

## **Developing NIPD for fetal aneuploidy**

Helen White, PhD

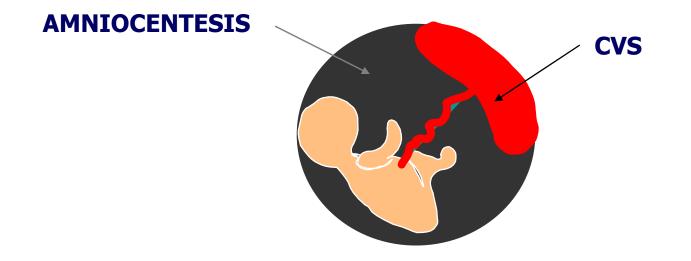
**Senior Scientist** 

National Genetics Reference Lab (Wessex)

## **Outline of talk**

- Current prenatal screening for aneuploidy
- How can cell free fetal nucleic acids be used for DS testing?
- New non-invasive techniques for detection of DS
  - Quantitative SNP analysis from cffRNA
  - Shot gun sequencing of cfDNA
  - Epigenetics
  - Digital PCR
- RAPID: Plans to develop NIPD for aneuploidy

### **Current prenatal screening for aneuploidy**



- Prenatal screening for is offered to all pregnant women
- Undertaken in two phases:

screening and risk assessment

invasive prenatal diagnosis of high risk cases

• Gold standard for diagnosis of chromosomal abnormalities is karyotyping

### **Current prenatal screening for aneuploidy**

Due to a small but significant risk to the pregnancy, many women are reluctant to opt for these procedures

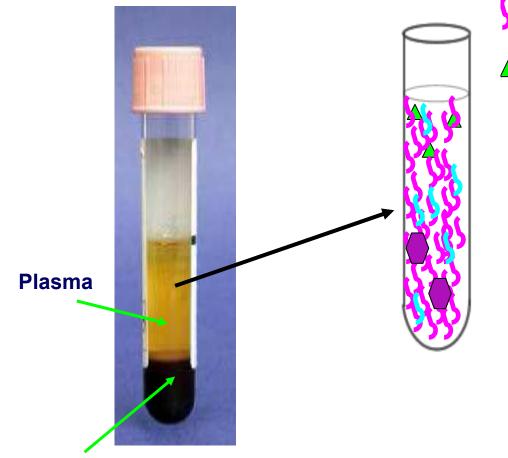
In 2006-7:

~700,000 pregnant women a year underwent antenatal screening

20,000 amniocentesis and 5,200 CVS were performed

Estimated associated procedural related pregnancy loss of ~250

## Extraction of cell free fetal nucleic acids from maternal plasma



Cell free maternal DNA (96.6%)

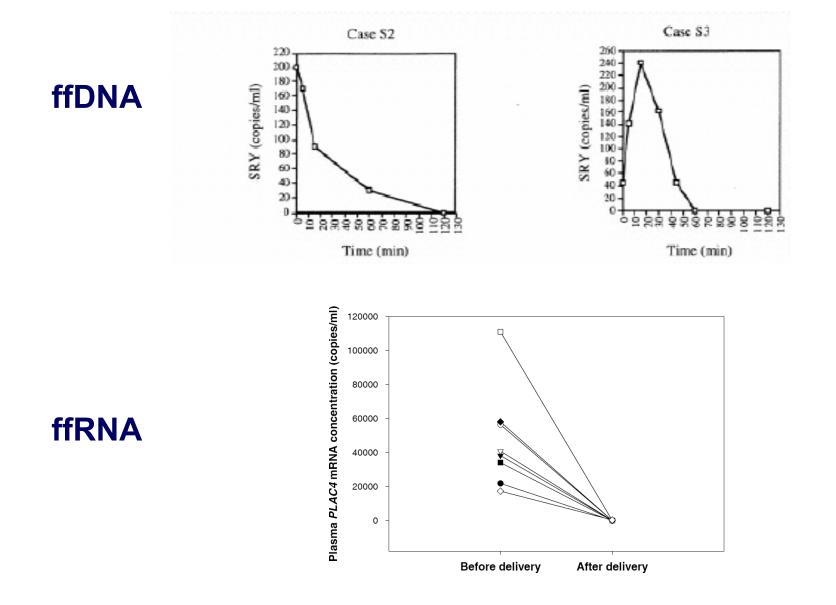
▲ Cell free fetal DNA (3.4%)

