

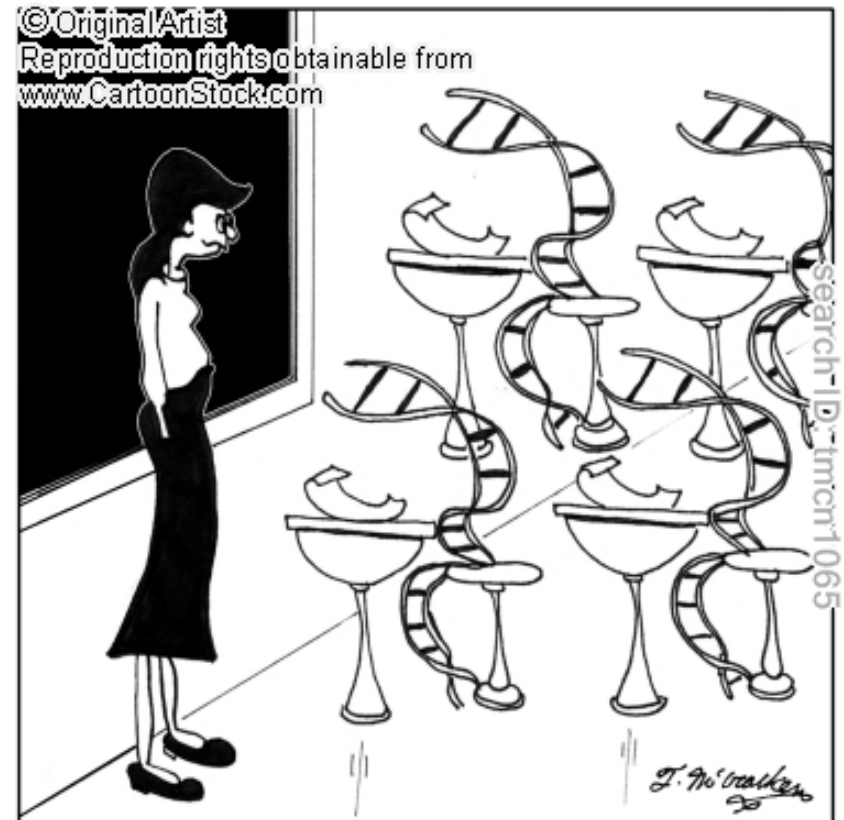


Updates in Genetic Screening

**Pregnancy Care ECHO
December 18th**

Amber Mathiesen, M.S., L.C.G.C

- Screening options:
 - cfDNA testing
 - ACOG statement
- Carrier testing:
 - Ancestry based
 - Enhancements to fragile X and SMA
 - Expanded carrier testing



DNA Testing.

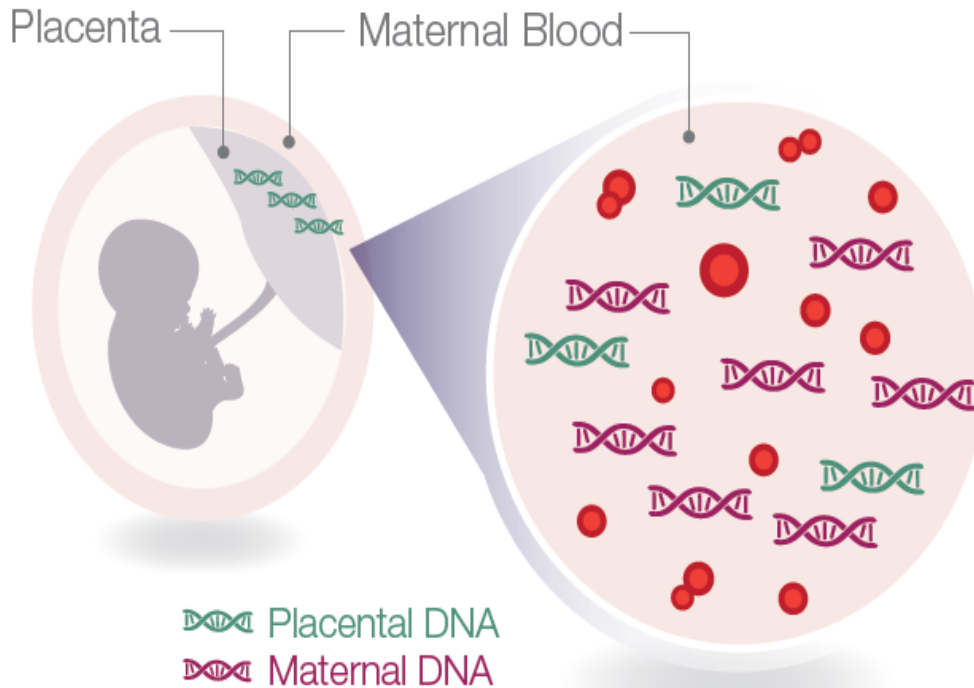


Advancements in Prenatal Screening

1960's	1980s	1988	1996	1997	2011-2012
Maternal Age	MSAFP	Triple Screen	Quad Screen	FTS NT/Serum	cfDNA SNP
Detection Rate					
27%	36%	60-74%	70-81%	80-95%	92->99%
Chromosome Abnormalities Screened					
All	T21	T21 T18	T21 T18	T21 T18 T13*	T21 T18 T13 SCA Triploidy* Microdeletions*

* some

Cell-free DNA (cfDNA)

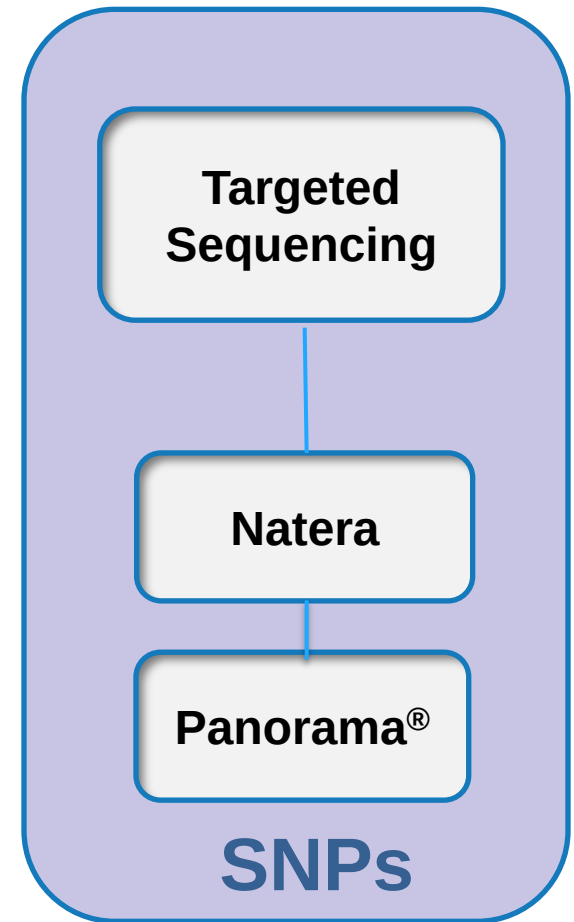
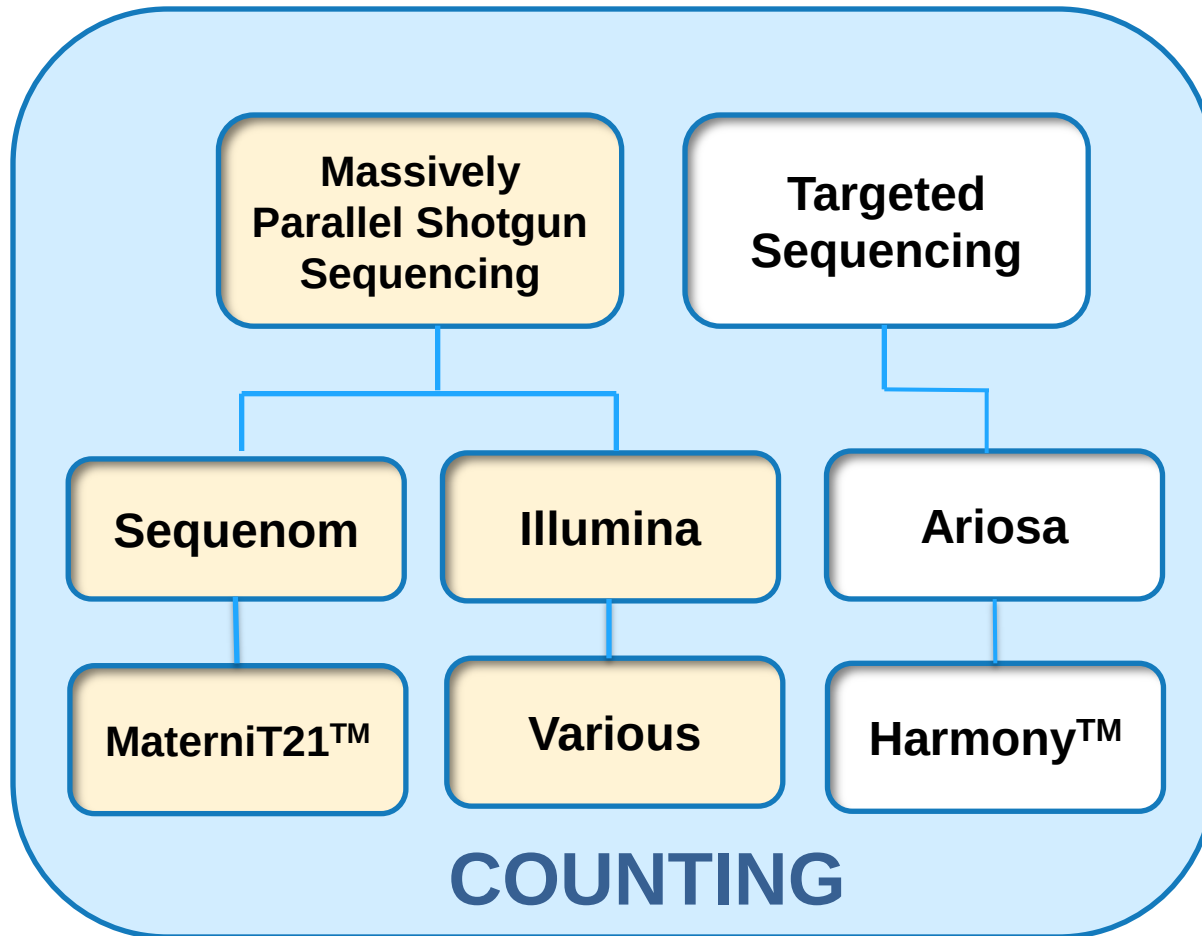


cfDNA comes from apoptotic cells derived from:

