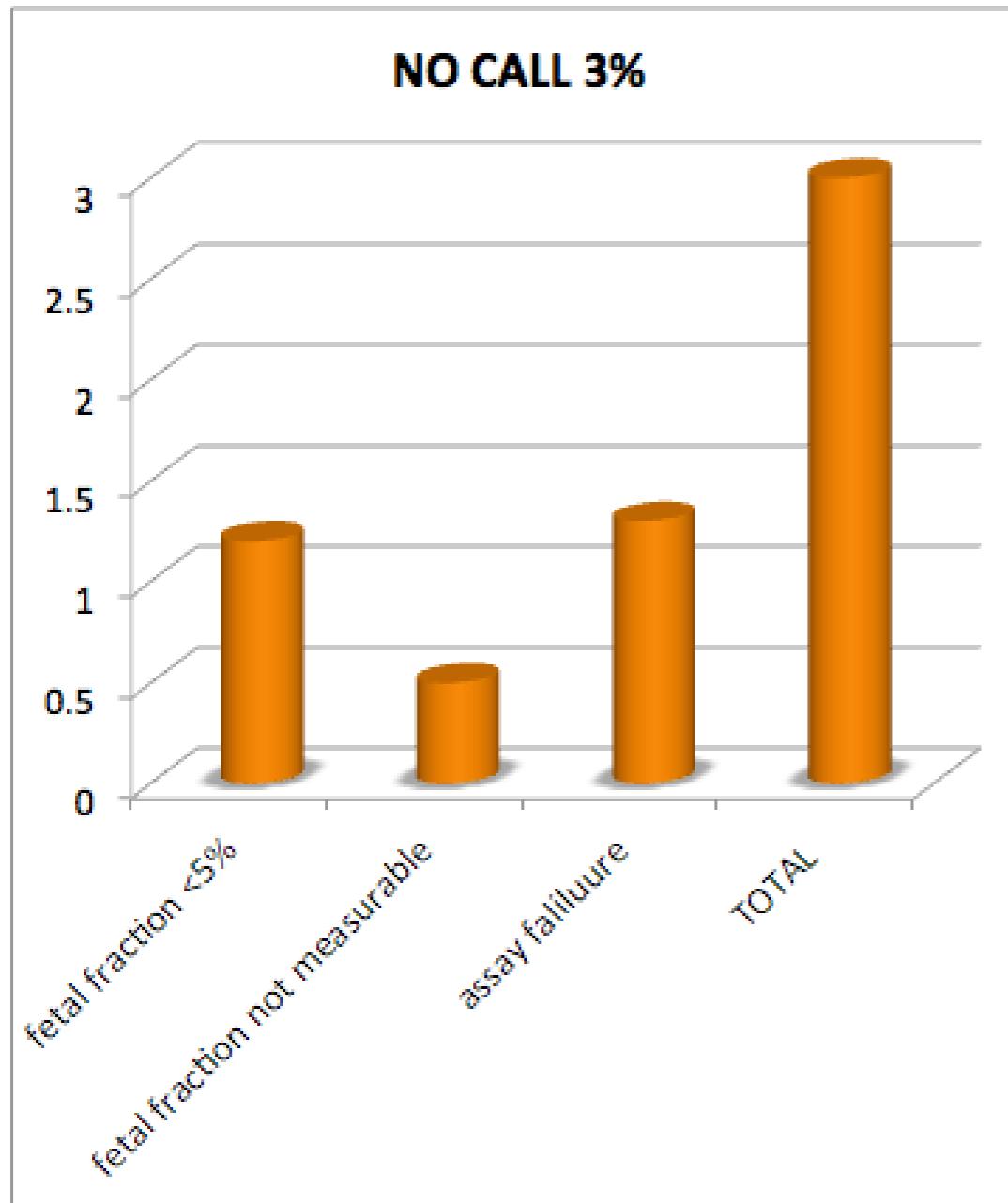
CSS 0.06%

Standard 5.4%



Cell-free DNA Analysis for Noninvasive Examination of Trisomy

Mary E. Norton, M.D., Bo Jacobsson, M.D., Ph.D., Geeta K. Swamy, M.D., Louise C. Laurent, M.D., Ph.D.,
Angela C. Ranzini, M.D., Herb Brar, M.D., Mark W. Tomlinson, M.D., Leonardo Pereira, M.D., M.C.R.,
Jean L. Spitz, M.P.H., Desiree Hollomon, M.S.N., M.P.H., Howard Cuckle, D.Phil., M.B.A.,
Thomas J. Musci, M.D., and Ronald J. Wapner, M.D.