Amount of cell free fetal DNA extracted is equivalent to 25 cells / ml plasma

Cell free maternal RNA Cell free fetal RNA

Maternal blood cells

### Clearance of cell free fetal nucleic acids after delivery



## How can cell free fetal nucleic acids be used for non-invasive Down syndrome testing?

#### Major technical challenge

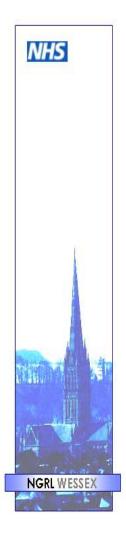
Background of cell free maternal **DNA** means direct quantification of fetal chromosome copy number is problematic and technically demanding

Ideally need:

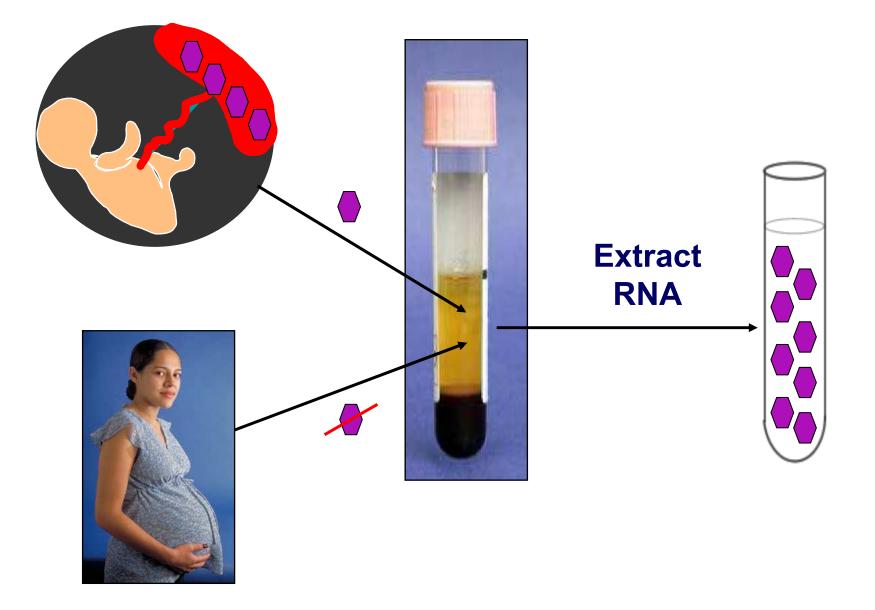
targets that are free from maternal background interference