- Maternal Circulation
 - Adipocytes
 - White Blood Cells
- Fetal
 - Placental cells (trophoblasts) in the maternal circulation



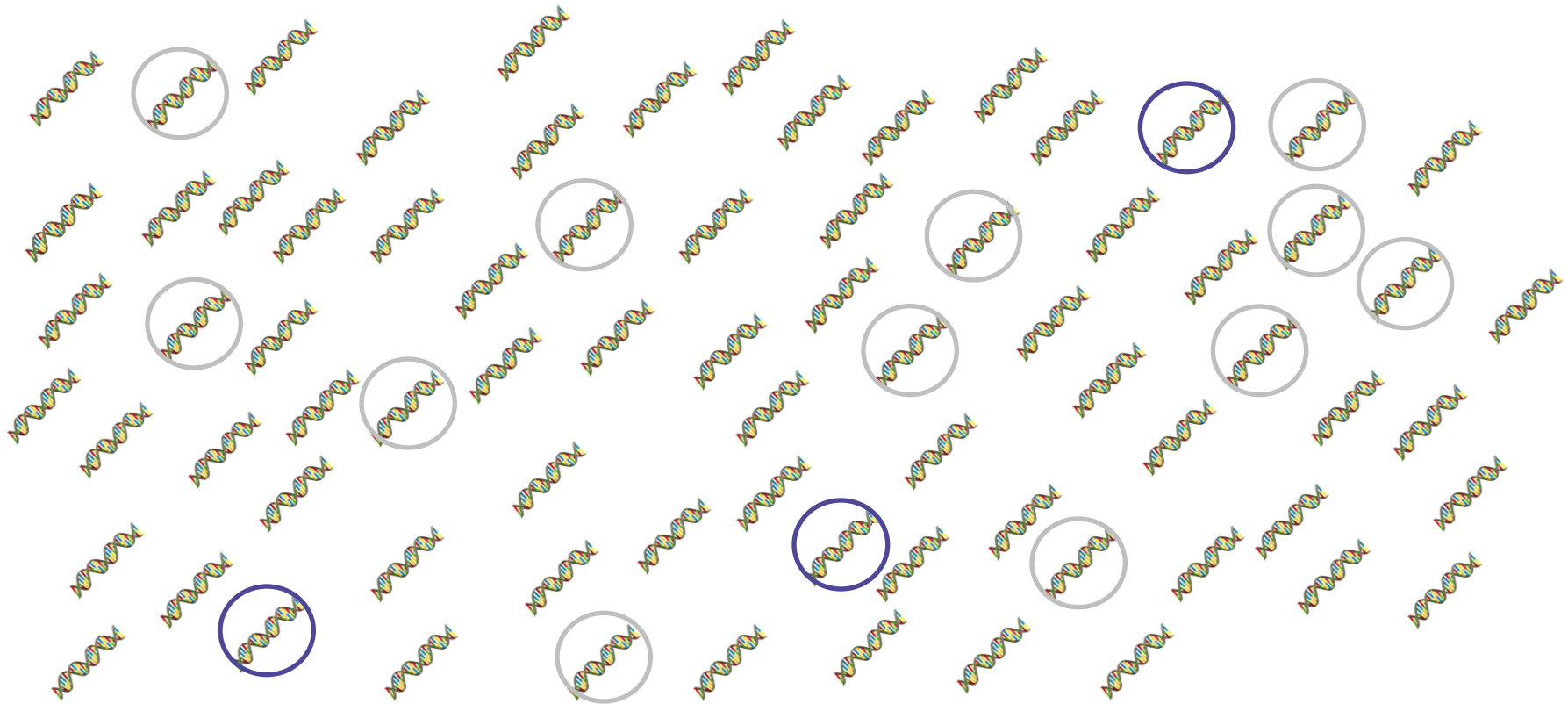
Differentiating Methodologies



Counting

Chromosome
21

Chromosome
3



Counting

Chromosome 21

Chromosome 3

Expected Amount:

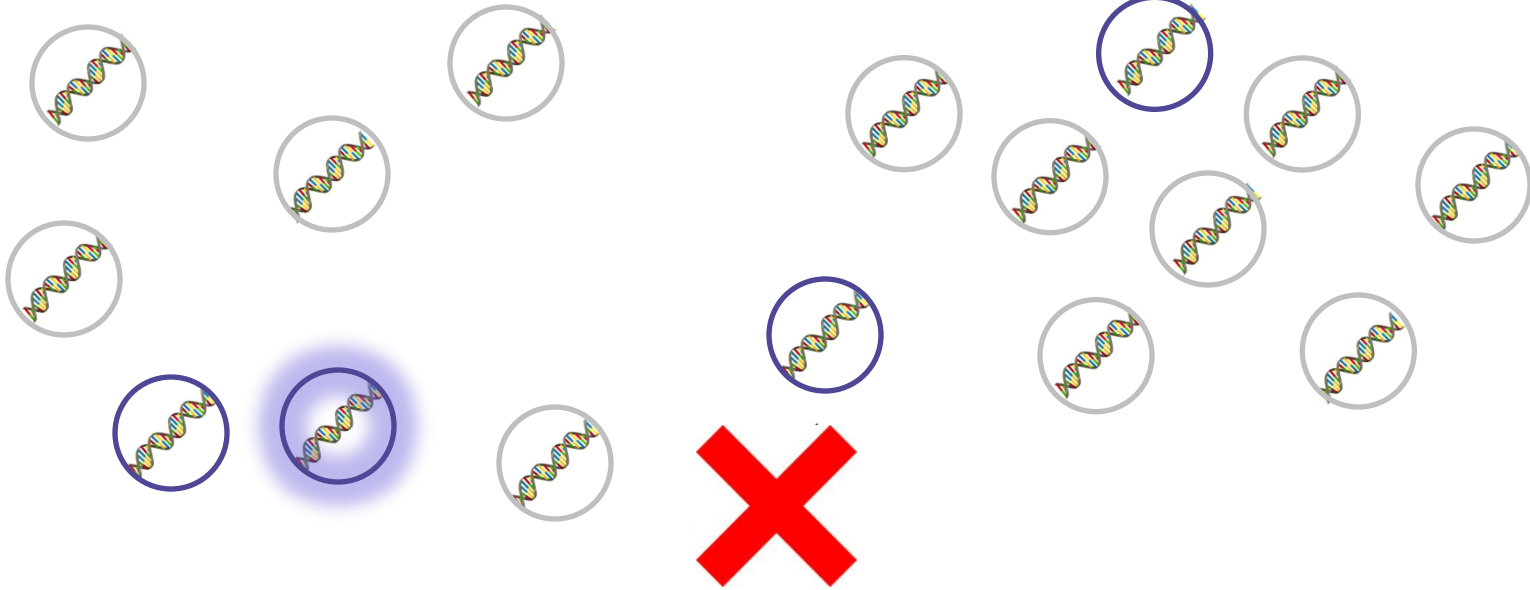
20%

80%

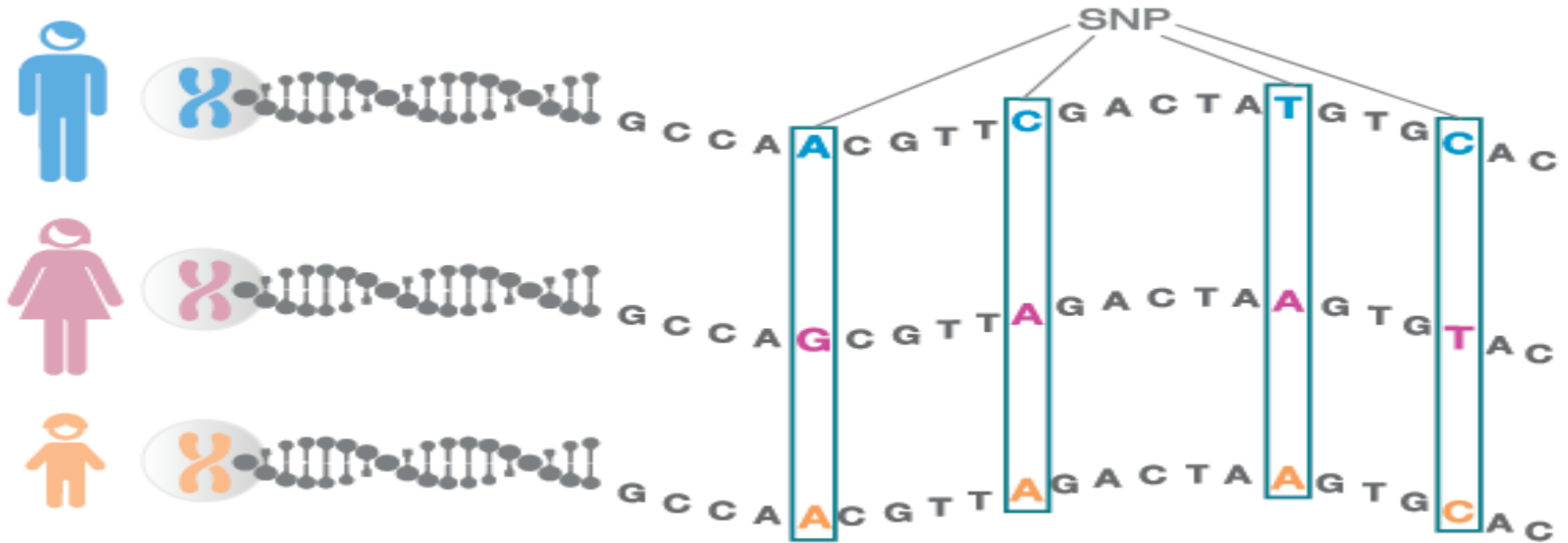
Observed Amount:

25%

75%



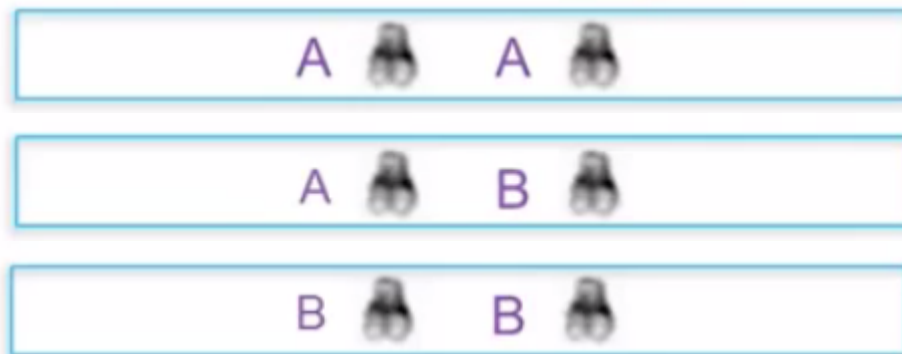
SNP = Single Nucleotide Polymorphism



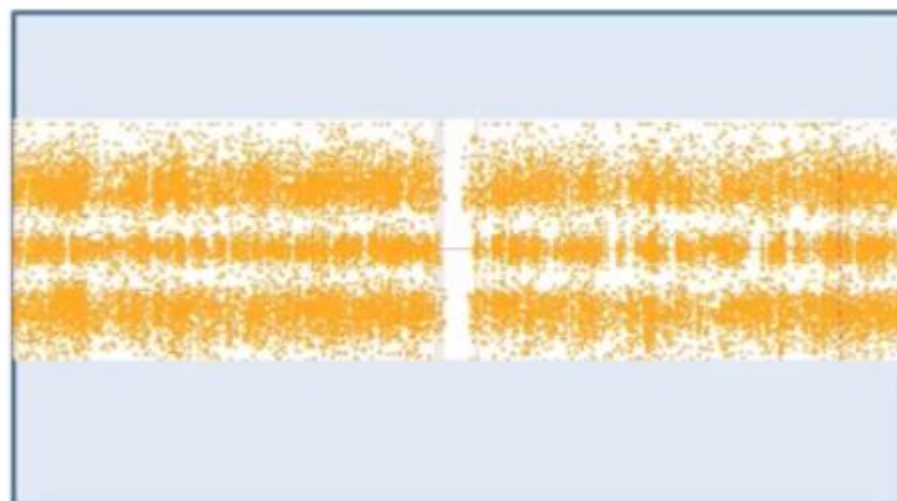
base pair (nucleotide) - A, T, C, or G – is changed.