CSS, chromosome- specific sequencing

In the group with no results on cfDNA testing,
there were:
3 trisomy 21,
1 trisomy 18,
2 trisomy 13,
4 triploidy,
1 trisomy 16 mosaic,
1 deletion 11p,
1 w structurally abnormal chromosome

The prevalence of aneuploidy in this group 2.7%

The prevalence of aneuploidy in the overall cohort 0.4%

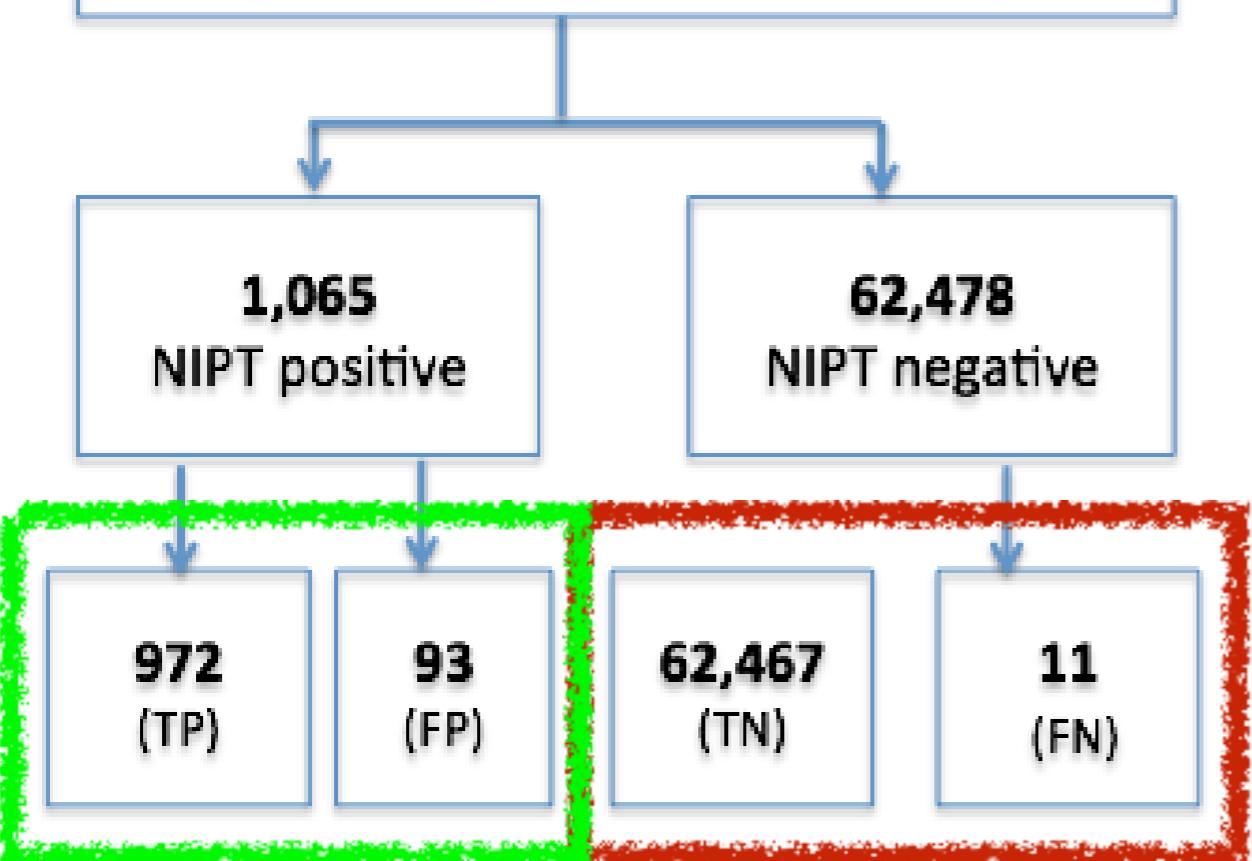
massive parallel genome sequencing

performance for aneuploidy

178,719 NIPT tests by Aug 2013

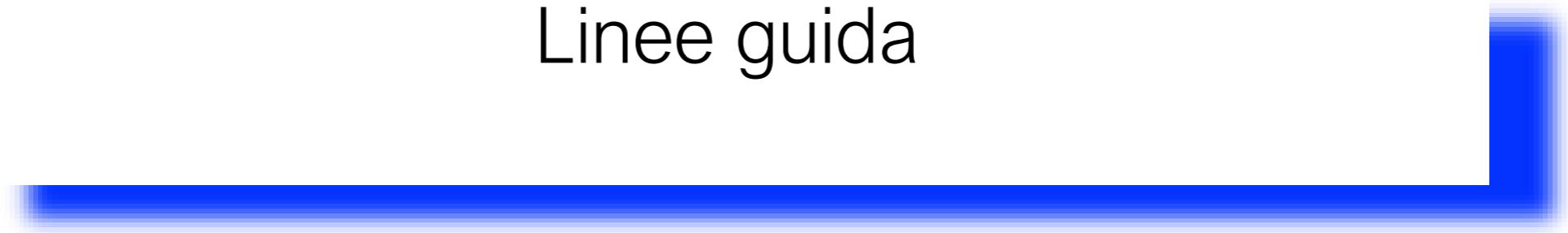
- Resampling/retesting: 2.8%;
- No call rate: 0.069%;

63,543 singleton pregnancies
passed EDC + NIPT calls + follow-up
(fetal karyotypes or outcomes)



T21	
Specificity	99.95%
Sensitivity	99.22%
PPV	95.73%
NPV	99.99%
T18	
Specificity	99.95%
Sensitivity	97.34%
PPV	84.72%
NPV	99.99%
T13 with 95% CI	
Specificity	99.96% (99.95-99.97%)
Sensitivity	100.00% (87.2-100%)
PPV	51.92% (38.3-65.5%)
NPV	100.00% (99.99-100%)

Linee guida





COMMITTEE OPINION

Number 640, September 2015

(This Committee Opinion Replaces Committee Opinion Number 545)

Committee on Genetics

Society for Maternal-Fetal Medicine

This document reflects emerging clinical and scientific advances as of the date issued and is subject to change. The information should not be construed as dictating an exclusive course of treatment or procedure to be followed.

Recommendations

- Cell-free DNA screening does not assess risk of fetal anomalies . Patients who are undergoing cell-free DNA screening should be offered ultrasound evaluation for risk assessment.

Women whose results are not reported, indeterminate, or uninterpretable (a “no call” test result) from cell-free DNA screening should receive further genetic counseling and be offered comprehensive ultrasound evaluation and diagnostic testing because of an increased risk of aneuploidy.

a diagnostic test should be recommended for a patient who has a positive cell-free DNA test result.

-



The American College of
Obstetricians and Gynecologists