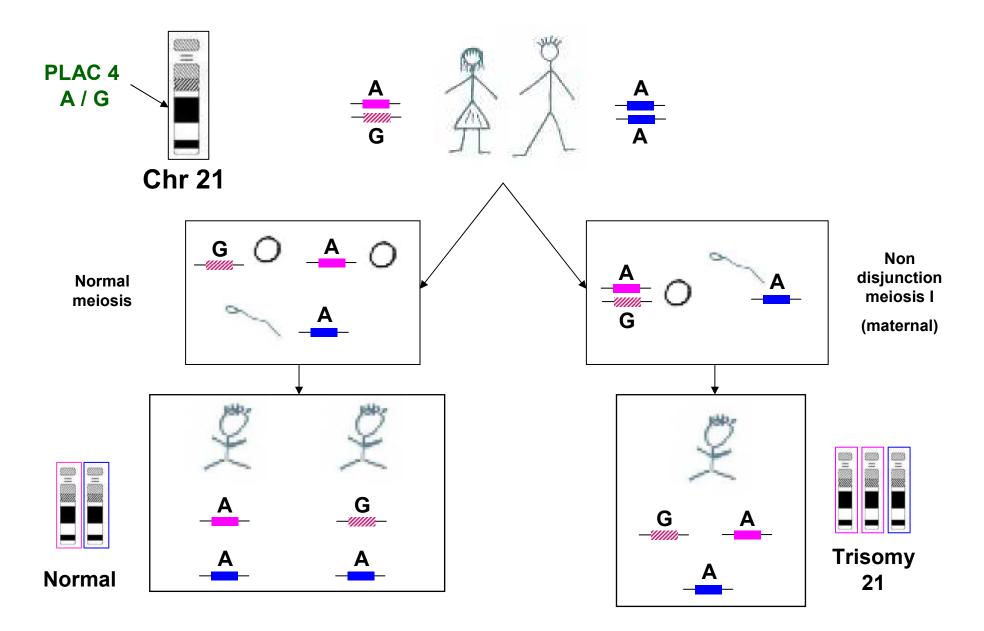
and / or

technologies that enable accurate & large scale copy number 'counting'



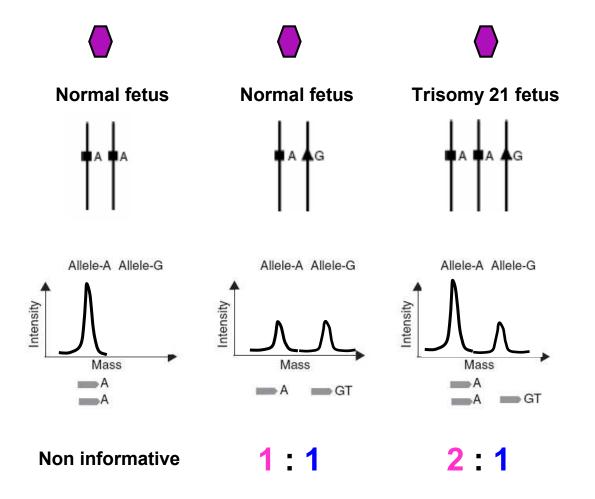
## Quantitative analysis of Single Nucleotide Polymorphisms in fetal specific mRNAs





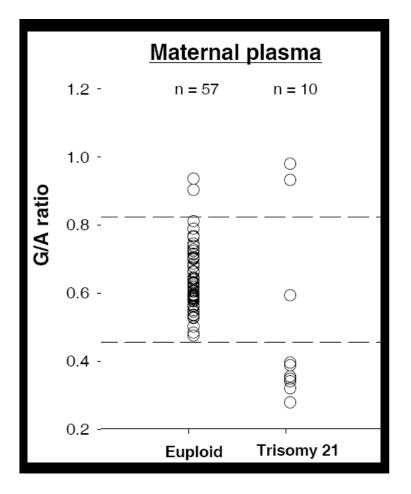
#### Analysis by MALDI-TOF (mass spectrometry)

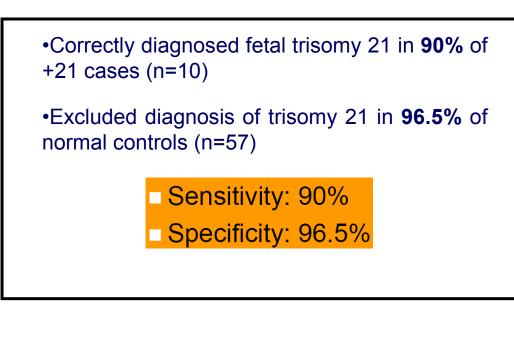
- PLAC4 mRNA ( ) is derived exclusively from fetal chromosome 21
- PLAC4 mRNA expressed in the placenta and is found in the plasma of pregnant women



Lo et al., Nature Med & PNAS, 2007

Analysis by MALDI-TOF (mass spectrometry)





Lo et al., Nature Med & PNAS, 2007

#### ADVANTAGES

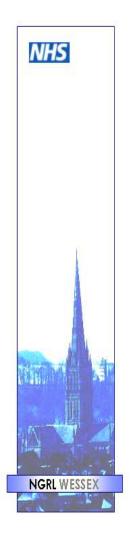
- Diagnostic sensitivity and specificity using one marker are comparable to multiple marker screening tests for DS currently in practice (if informative)
- Test is insensitive to gestational age and can be offered early in pregnancy
- Target free of maternal background
- Compatible with high throughput screening