- These are normal genetic changes that occur in every person

SNP



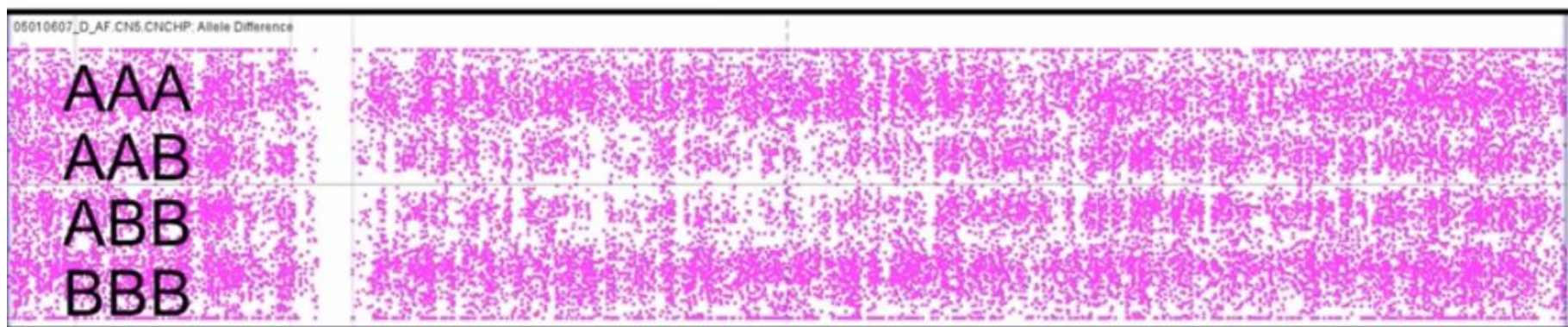
AA
AB
BB



Disomy
2 Chromosomes



SNP



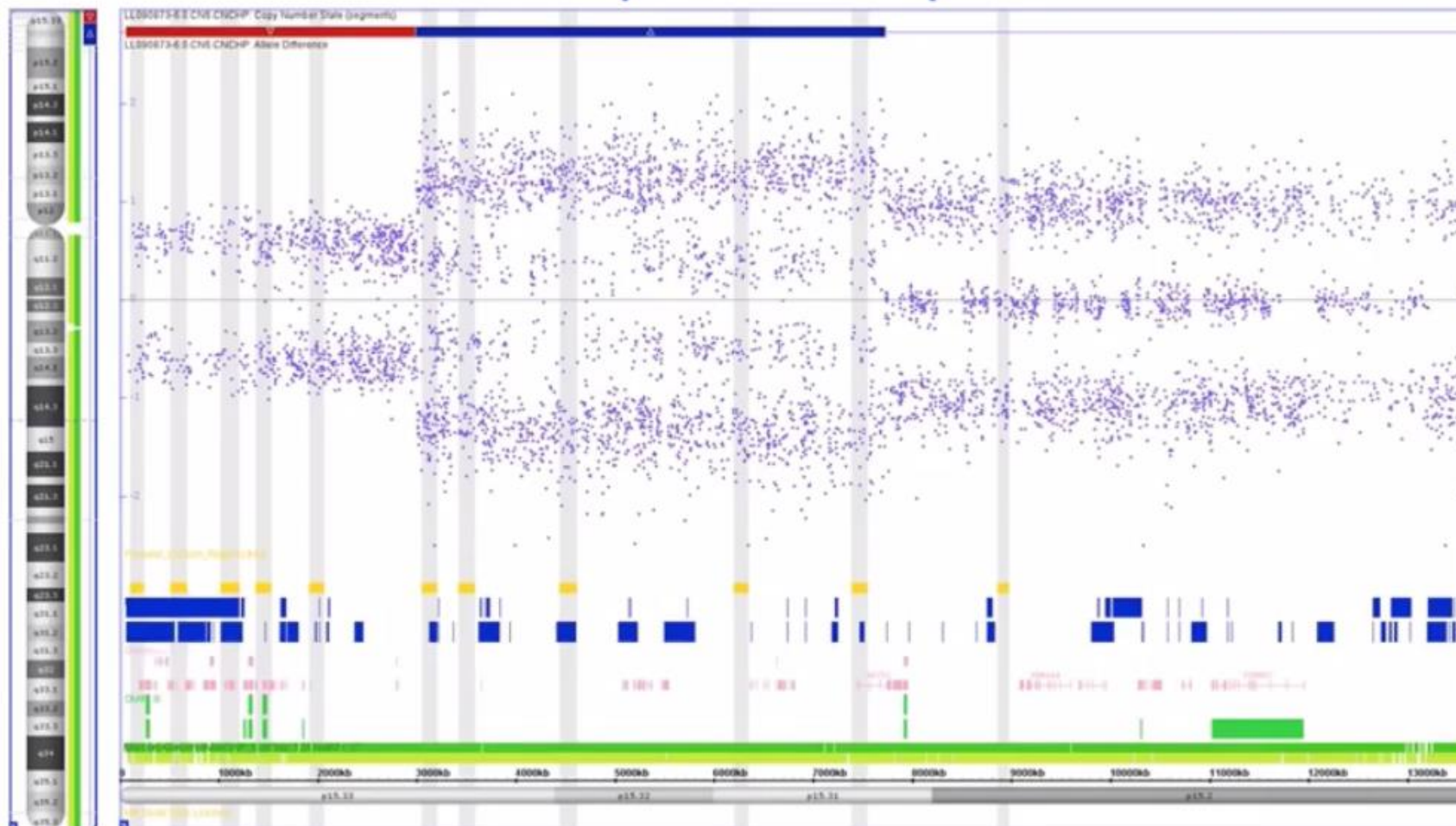
Trisomy
3 Chromosomes

SNP

Deletion

Duplication

Disomy



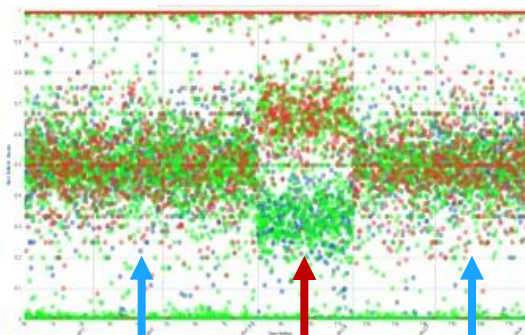
Maternal
blood



Maternal +
Fetal
cfDNA



SNP Sequencing of Maternal + Fetal Genotype



Disomy

Trisomy

Disomy

Advanced Bioinformatics



Report

Patient Information

Patient Name	Jane Doe
DOB	12/15/1978
Maternal Age at EDD	37
Estimated Weight	155 pounds (70 kg)
Height	5'6"
Parity	0/0/0/3
Previous Screen	None
Assessment ID	ABCDEF123
Referring Doctor	Dr. Smith MD
Case #	12345678

Provider Information

Ordering Physician	Dr. Matthew Goldstein, MD (MCHS001)
OB/GYN Information	Maternal ID: M123456789, Maternal Signature: [Signature]
Specimen Date	12/15/2023
Specimen Received	12/15/2023
Lab Name	LabCorp
Referral Clinic	Obstetrics

Test Results Summary

Test	Result	Reference Range	Notes
1st Trimester Screening	0.83%	0.05% - 3.00%	Low risk
2nd Trimester Screening	0.05%	0.02% - 0.10%	Low risk
3rd Trimester Screening	0.02%	0.01% - 0.05%	Low risk
4th Trimester Screening	0.01%	0.00% - 0.02%	Low risk
5th Trimester Screening	0.00%	0.00% - 0.01%	Low risk
6th Trimester Screening	0.00%	0.00% - 0.00%	Low risk
7th Trimester Screening	0.00%	0.00% - 0.00%	Low risk
8th Trimester Screening	0.00%	0.00% - 0.00%	Low risk
9th Trimester Screening	0.00%	0.00% - 0.00%	Low risk
10th Trimester Screening	0.00%	0.00% - 0.00%	Low risk

Summary of Findings: The patient is a 37-year-old female with a low risk of chromosomal abnormalities. The results of the 1st, 2nd, 3rd, 4th, 5th, 6th, 7th, 8th, 9th, and 10th trimester screenings are all within the reference range, indicating a low risk of chromosomal abnormalities. The patient is currently at 37 weeks gestation and is in good health. The results of the 1st trimester screening are consistent with the results of the 2nd, 3rd, 4th, 5th, 6th, 7th, 8th, 9th, and 10th trimester screenings. The patient is currently at 37 weeks gestation and is in good health. The results of the 1st trimester screening are consistent with the results of the 2nd, 3rd, 4th, 5th, 6th, 7th, 8th, 9th, and 10th trimester screenings.