WOMEN'S HEALTH CARE PHYSICIANS



Society for
Maternal-Fetal
Medicine

COMMITTEE OPINION

Number 640, September 2015

(This Committee Opinion Replaces Committee Opinion Number 545)

Committee on Genetics

Society for Maternal-Fetal Medicine

This document reflects emerging clinical and scientific advances as of the date issued and is subject to change. The information should not be construed as dictating an exclusive course of treatment or procedure to be followed.

Recommendations

- A discussion of the risks, benefits, and alternatives ...should occur with all patients.
- Given the performance of conventional screening methods, the limitations of cell-free DNA screening performance, and the limited data on cost-effectiveness in the low-risk obstetric population, **conventional screening methods remain the most appropriate choice for first-line screening for most women in the general obstetric population.**
- Although any patient may choose cell-free DNA analysis as a screening strategy for common aneuploidies regardless of her risk status, the patient choosing this testing should understand the limitations and benefits of this screening paradigm...
- **The cell-free DNA test will screen for only the common trisomies and, if requested, sex chromosome composition.**
-



Non-invasive Prenatal Testing for Chromosomal Abnormality using Maternal Plasma DNA

In April 2013, the **International Society for Prenatal Diagnosis** published a position statement which concluded that maternal plasma DNA testing may be considered **in women classified as high-risk...** by previous screens or assessed as high-risk maternal age, the presence of an ultrasound finding suggestive of trisomy 21, 18 or 13,



7.2.1 Contingent (step-wise) screening

The use of maternal plasma MPS within a contingent approach would mean continuing combined testing and then to offer the maternal plasma DNA test to the subgroup (perhaps 20%) identified as higher risk.

