



**Preconception genetic carrier screening and PGT**



Margareta D. Pisarska MD  
Director, Division of Reproductive Endocrinology and Infertility  
Professor, Department of Ob/Gyn and Biomedical Sciences  
Cedars-Sinai Medical Center  
Professor, David Geffen School of Medicine at UCLA

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**Disclosures**

- *Ferring*
- *Natara*

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**Objectives**

- *Address preconception genetic carrier screening*
- *Preimplantation Genetic Testing*
- *Prenatal Genetic Screening and Testing*
- *Utilization of genomics and technologies for pregnancy well being*

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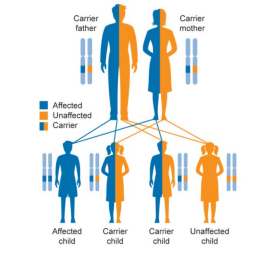
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### Carrier screening – identification of autosomal recessive disorders

Cross: Aa x Aa

	A	a
A	AA	Aa
a	Aa	aa

Autosomal Recessive Inheritance



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### Carrier screening for genetic conditions ACOG 2017



The American College of Obstetricians and Gynecologists  
www.acog.org | www.aog.org

#### COMMITTEE OPINION

Number 691 • March 2017  
Replaces 2005

Replaces Committee Opinion Number 238, October 2005  
Committee Opinion Number 691, May 2008  
Committee Opinion Number 452, October 2009  
Committee Opinion Number 493, October 2012  
Committee Opinion Number 686, April 2015

**Committee on Genetics**  
This committee opinion was developed by the American College of Obstetricians and Gynecologists' Committee on Genetics in consultation with its members.  
It is intended to assist obstetricians, gynecologists, and their patients in making decisions about genetic testing.  
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 endorsing one exclusive course of treatment or procedure to be followed.

**Carrier Screening for Genetic Conditions**

- Carrier screening to all couples, regardless of their race/ethnicity (ie, pan-ethnic carrier screening)
  - cystic fibrosis (CF)
  - spinal muscular atrophy (SMA)
- Carrier screening based on certain races/ethnicities
  - alpha thalassemia
  - Hb beta chain-related hemoglobinopathy (sickle cell disease)
  - Tay-Sachs disease
  - Canavan disease
  - familial dysautonomia

Cedars Sinai ACOG Committee Opinion- Carrier Screening for Genetic Conditions No. 691, March 2017

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### Preconception genetic carrier screening ACMG 2021

- Clinical utility is measured by the fact that individuals or couples are informed and may alter reproductive decision making because of the carrier screening results.
- Clinical utility is represented by its ability to provide individuals an opportunity to discuss their risks and consider reproductive options that are available pre-pregnancy, during pregnancy, or after birth. Availability of reproductive options may depend on various socioeconomic, legal, and cultural factors in different regions.
- Examples of reproductive options include:
  - In vitro fertilization with preimplantation genetic testing for monogenic conditions
  - Use of donor gamete/embryo
  - Adoption
  - Prenatal diagnosis using chorionic villus sampling or amniocentesis followed by a decision to either prepare for an affected child including special care after birth or terminate the pregnancy
  - A decision not to have children

Cedars Sinai Gregg AR, et al. ACMG Professional Practice and Guidelines Committee Genetics in Medicine (2021) 23:1793–1806; https://doi.org/10.1038/s41436-021-01203-

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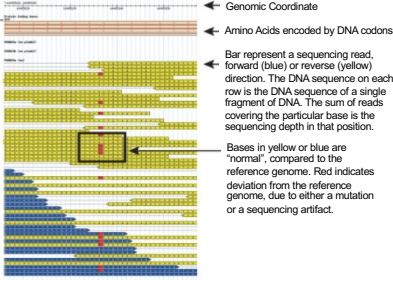
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### Next generation sequencing

- NGS platforms perform sequencing of millions of small fragments of DNA in parallel.
- Bioinformatics analyses are used to piece together these fragments by mapping the individual reads to the human reference genome.
- Each of the three billion bases in the human genome is sequenced multiple times, providing high depth to deliver accurate data and an insight into unexpected DNA variation.
- NGS can be used to sequence entire genomes or specific areas of interest, including all 22 000 coding genes (a whole exome) or small numbers of individual genes.



← Genomic Coordinate

← Amino Acids encoded by DNA codons

← Bar represent a sequencing read, forward (blue) or reverse (yellow) direction. The DNA sequence on each row is the DNA sequence of a single fragment of DNA. The sum of reads covering the particular base is the sequencing depth in that position.