Reported Detection Rates

	Sequenom (MaterniT21)	Illumina (Verify)	Ariosa (Harmony)	Natera (Panorama)
Trisomy 21	99.1% 0.1%	>99.9% 0.2%	>99% <0.1%	>99% 0%
Trisomy 18	99.9% 0.4%	97.4% 0.4%	>98% <0.1%	96.4% <0.1%
Monosomy X	94.4% 0.6%	95% 1.0%	91.5% 0%	>99.9% 0%
Sex chromosome	96.2%	67-100%	99%	92.9% <0.1%
Female	99.1% 0.5%	97.6% 0.8%	99% 0%	>99.9% 0%
Male	99.4% 0.9%	99.9% 1.1%	100% 1%	>99.9% 0%
Triploidy	unable to detect	unable to detect	Unable to detect	>99%



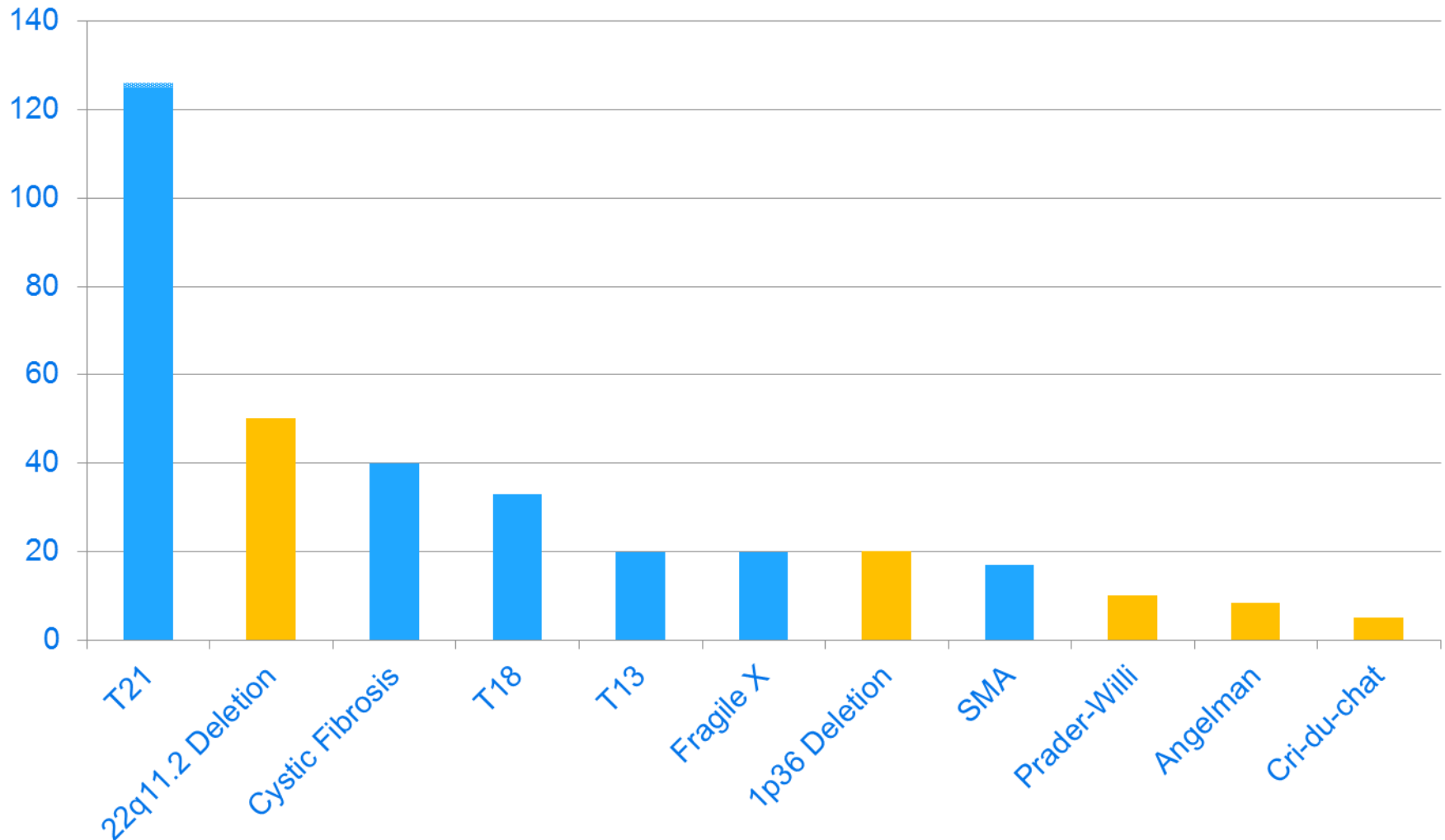
Microdeletions

22q11 deletion (associated with DiGeorge syndrome)
15q11 deletion (associated with Prader-Willi / Angelman syndrome)
11q23 deletion (associated with Jacobsen syndrome)
8q24 deletion (associated with Langer-Giedion syndrome)
5p15 deletion (associated with Cri-du-chat syndrome)
4p16 deletion (associated with Wolf-Hirschhorn syndrome)
1p36 deletion syndrome



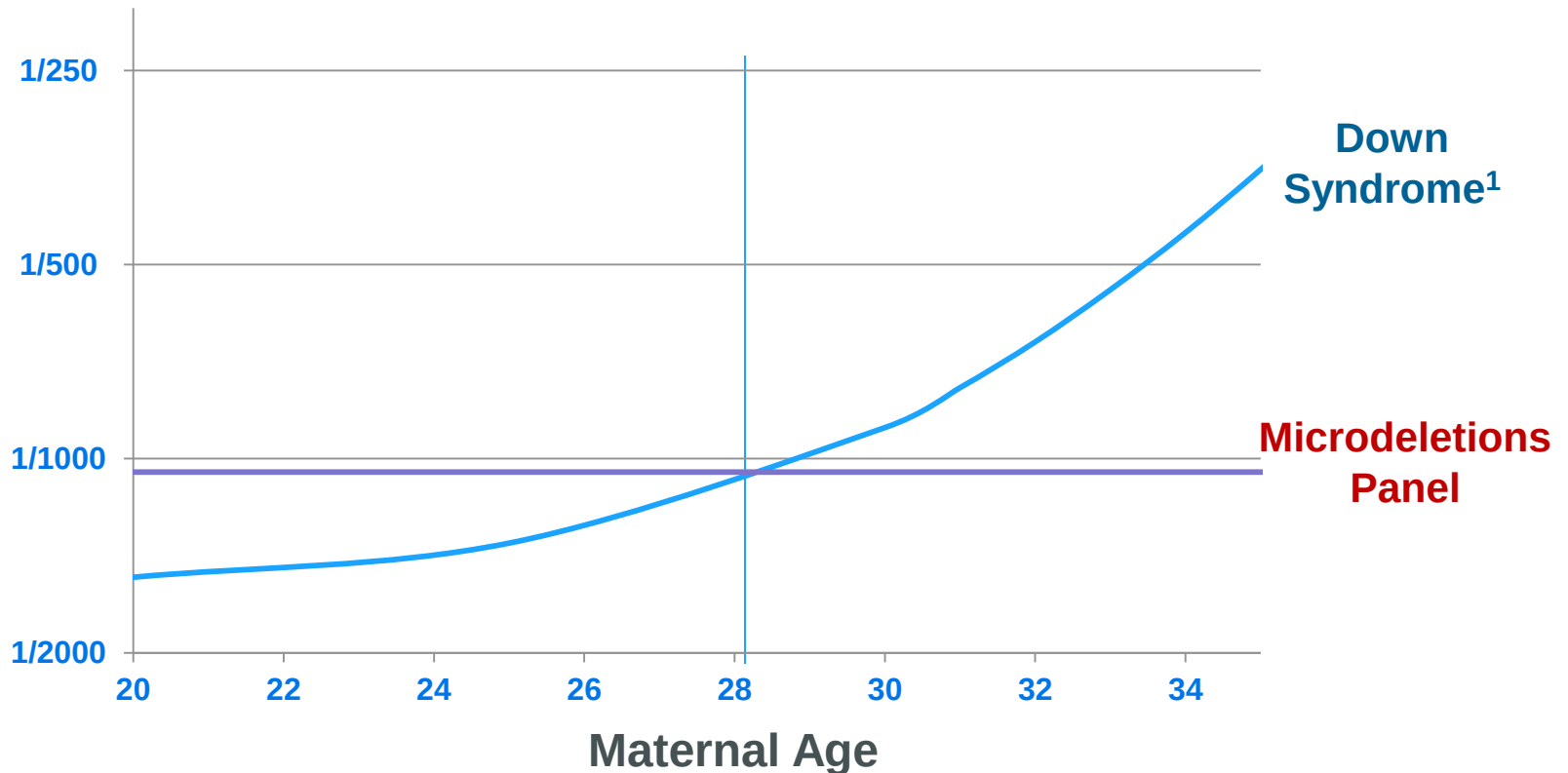
Microdeletions

Incidence out of 100,000 Live Births



Microdeletions

More Common Than Down Syndrome in Younger Women



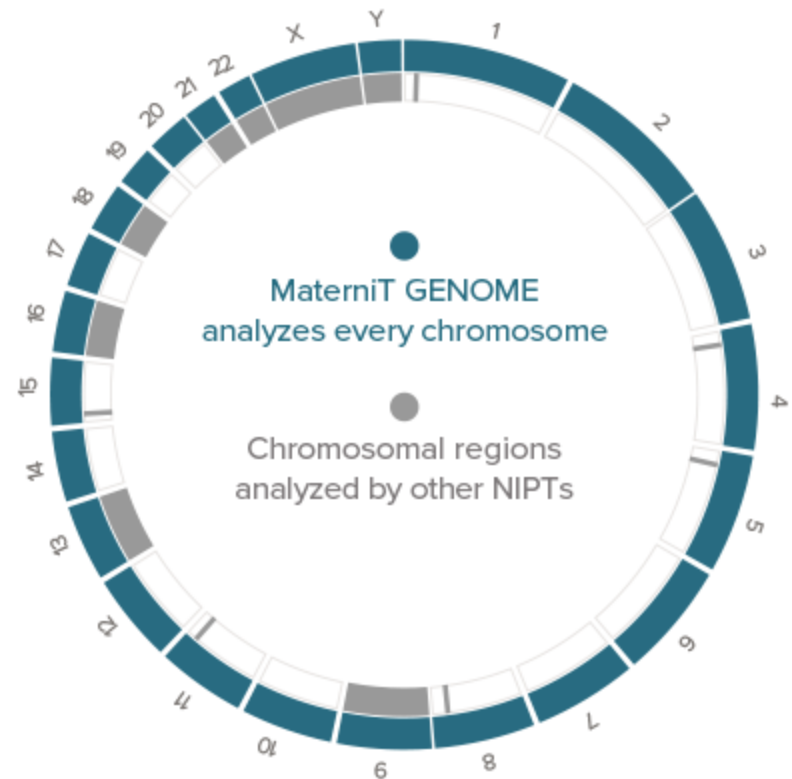
¹Snijders, et al. Ultrasound Obstet Gynecol 1999;13:167–170.