The cut-off to offer maternal plasma MPS could be set to manage the numbers and resulting costs and this approach could significantly reduce the invasive procedure numbers (by greatly reducing the false-positive rate).

Possible sources of error

5.1 Early gestational age

5.2 Maternal obesity

5.3 Multiple pregnancies

monochorionic (and so monozygotic), both fetuses will be affected or unaffected. Since the amount ofcffDNA is approximately double that of a singleton pregnancy,⁵¹cffDNA aneuploidy testing will not only be possible but probably more effective than in singletons.

dichorionic, and so may be discordant, maternal plasma DNA testing would, in theory, not be as straightforward. The complexity introduced by twin pregnancies suggests that, prior tocffDNA testing, a good quality ultrasound scan would be a valuable first step in all pregnancies, to detect empty pregnancy sacs, for example, with fetal medicine counselling when one is suspected.

5.4 Placental mosaicism.

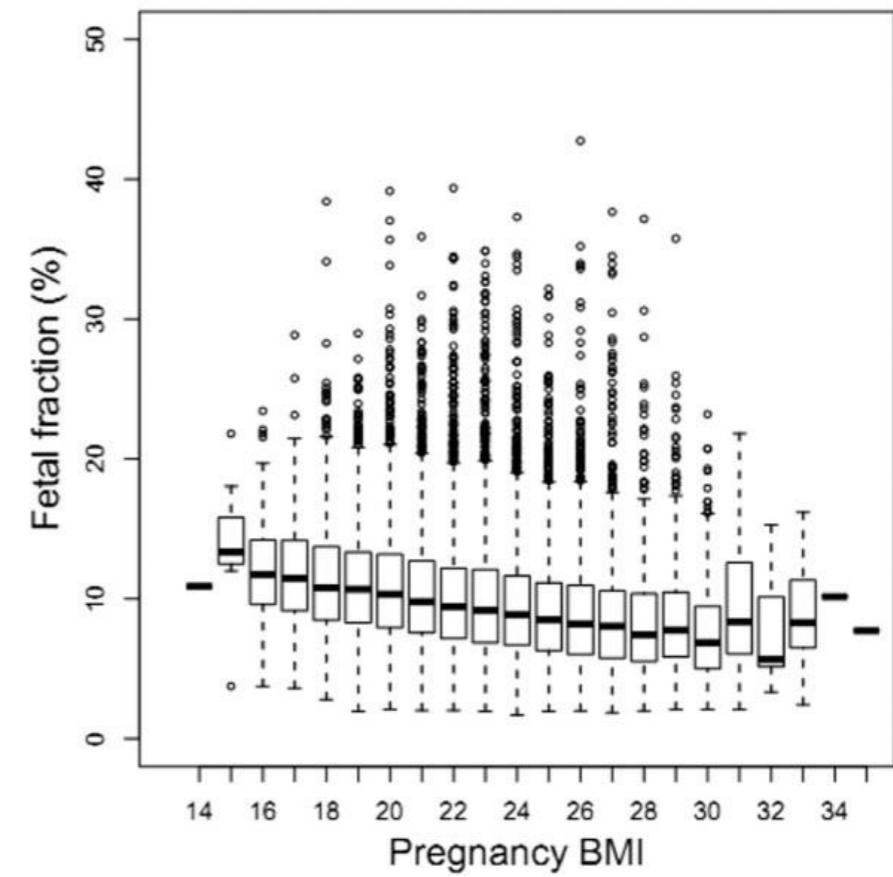
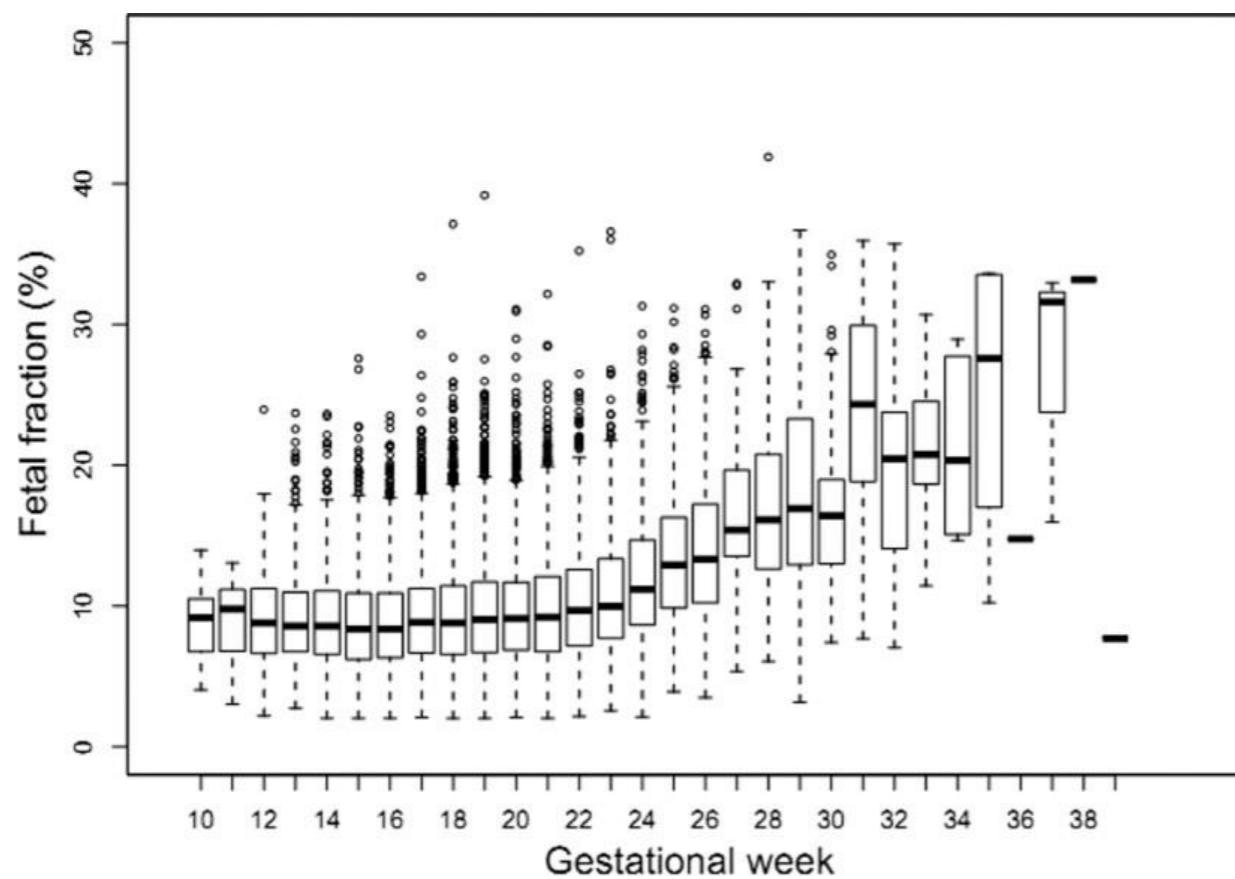
approximately 1% of CVS samples, a phenomenon often called ‘confined placental mosaicism’.

5.5 Maternal conditions

Maternal chromosomal abnormalities, including mosaicism or malignant disease, could be very rare causes of discordant results.

Effects of Maternal and Fetal Characteristics on Cell-Free Fetal DNA Fraction in Maternal Plasma

Reproductive Sciences
1-7
© The Author(s) 2015
Reprints and permission:
sagepub.com/journalsPermissions.nav
DOI: 10.1177/1933719115584445
rs.sagepub.com



I Centri che erogano il test devono:

- a) avere competenze nella diagnosi ecografica;
- b) avere competenza nella diagnosi prenatale;
- c) essere in grado di offrire la consulenza pre-test e post-test;
- d) essere collegati con il laboratorio di cui al punto 8.