#### DISADVANTAGES

- Fetus has to be informative for SNP analysed
- RNA can be unstable implications for sample collection

#### FUTURE REQUIREMENTS

- Identification of more polymorphic loci to increase informative cases
- Multi centre large scale validation required
- Expand testing to include fetal specific transcripts from chromosomes 18 & 13

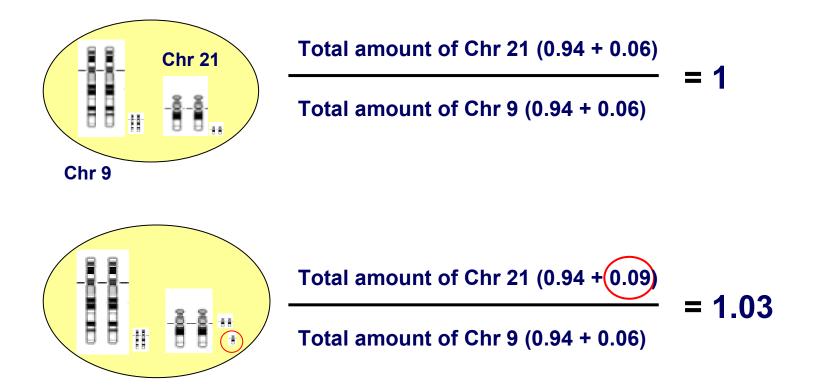


## Shot gun sequencing

# A method for digital quantification of DNA

## DNA testing preferable: is universal i.e. polymorphism independent

- Only 3-6% of the cell free DNA fraction is fetal
- Expected enrichment of chromosome 21 lies within the range of 1.5% to 3%



## Noninvasive diagnosis of fetal aneuploidy by shotgun sequencing DNA from maternal blood

H. Christina Fan\*, Yair J. Blumenfeld<sup>†</sup>, Usha Chitkara<sup>†</sup>, Louanne Hudgins<sup>‡</sup>, and Stephen R. Quake\*§

\*Department of Bioengineering, Stanford University and Howard Hughes Medical Institute, 318 Campus Drive, Clark Center, Room E300, Stanford, CA 94305; <sup>†</sup>Division of Maternal and Fetal Medicine, Department of Obstetrics and Gynecology, Stanford University, 300 Pasteur Drive, Room HH333, Stanford, CA 94305; and <sup>‡</sup>Division of Medical Genetics, Department of Pediatrics, Stanford University, 300 Pasteur Drive, Stanford, CA 94305

Communicated by Leonard A. Herzenberg, Stanford University School of Medicine, Stanford, CA, August 22, 2008 (received for review July 13, 2008)