²Combined prevalence using higher end of published ranges from Gross et al. Prenatal Diagnosis 2011; 39, 259-266; and www.genetests.org. Total prevalence may range from 1/1071 - 1/2206.

Sequenom's MaterniT Genome

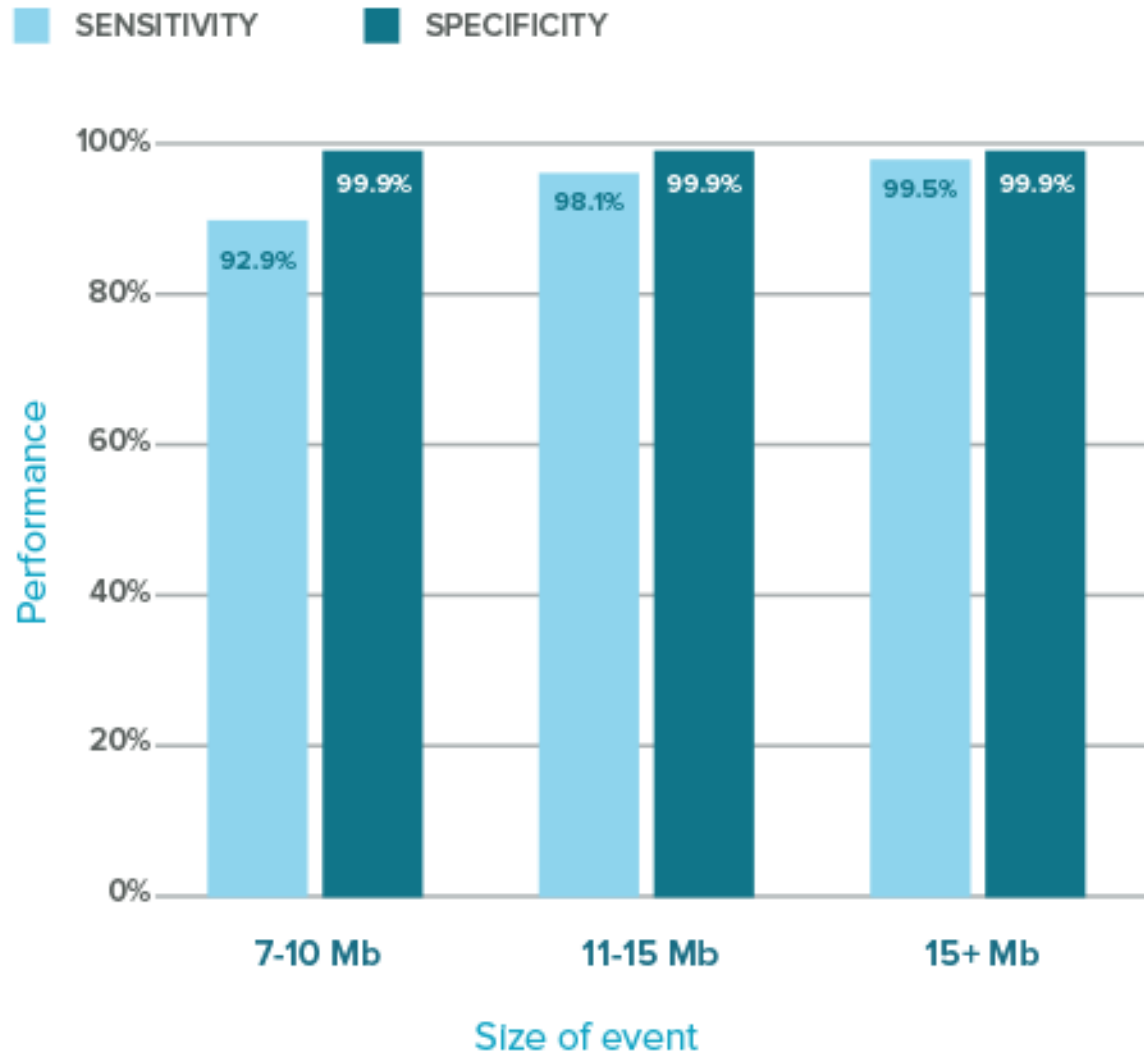
Genome-wide deletions or duplications of 7 Mb and greater, and also detects select microdeletions below 7 Mb.

	Fetal karyotype test	MaterniT GENOME test
Analyzes every chromosome	Yes	Yes
Requires an invasive procedure	Yes	No
Detects large, unbalanced translocations	Yes	Yes
Detects marker chromosomes	Yes	Yes
Detects balanced translocations or inversions	Yes	No
Detects chromosome gains or losses as small as 7 mb	No	Yes
Detects select microdeletions	No	Yes
Detects Triploidy	Yes	No
Considered diagnostic	Yes	No