← Bases in yellow or blue are "normal", compared to the reference genome. Red indicates deviation from the reference genome, due to either a mutation or a sequencing artifact.

Sam Behjati and Patrick S Tarpey Arch Dis Child Educ Pract Ed. 2013 Dec; 98(6): 236-238.

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### Next Generation Sequencing for carrier screening - 2021

- low cost
- high throughput identification of sequence variants across many genes simultaneously
- Allows equitable opportunities for patients to learn their reproductive risks using next-generation sequencing technology
- An improved understanding of this risk allows patients to make informed reproductive decisions
- Reproductive decision making is the established metric for clinical utility of population-based carrier screening
- Standardization of the screening approach will facilitate testing consistency



Gregg AR, et al. ACMG Professional Practice and Guidelines Committee Genetics in Medicine (2021) 23:1793-1806; https://doi.org/10.1038/s41436-021-01203-8

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### Carrier Screening for Genetic Conditions American College of Medical Genetics and Genomics (ACMG) 2021

- ACMG Goals
  - Develop carrier screening that is ethnic and population neutral and more inclusive of diverse populations to promote equity and inclusion

Public Database gnomAD v2.0.2<sup>18</sup>; 415 autosomal recessive

Carrier frequency in gnomAD at least 1/200 for six ancestral populations where Pathogenic and Likely Pathogenic variants were considered<sup>18</sup>

86 genes

Carrier frequency known to be at least 1/200 (however not captured in gnomAD v 2.0.2)

Genes with at least a 1/200 carrier frequency of pathogenic or likely pathogenic variants in a subpopulation that has at least 1% representation in the US including US territories.

9 genes

X-linked phenotypes (N=355) were identified in the OMIM database (November 30, 2020)<sup>19</sup> (Table S2)

Prevalence of the OMIM phenotypes (Table S2) were determined using OMIM<sup>20</sup>, Orphanet<sup>21</sup>, MedlinePlus<sup>22</sup>; prevalence required was at least 1/40,000

16 genes

\*All conditions included with at least moderate severity<sup>18,19</sup>

Gregg AR et al. ACMG Professional Practice and Guidelines Committee Genetics in Medicine (2021) 23:1793-1806; https://doi.org/10.1038/s41436-021-01203-8

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### Preconception genetic carrier screening ACMG

Tier 4<sup>+</sup>  
<1/200 carrier frequency (includes Tier 3)  
genes/condition will vary by lab

Tier 3<sup>+</sup>  
≥ 1/200 carrier frequency (includes Tier 2)  
includes X-linked conditions

Tier 2<sup>+</sup>  
≥1/100 carrier frequency (includes Tier 1)

Tier 1\*  
CF + SMA + Risk Based Screening

Tier 4 screening should be considered for a pregnancy that stems from a known or possible consanguineous relationship (second cousins or closer) or when a family or personal medical history warrants.

All pregnant patients and those planning a pregnancy should be offered Tier 3 carrier screening which tests for 112 genetic conditions

Limiting the carrier frequency to ≥1/100 creates missed opportunities to identify couples at risk for serious conditions

Carrier screening for two common conditions using a carrier frequency threshold of 1/100 may not be equitable across diverse populations.

Gregg AR et al. ACMG Professional Practice and Guidelines  
Committee Genetics in Medicine (2021) 23:1793-1806;  
<https://doi.org/10.1038/s41436-021-01203-3>

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### Carrier screening panel that supports equity across diverse populations

- Using evidence-based interpretations of both ACOG and ACMG criteria and leveraging carrier frequency data from >460,000 individuals across 11 ethnicities (self-reported) which identified 176 conditions and applied criteria from ACOG frequency threshold of ≥1 in 100 and ACMG threshold of ≥1 in 200.
- Forty conditions had carrier frequencies of ≥1 in 100 and 75 had carrier frequencies ≥1 in 200
- Following severity criteria a conservative equitable panel consisting of 37 conditions and a more permissive panels and ≥1 in 200 consists of 74 conditions.

number of conditions: 40 (1 in 100), 75 (1 in 200)

Taber, et al. Genetics in Medicine (2022) 24, 201-213

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### 176 Panel

Moderate or higher gene-disease association – 175 of 176 conditions (99.4%) Captures 99.8% of carriers and >99.9% of ARCs compared with a 176-condition panel

Moderate severity - 175 of 176 conditions (99.4%)