#### • Shotgun sequenced plasma DNA samples from 18 pregnant women:

9 trisomy 21

2 trisomy 18

1 trisomy 13

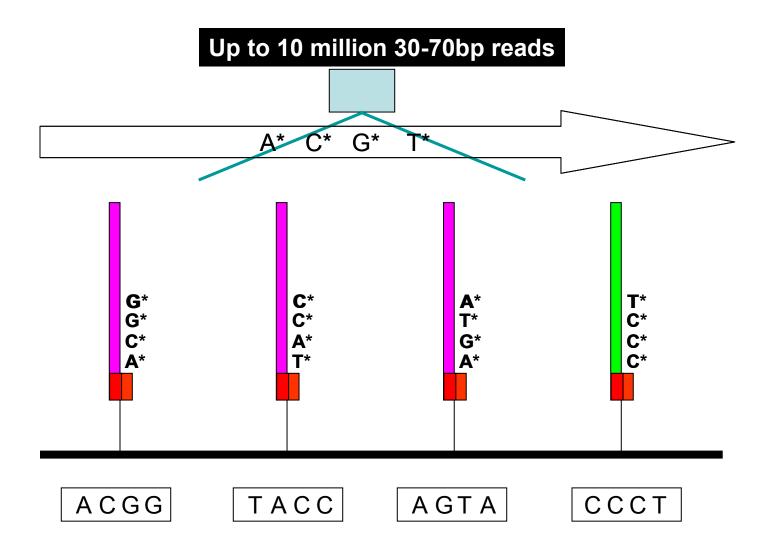
6 normal

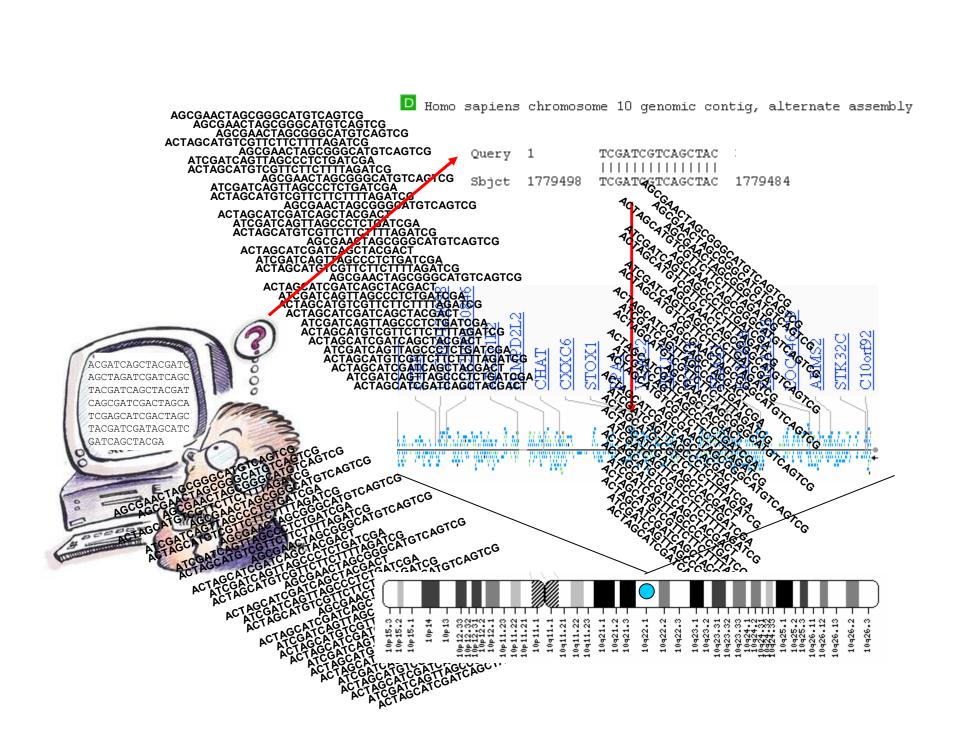
1 genomic DNA sample from a male control

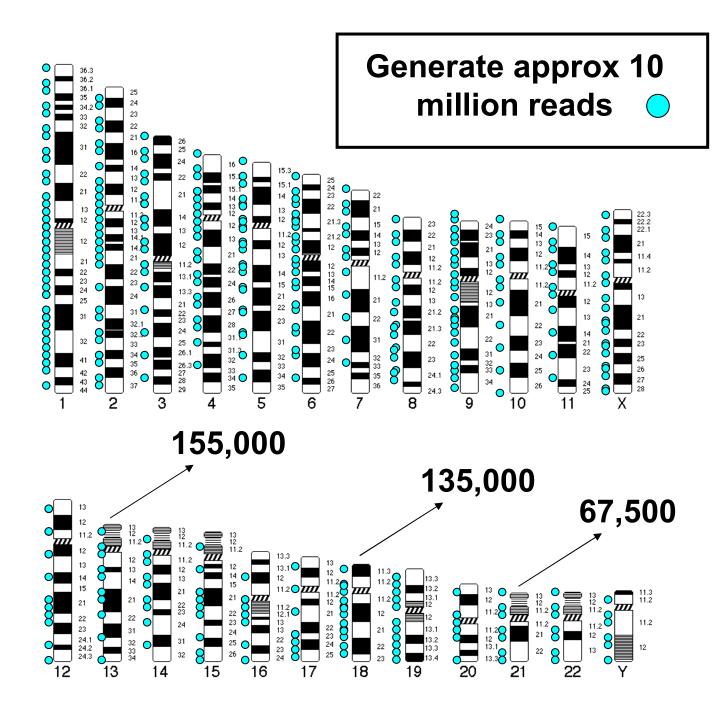
• Gestational age 10 – 35 weeks (earliest trisomy case 14 weeks)