Sequenom's MaterniT Genome

GENOME-WIDE PERFORMANCE



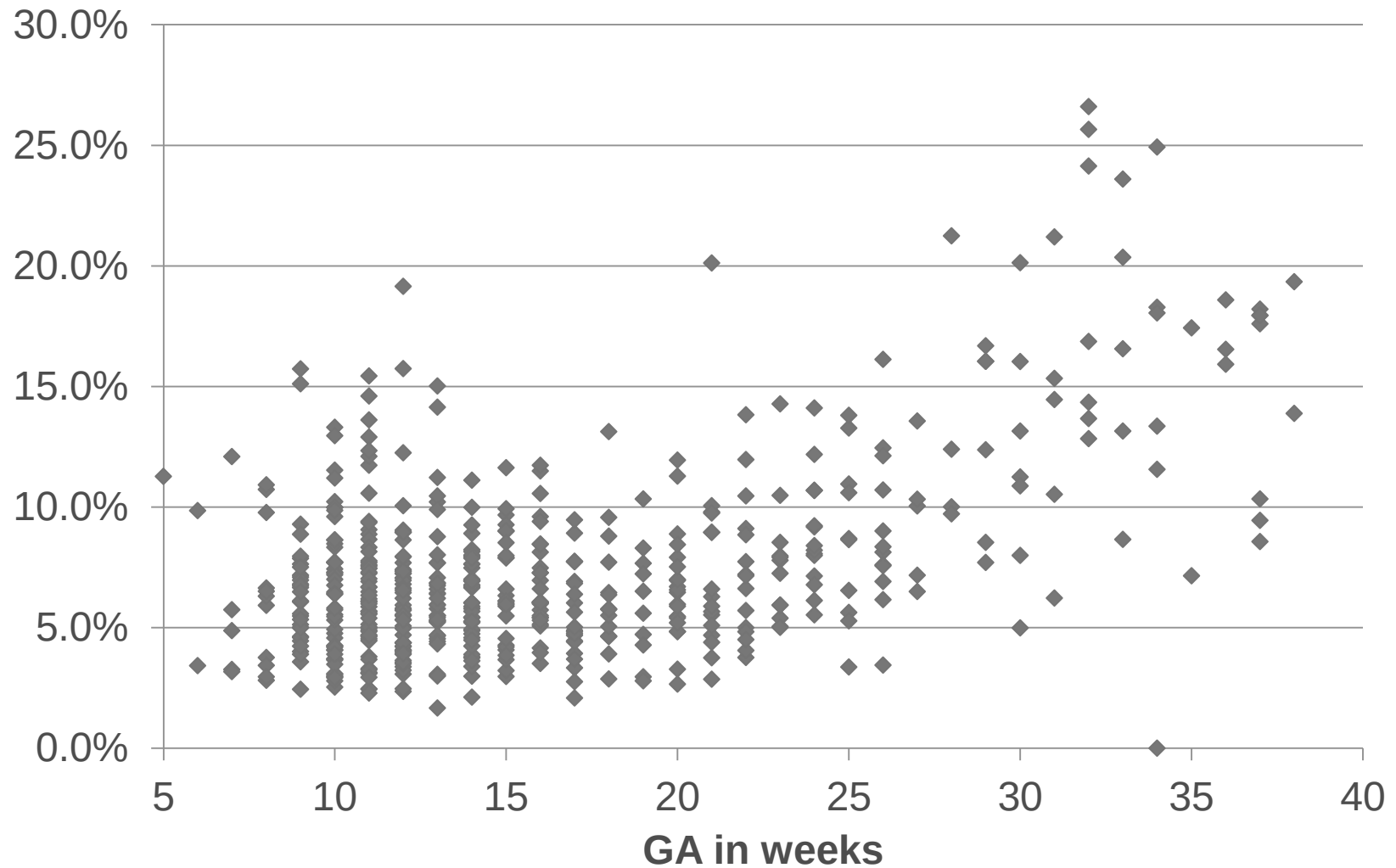


Known Causes of Discrepancies

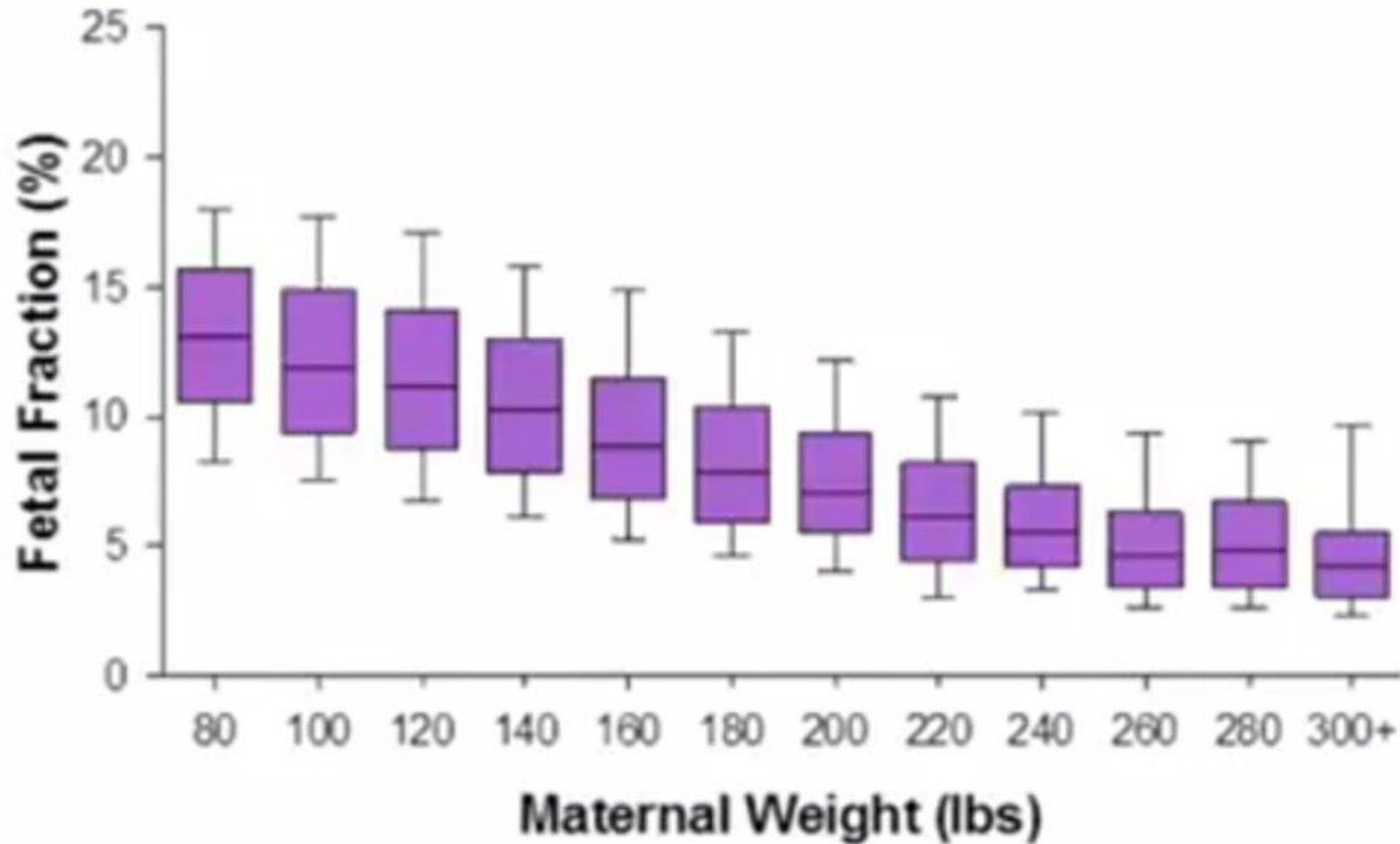
Cause	Example	Reference
True fetal mosaicism	cfDNA: + or - Invasive: two cell lines	Canick et al., Prenatal Diagn 2013
Confined placenta mosaicism	cfDNA: - Invasive: 47,XY,+21	Wang et al., Prenatal Diagn, 2013
Twin pregnancy (vanishing twin)	cfDNA: + liveborn twin 46,XX Vanishing 47,XX, +21	Gromminger et al., JCM 2014
Maternal chromosome abnormality	47,XXX	Wang et al., Clin Chem 2014
Maternal somatic cell variation	cfDNA: + Maternal loss of X (45,X)	Wang et al., Clin Chem 2014
Maternal malignancy	cfDNA: +13 Invasive: 46,XY Cancer found in mom	Osborne et al., Prenat Diagn, 2013
Low fetal fraction	cfDNA: - Invasive: 47,XY, +21	Allen et al., ACMG meeting; abstract, 2013



Fetal Fraction Related to GA



Fetal Fraction Maternal Weight



Fetal fraction is inversely proportional to maternal weight:

↑ maternal DNA + ↓ fetal DNA



Fetal Fraction and Aneuploidy

- 17/31 samples with low fetal fraction ($<3.4\%$) were aneuploid
- 119/1015 samples with above 3.4% fetal fraction were aneuploid

Translated to OR of 9.2X risk of aneuploidy with low fetal fraction.

Pergament E, Cuckle H, Zimmermann B, Banjevic M, Sigurjonsson S, Ryan A, et al. Single-nucleotide polymorphism-based noninvasive prenatal screening in a high-risk and low-risk cohort. *Obstet Gynecol* 2014;124:210–8.



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



Society for
Maternal-Fetal
Medicine

(Published Electronically Ahead of Print on June 26, 2015)

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.

Cell-free DNA Screening for Fetal Aneuploidy

ABSTRACT: Noninvasive prenatal screening that uses cell-free DNA from the plasma of pregnant women offers tremendous potential as a screening method for fetal aneuploidy. A number of laboratories have validated different techniques for the use of cell-free DNA as a screening test for fetal aneuploidy. All tests have a high sensitivity and specificity for trisomy 18 and trisomy 21, regardless of which molecular technique is used. 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. Patients should be counseled that cell-free DNA screening does not replace the precision obtained with diagnostic tests, such as chorionic villus sampling or amniocentesis and, therefore, is limited in its ability to identify all chromosome abnormalities. Cell-free DNA screening does not assess risk of fetal anomalies such as neural tube defects or ventral wall defects. Patients who are undergoing cell-free DNA screening should be offered maternal serum alpha-fetoprotein screening or ultrasound evaluation for risk assessment. The cell-free DNA screening test should not be considered in isolation from other clinical findings and test results. Management decisions, including termination of the pregnancy, should not be based on the results of the cell-free DNA screening alone. Patients should be counseled that a negative cell-free DNA test result does not ensure an unaffected pregnancy. 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.

- 
- Offer to patients with high risk
 - Advanced maternal age;
 - Fetal ultrasound abnormality;
 - Personal or family history of Down syndrome or other chromosomal aneuploidy; and/or
 - Positive serum screening test
 - Low risk population?
 - All women should receive information on all testing options
 - Any women can choose any method regardless of risk status
 - Conventional screening most appropriate for low risk given limitations (adverse pregnancy outcomes and PPV)

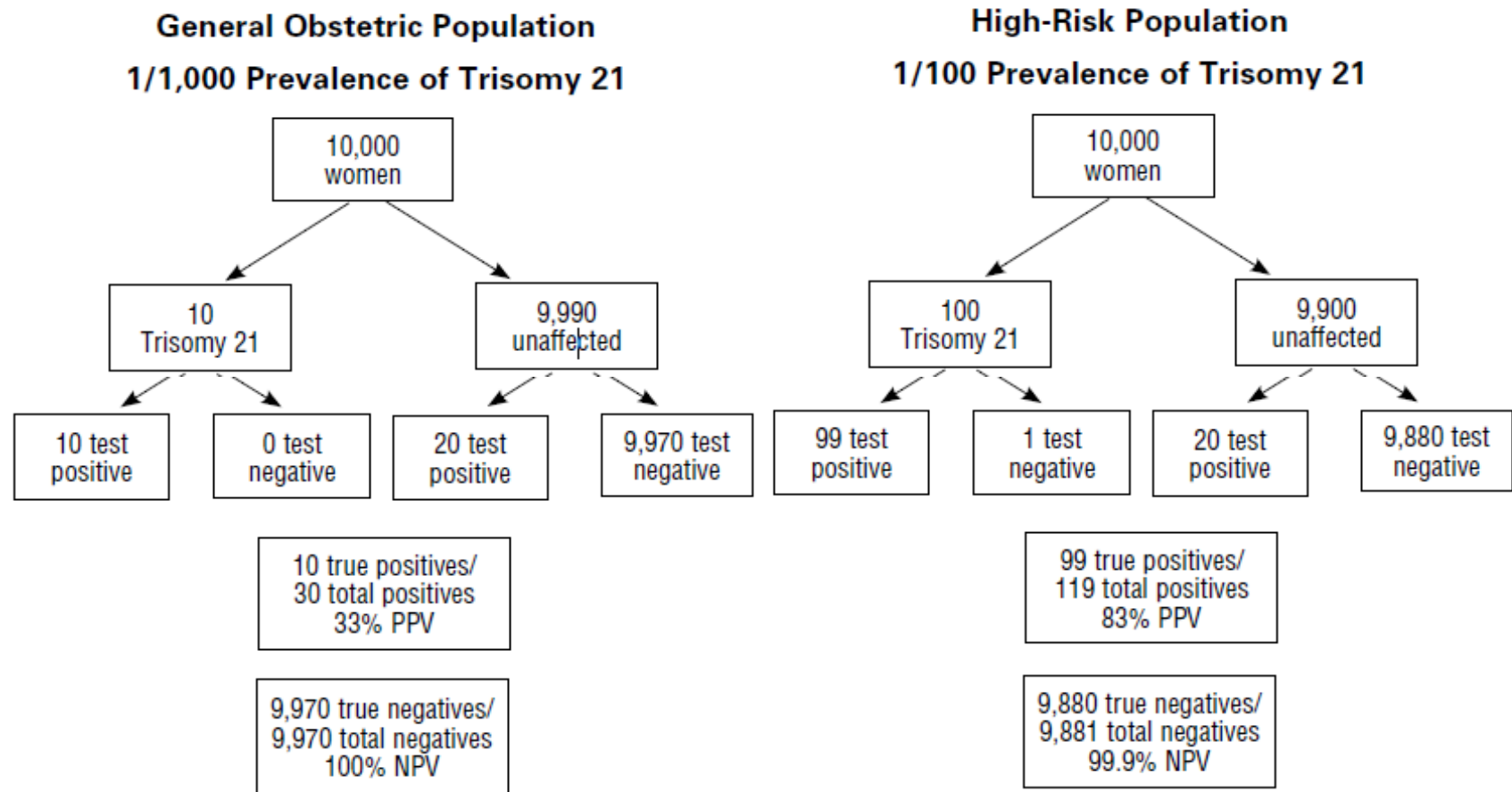


Fig.1. The importance of population prevalence on the predictive value for a screening test: an illustration with cell-free DNA.

Abbreviations: NPV, negative predictive value; PPV, positive predictive value



Test	DR	FPR	PPV (high risk 1/100)	PPV (low-risk 1/500)
FTS (NT, PAPP-A, hCG)	80%	3%	21%	5%
Quad	60%	3%	17%	4%
Sequential	93%	3%	24%	6%
cfDNA (all methods)	99.3%	0.1%	91%	67%

DR, FPR, for conventional screening: Benn et al. Prenat Diagn. 2013; 31:519-22
Based on meta-analysis of 19 validation studies (all methods)

- Microdeletions
- Multiple pregnancies
- Recommend diagnostic testing for positive results
- Management decisions, including TOP, should not be made on cfDNA testing alone
- Should be offered MS-AFP
- If anomaly identified offer diagnostic testing
- Women with failed results should receive further counseling and offered ultrasound and diagnostic testing.

Carrier Testing



HETEROZYGOATS

Just allele uneven.