I Laboratori che eseguono il test devono:

- a) essere certificati;
- b) partecipare a programmi di controllo della qualità,
- c) essere dotati di personale con competenza specifica nelle tecniche di NGS.

Nel caso in cui i laboratori sviluppino protocolli originali, è necessario che le tecniche e le procedure bioinformatiche siano rese pubbliche e disponibili alla validazione scientifica.



Per l'analisi delle aneuploidie mediante NIPT si utilizzano tre principali tecniche basate sulle tecniche di sequenziamento di seconda generazione (Next Generation Sequencing - NGS):

- 1) MPSS dell'intero genoma;
- 2) CSS di specifiche regioni;
- 3) SNP, cioè polimorfismi di singoli nucleotidi.



Ministero della Salute
Consiglio Superiore di Sanità

Gli studi relativi all'uso del NIPT come test di screening sono stati promossi e realizzati dalle aziende che, a partire dal 2012, hanno avviato la sua commercializzazione, con finalità cliniche:

**Sequenom (MPSS-MaterniT21), Verinata (MPSS- Verifi), BGI (MPSS - NiftY G-TEST). (CSS- Harmony),
Natera (SNP - Panorama),**



Ministero della Salute

Consiglio Superiore di Sanità

Il NIPT deve essere collegato e preceduto da un accurato controllo ecografico dopo l'XI settimana, effettuato da operatori accreditati nell'esame delle XI-XIV settimane.

Nel caso in cui i dati ecografici suggeriscano un aumento del rischio di patologia cromosomica nel feto, deve essere valutata l'opportunità di eseguire direttamente una diagnosi prenatale invasiva per lo studio del cariotipo fetale, integrato eventualmente da altre tecniche

analisi costi benefici su un modello locale

Fonte: CEDAP

GRAVIDE MILANO 2012			<i>Dati Nicolaides (in corsivo)</i>
Età materna	N. casi	N. Down attesi	Down attesi
<20	206	0,1	1/1600
20	147	0,1	1/1527
21	239	0,2	1/1483
22	259	0,2	1/1439
23	348	0,2	1/1395
24	417	0,3	1/1360
25	472	0,3	1/1352
26	559	0,5	1/1237
27	689	0,6	1/1123
28	801	0,8	1/1009
29	918	1,0	1/903
30	1146	1,3	1/895
31	1332	1,7	1/776
32	1421	2,2	1/659
33	1439	2,6	1/547
34	1600	3,6	1/446
< 35 anni	11.993	16	

GRAVIDE MILANO 2012			<i>Dati Nicolaides (in corsivo)</i>
Età materna	N. casi	N. Down attesi	Down attesi
35	1676	4,7	1/356
36	1488	5,3	1/280
37	1575	7,2	1/218
38	1390	8,3	1/167
39	1192	9,3	1/128
40	972	10,0	1/97
41	716	9,8	1/73
42	421	7,7	1/55
43	305	7,4	1/41
44	180	6,0	1/30
45	112	4,9	1/23
46	60	3,5	1/17
47	33	2,1	1/16
48	14	0,9	1/15
49	15	1,5	1/10
≥50	12	1,2	1/10
≥ 35 anni	10.161	90	

TEST	Popolazione gravide Milano 2012	Costo totale	N. Down diagnosticati	N. Down non diagnosticati	N. Aborti Previsti
Età materna	10.161 (≥ 35)	€4.064.400	90/106	16/106	51-102
Test combinato	22.154	€2.658.480	95/106	11/106	6-12
Età materna + Test combinato	10.161 (≥ 35) 11.993 (< 35)	€5.722.880	104/106	2/106	54-108
Test del DNA fetale	22.154	€9.969.300	105/106	1/106	0,5-1
Test combinato(*) + Test del DNA fetale	22.154	€4.430.800	104/106	2/106	0,5-1
NOTE Costo test combinato 100€-Costo amniocentesi 400€-Costo test DNA fetale 400€-N. Down atteso sulla popolazione gravide Milano 2012 106-Rischio aborto da amniocentesi 0,5/1%					
(*) rischio intermedio 1/100-1/2000 = 20%					