ACOG severity criterion, 165 of 176 - Captures 94.2% of carriers and 92.3% of ARCs

Age of onset (infancy/childhood) - 165 of 176 conditions (93.8%)

Gene-disease association: Definitive, Moderate, Limited

Severity: Profound, Severe, Moderate, Mild

Number of severity criteria met (not including "Onset early in life"): 3, 2, 1

Onset early in life: Infancy/childhood (0 -12 years), Adolescence (13 -21 years)

Taber, et al. Genetics in Medicine (2022) 24, 201-213

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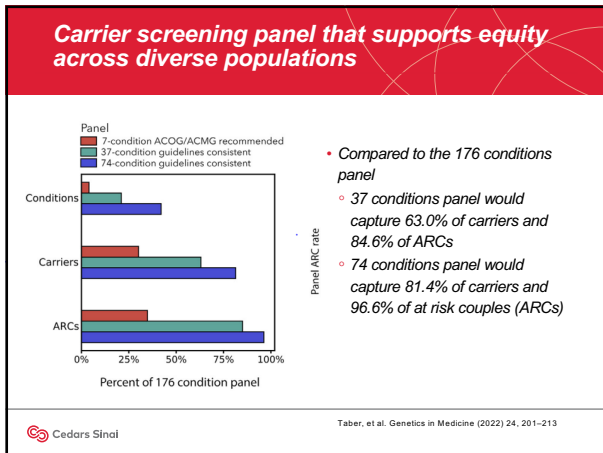
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### Genetic carrier screening – Impact on decision making (3 studies)

- 47% - screening was to spare a future child a life with a severe disorder
- Higher anxiety in high-risk and pregnant respondents
- 100% would opt for the test again
- Reproductive decision making was more common when patients received results **before an established pregnancy** (62–77%).
  - The most common decisions were
    - 59% in vitro fertilization with preimplantation genetic diagnosis
    - 20% diagnostic test during pregnancy
    - 7.7% use of a donor gamete
    - 5.1% consider adoption
- Testing during pregnancy
  - 16-36% had an affected fetus of those performing diagnostic testing
  - 40-67% discontinued their pregnancy

Ivy van Dijke, et al European Journal of Human Genetics (2021) 29:1252–1258;  
Ghossai, C. E., et al. J. Genet. Couns. 27, 616–625 (2018).  
Johansen Taber, K. A., et al Genet. Med. 21, 1041–1048 (2019)  
Gregg AR, et al ACMG Professional Practice and Guidelines Committee Genetics in Medicine (2021) 23:1793–1806

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### Carrier screening ACMG – Recommendations Tier 3 or Tier 4

- Carrier screening (Tier 3) is optional and can be performed at any time
- Preconception screening is recommended over prenatal screening
  - less stressful on patients with positive screening
  - allows for the full complement of reproductive decision making
- If done in pregnancy, concurrent partner testing should be offered
- When a reproductive partner has changed, carrier screening should be readdressed
- Carrier screening is not a test for all genetic conditions
  - will not identify de novo variants in the offspring
  - does not replace newborn screening
- When Tier 1 or Tier 2 carrier screening was performed in a prior pregnancy, Tier 3 screening should be offered
- Consanguineous couples should have Tier 4 screening
- If family history warrants, additional genes may be considered
- Negative test reduces but does not eliminate the risk of an affected child

Gregg AR et al. ACMG Professional Practice and Guidelines Committee Genetics in Medicine (2021) 23:1793–1806;  
https://doi.org/10.1038/s41436-021-01203-

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### Carrier screening- Greater Expanded Panel (176 plus)

- Larger panels that include ACOG and ACMG criteria should be considered
  - More ethnically inclusive panel
  - Moderate or higher gene–disease association – 175 of 176 conditions (99.4%)
  - Moderate to severe disease severity - 175 of 176 conditions (99.4%)
  - ACOG severity criterion
    - Determinantal effect on quality of life
    - Cognitive or physical impairment
    - Surgical or medical intervention
  - Onset early in life

Cedars Sinai Taber, et al. Genetics in Medicine (2022) 24, 201–213

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### Preimplantation genetic testing PGT-A, PGT-SR, PGT-M

**PGT-A ANEUPLOIDY**

Screens for presence of all 23 pairs of Chromosomes

Trisomy 21, 18, 13, Monosomy 7, X

**PGT-SR STRUCTURAL REARRANGEMENTS**

Translocations, Inversions, Deletions, Duplications, Insertions

**PGT-M MONOGENIC**

Used to detect known inherited disorders

Cystic Fibrosis, Tay Sachs, Sickle Cell, BRCA

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### Preimplantation genetic testing platforms