Current Guidelines – Ancestry

ACOG Guidelines

- African Americans,
- Southeast Asians,
- Mediterranean

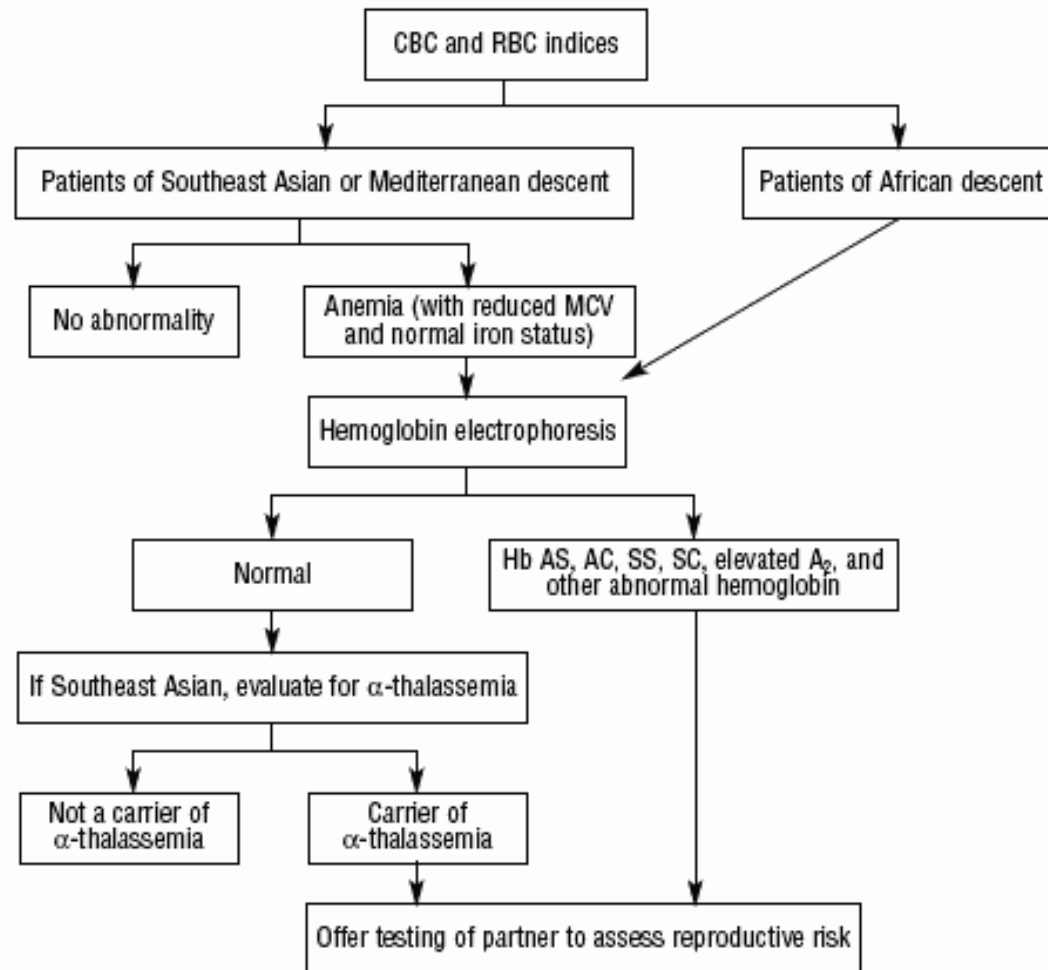


Figure 1. Specialized antepartum evaluation for hematologic assessment of patients of African, Southeast Asian, or Mediterranean descent. Patients of Southeast Asian or Mediterranean descent should undergo electrophoresis if their blood test results reveal anemia. Abbreviations: CBC, complete blood count; Hb, hemoglobin; MCV, mean corpuscular volume; RBC, red blood cell.



Current Guidelines – Ancestry

Gaucher Type I

*!Tay-Sachs

*Cystic Fibrosis

*Familial dysautonomia

*Canavan

Niemann-Pick A

Fanconi anemia C

Bloom syndrome

Mucopolysaccharidosis IV



Biochemical screening of hexosaminidase is the most sensitive screening in all populations.

*= For Ashkenazi Jewish ACOG recommended, ACMG recommends all

! = offer in those of French Canadian and Cajun ancestry

Current Guidelines – Panethnic



The American College of Obstetricians and Gynecologists
Women's Health Care Physicians

COMMITTEE OPINION

Number 486 • April 2011

(Replaces No. 325, December 2005)

Committee on Genetics

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.

Update on Carrier Screening for Cystic Fibrosis

ABSTRACT: In 2001, the American College of Obstetricians and Gynecologists and the American College of Medical Genetics introduced guidelines for prenatal and preconception carrier screening for cystic fibrosis. The American College of Obstetricians and Gynecologists' Committee on Genetics has updated current guidelines for cystic fibrosis screening practices among obstetrician–gynecologists.

Every Patient

Current Guidelines – Targeted

- Fragile X
 - Pan-Ethnic carrier frequency: all women have ~1/250 or greater risk to be carriers
 - Severe disorder with no cure
- Offer testing:
 - Family history of fragile X
 - Family history of ID of unknown etiology
 - Family history of autism
 - Personal history of POI
 - Personal history of ataxia, esp male





CGG Repeat

- Trinucleotide repeat – CGG
 - **Expansion of CGG repeats within the gene occurs when inherited through the mother**

Fragile X Result	CGG Repeat Sizes
Normal	<45
Intermediate	45 – 54
Premutation (Carrier)	55 - 200
Full Mutation	>200



Anticipation

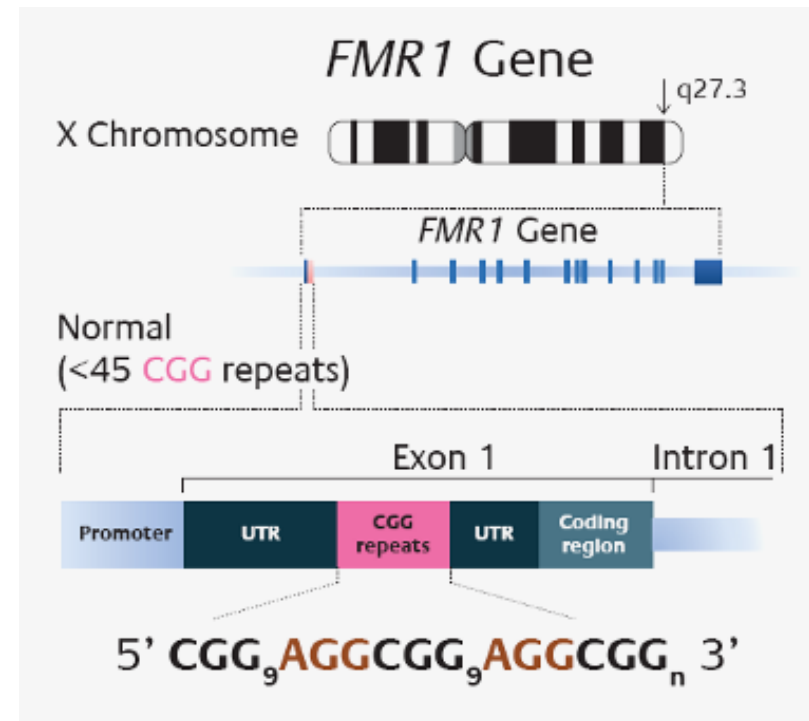
Number of Maternal Premutation CGG Repeats	Number which expanded to full mutations
55-59	1 (4%)
60-69	6 (5%)
70-79	28 (31%)
80-89	81 (58%)
90-99	89 (80%)
100-200	193 (98%)

CGG repeat length is not the only factor accounting for instability



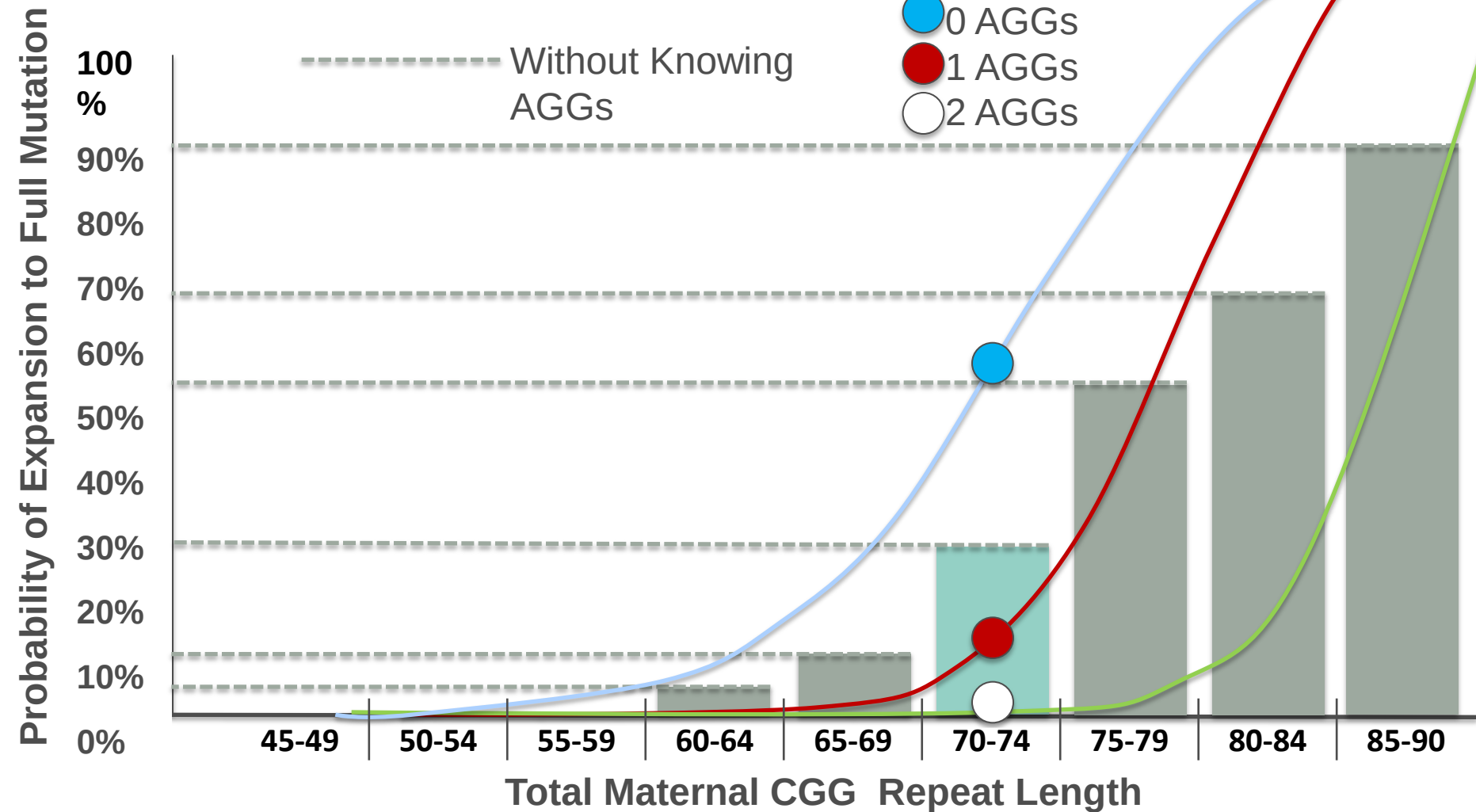
Stability of Allele

- AGG repeat
 - In the normal population, CGG repeats interrupted by AGG at positions 10 and 20
 - Lead to stability to the repeat – think of it as an anchor
 - Some laboratories offer AGG interruption number





Impact of AGGs



<http://asuragen.com/products-and-services/clinical-lab/xpansion-interpreter/>

Nolin et al., American Journal of Medical Genetics A. 2013;161(4):771-8.