Fan et al., PNAS Oct 2008

## **Sequencing by Synthesis**







## Noninvasive diagnosis of fetal aneuploidy by shotgun sequencing DNA from maternal blood

H. Christina Fan\*, Yair J. Blumenfeld<sup>†</sup>, Usha Chitkara<sup>†</sup>, Louanne Hudgins<sup>‡</sup>, and Stephen R. Quake\*§

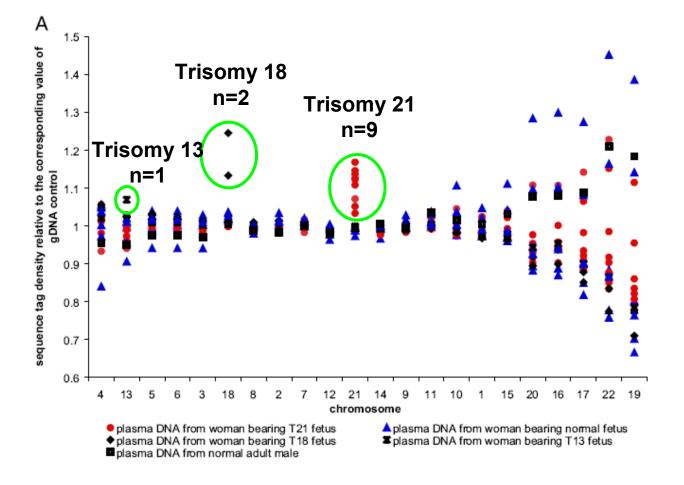
\*Department of Bioengineering, Stanford University and Howard Hughes Medical Institute, 318 Campus Drive, Clark Center, Room E300, Stanford, CA 94305; <sup>†</sup>Division of Maternal and Fetal Medicine, Department of Obstetrics and Gynecology, Stanford University, 300 Pasteur Drive, Room HH333, Stanford, CA 94305; and <sup>‡</sup>Division of Medical Genetics, Department of Pediatrics, Stanford University, 300 Pasteur Drive, Stanford, CA 94305

Communicated by Leonard A. Herzenberg, Stanford University School of Medicine, Stanford, CA, August 22, 2008 (received for review July 13, 2008)

- 5 million sequencing reads for each patient
- Compared density of reads on each chromosome to those obtained from a normal genomic DNA sample
- Compared density of Chr 21 reads from disomy and trisomy 21 samples

• Coverage of Chr 21 sequences in trisomy 21 was 4 – 18% higher than disomic cases

### **Results of shotgun sequencing of maternal plasma DNA**



Fan et al., PNAS Oct 2008

## Shot gun sequencing

#### ADVANTAGES

- Successful proof of principal study for detection of major trisomies; 13, 18 & 21
- Polymorphism independent and could be used in all pregnancies
- Has potential to detect unbalanced chromosome rearrangements

#### DISADVANTAGES

- Expensive and large amount of data processing interpretation.
- In current form would not be feasible to adapt to high throughput screening

#### FUTURE REQUIREMENTS

• Technological development required to produce machines and workflow protocols that could cope with a high throughput of samples at reduced cost

## **RAPID:** Plans to develop NIPD for aneuploidy

 Matched maternal blood samples and cultured fetal cells are currently being collected from women undergoing invasive prenatal testing in London and the Wessex region

• Assessment of analytical and clinical validity of tests for the noninvasive detection of DS (and trisomy 13 & 18):

- SNP allele ratios using MALDI-TOF mass spectrometry
- digital PCR
- development of targetted new generation sequencing assays
- cffDNA and cffRNA will be used to evaluate NIPD tests and cultured fetal cells will be used to confirm the NIPD result
- Sensitivity and specificity of each assay will be determined