- NGS allows for direct reading of sequenced DNA fragments and their quantification based on sequence read numbers
- whole chromosome aneuploidy (PGT-A)
- medium size deletions or insertions in chromosomes (PGT-SR)
- detection of single gene disorders (PGT-M)

ESHRE PGT Consortium good practice recommendations for the detection of structural and numerical chromosomal aberrations† ESHRE PGT-SR/PGT-A Working Group 2020 Sam Behjati and Patrick S Tarpey Arch Dis Child Educ Pract Ed. 2013 Dec; 98(6): 236–238.

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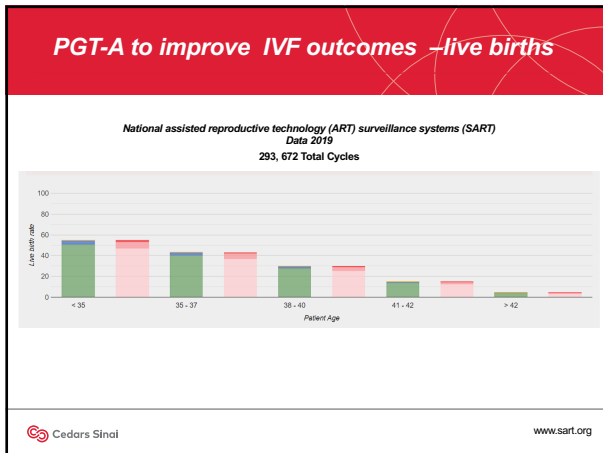
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**PGT-A - Time to pregnancy and advanced reproductive age > 37 yo**

- Analysis of data from national assisted reproductive technology (ART) surveillance systems
- PGT-A is not associated with improved rates of clinical pregnancy or live birth after fresh autologous blastocyst transfer among women aged <37 years
- PGT-A of embryos appeared to improve the likelihood of having a live birth among women >37 years
- Cycles that were intended for PGT-A were more likely to reach embryo transfer in all age groups, but more significantly in women aged >37
- RCT that focused on women with advanced maternal age (38-41 years old) demonstrated a significantly higher live birth rate with PGT-A group per cycle (36% vs 21.9%, P<0.031) and a lower miscarriage rate (2.7% vs 39%, P<0.007)

Chang et al. Fertil Steril. 2016; 105(2): 394-400.  
Practice Committees of the American Society for Reproductive Medicine and the Society for Assisted Reproductive Technology Fertil Steril 2016;109:429-38.

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**Live birth with and without PGT-A for < 38 yo (RCT)**

Yan, et al N Engl J Med 2021;385:2047-58

Outcome	PGT-A Group (N=456)	Conventional IVF Group (N=496)	Absolute Difference (95% CI)	Rate Ratio (95% CI)
<b>Primary outcomes</b>				
Cumulative live-birth rate—no. (%)	468 (77.2)	496 (81.8)	-4.6 (-9.2 to 0.0)	0.94 (0.89 to 1.00)
Singletons	469 (79.2)	478 (78.5)	-2.4 (-6.9 to 0.4)	0.97 (0.91 to 1.05)
Twin	4 (1.0)	18 (3.0)	-2.0 (-5.5 to 0.4)	0.33 (0.13 to 0.83)
<b>Secondary outcomes</b>				
Cumulative biochemical pregnancy—no. (%)	528 (86.8)	571 (94.2)	-7.4 (-10.7 to -4.2)	0.82 (0.80 to 0.96)
Cumulative clinical pregnancy—no. (%)	505 (83.3)	516 (83.7)	-4.4 (-12.1 to 4.2)	0.91 (0.87 to 0.95)
Cumulative ongoing pregnancy—no. (%)	479 (79.0)	514 (84.8)	-3.6 (-10.1 to 1.5)	0.93 (0.88 to 0.98)
<b>Birth weight</b>				
Singleton				
No. of observations	461	478		
Mean weight—g	3417.488	3449.488	-32 (-95 to 30)	
Twin				
No. of observations	12	36		
Mean weight—g	3906.714	3905.420	-105 (-444 to 235)	
<b>Cumulative pregnancy loss—no./total no. (%)</b>				
Biochemical	31/528 (5.9)	41/571 (7.2)	-3.3 (-4.2 to 1.6)	0.82 (0.52 to 1.29)
Clinical	44/505 (8.7)	72/571 (12.6)	-3.9 (-7.5 to -0.2)	0.69 (0.49 to 0.96)
First trimester	37/526 (7.0)	60/571 (10.5)	-3.5 (-6.8 to -0.1)	0.67 (0.45 to 0.99)
Second trimester	9/526 (1.7)	12/571 (2.1)	-0.4 (-2.0 to 1.2)	0.81 (0.35 to 1.92)
Good birth outcome—no. (%)	378 (62.4)	385 (63.5)	-1.2 (-4.6 to 4.3)	0.98 (0.90 to 1.07)
<b>Features of live births</b>				
Duration of pregnancy—wk	39.2±1.7	39.1±1.6	0.0 (-0.2 to 0.2)	
No. of embryos transferred	1.2±0.4	1.3±0.6	-0.2 (-0.2 to -0.1)	
No. of embryos transferred procedures	1.3±0.4	1.3±0.5	-0.1 (-0.2 to -0.1)	
Interval since randomization—mo	12.5±2.0	12.4±2.3	0.1 (-0.2 to 0.4)	
<b>Frozen embryos</b>				
No. of unused embryos	5.2±3.2	5.5±2.9	-0.3 (-0.6 to 0.1)	
No. of unused embryos in women without a live birth	4.4±2.8	4.9±2.9	-0.4 (-1.2 to 0.3)	