Coppinger et al., Platform Presentation, American College of Medical Genetics Annual Clinical Genetics Meeting. 2013



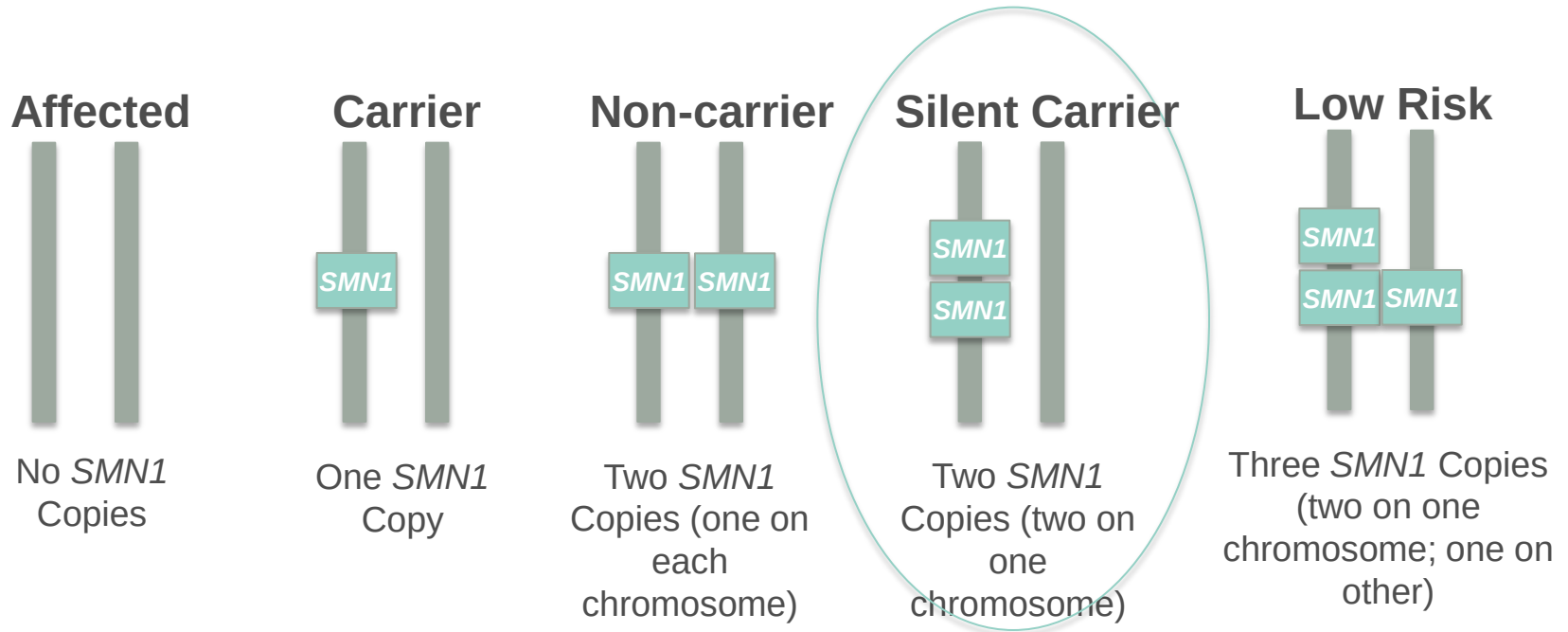
Current Guidelines – Targeted

- Spinal Muscular Dystrophy (SMA)
 - Severe Autosomal recessive Neuromuscular disease
 - Pan-ethnic carrier frequency: ~ 1:50
 - 1/6000 – 1/10,000
 - Standard SMA carrier screening determines SMN1 exon7 copy number
- Guidelines
 - ACOG – only with family history
 - ACMG – offer regardless of ancestry or family history



Gene Copy Number

- Routine carrier screening looks at *SMN1* gene copy number
- 2 copies of the *SMN1* gene could mean a non-carrier OR a silent carrier (2+0)





SNP Analysis for Silent Carriers

- SNP analysis
 - SNP in intron 7 of SMN1 (g.27134t>G) associated with “Silent Carriers”
 - If patient carries two SMN1 copies and the SNP is present, there is an increased likelihood of being a silent carrier
 - Ashkenazi Jewish or Asian – likely carrier
 - Caucasian, African American, Hispanic – risk increased
 - Routine screening would miss ~4% of carriers

Expanded Carrier Testing

Current Commentary

Expanded Carrier Screening in Reproductive Medicine—Points to Consider

A Joint Statement of the American College of Medical Genetics and Genomics, American College of Obstetricians and Gynecologists, National Society of Genetic Counselors, Perinatal Quality Foundation, and Society for Maternal-Fetal Medicine

Janice G. Edwards, MS, Gerald Feldman, MD, PhD, James Goldberg, MD, Anthony R. Gregg, MD, Mary E. Norton, MD, Nancy C. Rose, MD, Adele Schneider, MD, Katie Stoll, MS, Ronald Wapner, MD, and Michael S. Watson, MD

The Perinatal Quality Foundation and the American College of Medical Genetics and Genomics, in association with the American College of Obstetricians and Gynecologists, the Society for Maternal-Fetal Medicine, and the National Society of Genetic Counselors, have collaborated to provide education for clinicians and laboratories regarding the use of expanded genetic

carrier screening in reproductive medicine. This statement does not replace current screening guidelines, which are published by individual organizations to direct the practice of their constituents. As organizations develop practice guidelines for expanded carrier screening, further direction is likely. The current statement demonstrates an approach for health care providers and laboratories who wish to or who are currently offering expanded carrier screening to their patients.

(Obstet Gynecol 2015;125:653–62)

DOI: 10.1097/AOG.0000000000000666

From the American College of Medical Genetics and Genomics, Bethesda, Maryland; the American College of Obstetricians and Gynecologists and the Society for Maternal-Fetal Medicine, Washington, DC; the National Society of Genetic Counselors, Chicago, Illinois; and the Perinatal Quality Foundation, Oklahoma City, Oklahoma.

Supported by the Perinatal Quality Foundation. The authors thank Jean Lee Spitz, MPH, RDMS, Executive Director of the Perinatal Quality Foundation.

Ms. Edwards was the consensus facilitator.

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Financial Disclosure

Dr. Feldman is the director of a clinical laboratory offering molecular diagnostics, including carrier screening. Dr. Norton receives grant funding from Natera, Inc. and Ariosa Diagnostics, companies that perform noninvasive prenatal screening. She is also on the advisory board of Natera, Inc. Dr. Schneider is a consultant to Quest Diagnostics. Dr. Wapner is on the advisory boards for Baylor College of Medicine Medical Genetics Laboratories and CombiMatrix, which provide molecular diagnostic testing. He also receives federal funding for research related to prenatal diagnostic testing and receives research funding from Ariosa Diagnostics, Illumina, Natera, and Sequenom related to prenatal screening. The other authors did not report any potential conflicts of interest.

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ISSN: 0029-7844/15

Conditions included on a carrier screening panel should encompass one or more of the following criteria:

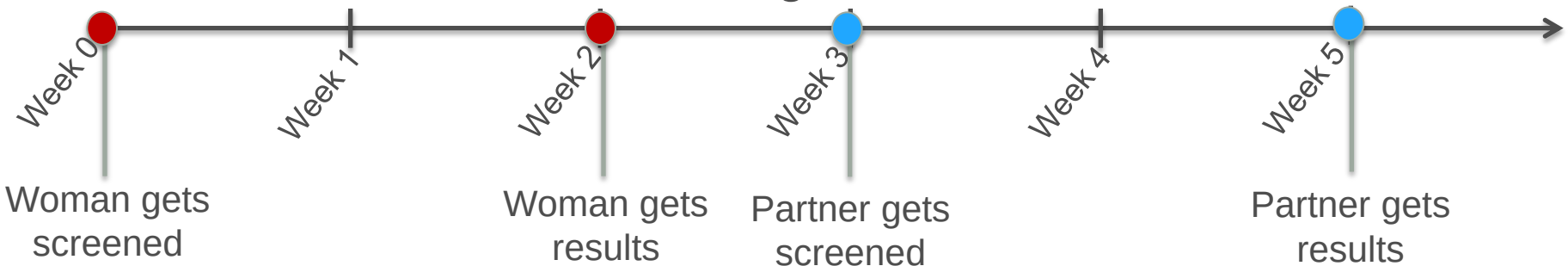
- Cognitive disability
- Need for surgical or medical intervention
- Effect on quality of life
- Prenatal diagnosis may result in prenatal intervention to improve perinatal outcomes, delivery management, or prenatal education of parents to prepare for special needs after birth



Concurrent Carrier Screening

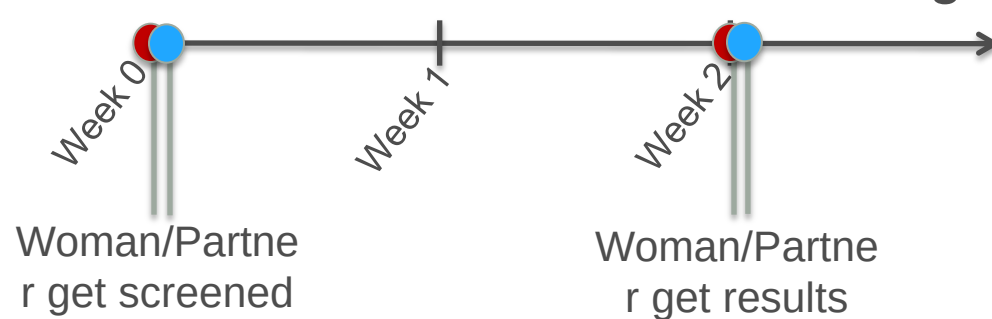
Speeds Up Identification of Potential Risk to the Fetus

Timeline of Individual Screening



Total turnaround time can be 5+ weeks...With two positives, is there still time for CVS or amnio? And what about the emotional stress of waiting?

Timeline of Concurrent Screening



Done in about 2 weeks =
more time for CVS or amnio
AND reduced emotional
burden



Considerations

- Hex A and CBC/hemoglobin electrophoresis still most appropriate.
- Recommend pre and post test counseling
 - Varying degrees of severity and inheritance
 - Risk reduction not elimination
 - Patient could be found with two mutations
 - Paternal information is needed for accurate risk
 - Sequencing possible but not routinely recommended
 - Offer follow-up if applicable



If You Would Like to Refer a
Patient or Have a Question ...

We'd be happy to help!
Call Perinatal Genetics at 581-7825