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### Live birth with and without PGT-A for < 38 yo (RCT)

ORIGINAL ARTICLE

Live Birth with or without Preimplantation Genetic Testing for Aneuploidy

**Table S3. The rates of pregnancy, pregnancy loss and live birth after the first embryo transfer between PGT-A and IVF.**

Outcome	PGT-A group (N=576)	IVF group (N=594)	Absolute Difference (95% CI)	Rate Ratio for PGT-A vs. IVF (95% CI)
Primary outcome: live birth-no. (%)	382/576 (66.3)	369/594 (62.1)	4.2 (-1.3, 9.7)	1.07 (0.98, 1.16)
Singleton	376/576 (65.3)	357/594 (60.1)	5.2 (-0.4, 10.7)	1.09 (0.99, 1.19)
Twin	6/576 (1.0)	12/594 (2.0)	-1.0 (-2.4, 0.4)	0.52 (0.19, 1.36)
Secondary outcomes				
Biochemical pregnancy-no. (%)	451/576 (78.3)	462/594 (77.8)	0.5 (-4.2, 5.3)	1.01 (0.95, 1.07)
Clinical pregnancy-no. (%)	422/576 (73.3)	427/594 (71.9)	1.4 (-3.7, 6.5)	1.02 (0.95, 1.09)
Ongoing pregnancy-no. (%)	393/576 (68.2)	384/594 (64.6)	3.6 (-1.8, 9.0)	1.06 (0.97, 1.15)
Pregnancy loss-no./total no. (%)				
Biochemical pregnancy loss	26/451 (5.8)	33/462 (7.1)	-1.4 (-4.6, 1.8)	0.81 (0.49, 1.33)
Clinical pregnancy loss	39/451 (8.7)	55/462 (11.9)	-3.3 (-7.2, 0.7)	0.73 (0.49, 1.07)
First trimester	30/451 (6.7)	44/462 (9.5)	-2.9 (-6.4, 0.7)	0.70 (0.45, 1.09)
Second trimester	9/451 (2.0)	11/462 (2.4)	-0.4 (-2.3, 1.5)	0.84 (0.35, 2.00)

No adjustment was made for multiplicity of secondary outcomes. 95% CIs should not be used to infer definitive treatment outcomes.

- No difference even after the first IVF cycle

Cedars Sinai Yan, et al N Engl J Med 2021;385:2047-58

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### Live birth with and without PGT-A for < 38 yo (RCT)

ORIGINAL ARTICLE

Live Birth with or without Preimplantation Genetic Testing for Aneuploidy

**Table S2. Live birth rate after each embryo transfer cycle.**

Outcome	PGT-A group (N=606)	IVF group (N=606)	Absolute Difference (95% CI)	Rate Ratio for PGT-A vs. IVF (95% CI)
Live birth after 1 <sup>st</sup> embryo transfer-no. (%)	382/576 (66.3)	369/594 (62.1)	4.2 (-1.3, 9.7)	1.07 (0.98, 1.16)
Live birth after 2 <sup>nd</sup> embryo transfer-no. (%)	7 (1.1) (0.2, 2)	10 (1.7) (0.5, 2.9)	7.0 (-4.2, 18.2)	1.13 (0.93, 1.36)
Live birth after 3 <sup>rd</sup> embryo transfer-no. (%)	5 (0.8) (0.0, 1.6)	11 (1.8) (0.8, 2.8)	1.2 (-43.8, 46.3)	1.03 (0.33, 3.19)
Live birth conceived naturally-no.	10	2	-	-

No adjustment was made for multiplicity of secondary outcomes. 95% CIs should not be used to infer definitive treatment outcomes.

- More women in the conventional-IVF group underwent a second or third embryo-transfer cycle:
  - Second Cycle - 192 women in the conventional-IVF group and 119 in the PGT-A group
  - Third Cycle - 49 women in the conventional-IVF group and 5 in the PGT-A group

Cedars Sinai Yan, et al N Engl J Med 2021;385:2047-58

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### PGT-A Retrospective Cohort Study

2464 PGT-A, 190,010 cycles

Maternal Age Group	IVF non-PGT-A	PGT-A
UNDER 35	1.27	1.15
35-37	1.34	1.08
38-39	1.46	1.11
40-42	1.67	1.08
43-44	1.77	1.12
OVER 44	1.67	1.05

- Fewer embryos are required to achieve a pregnancy following PGT-A compared to regular IVF

Sanders et al Journal of Assisted Reproduction and Genetics (2021) 38:3277-3285

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### PGT-A Retrospective Cohort Study

2464 PGT-A 190,010 cycles

Maternal Age Group	Non PGT-A (%)	PGT-A (%)
Under 35	~15	~25
35-37	~18	~28
38-39	~12	~25
40-42	~8	~22
43-44	~5	~18
Over 44	~3	~15

- PGT-A versus non PGT-A
  - Live birth rates were significantly higher in all age groups
  - Mostly single embryo transfers (SET)
  - Less number of transfers per live birth, particularly if over 40 years

Cedars Sinai | Sanders et al Journal of Assisted Reproduction and Genetics (2021) 38:3277-3285

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### PGT-A: Recommendations

- Recommendations**
  - Shortened time to pregnancy and increased success for women over 37 yo
  - Potential benefit in select populations of younger reproductive age women
  - Selection of embryo for elective single embryo transfer – Decrease risk of multiple gestations
  - Beneficial if proceeding with PGT-M or PGT-SR
  - Potential benefit for long term fertility preservation
  - Cost benefit – minimize number of frozen embryo transfer cycles?
- Considerations**
  - Would embryos that don't survive to the stage of biopsy for genetic testing lead to successful pregnancies
  - Are false positive test results possible (mosaicism 3-20%) that could lead to a healthy genetically normal pregnancy?
- Counseling is necessary for shared decision making for PGT**

Cedars Sinai | Practice Committees of the American Society for Reproductive Medicine and the Society for Assisted Reproductive Technology Fertil Steril 2018;109:429-36, 2018; Practice Committee and Genetic Counseling Professional Group (GCPG) of the American Society for Reproductive Medicine. Fertil Steril 2020;114:248-54

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### Genetics and Pregnancy – Purpose

- Prenatal testing for chromosomal abnormalities are designed to provide an accurate assessment of a patient's risk of carrying a fetus with a chromosomal disorder.
- Testing for chromosomal abnormalities should be an informed patient choice based on adequate and accurate information.
- All patients should be offered both screening and diagnostic tests, and all patients have the right to accept or decline testing after counseling.

Cedars Sinai | ACOG Practice Bulletin Screening for Fetal Chromosomal Abnormalities. 2018  
Photo: <https://www.theguardian.com/world/2021/aug/03/hopes-uk-trial-will-alleviate-pregnant-womens-covid-vaccine-concerns>

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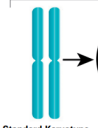
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## Pregnancy – Genomic testing capabilities

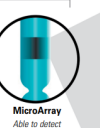
- Recommendations

**DIAGNOSTIC CAPABILITY OF PRENATAL GENETIC TESTS**

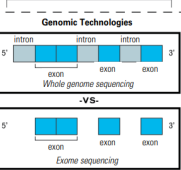
*Currently Available*                      *On the Horizon*



**Standard Karyotype**  
Able to detect large, extra or missing chromosomes (i.e. Down syndrome)



**MicroArray**  
Able to detect small deletions or duplications (i.e. 22q11 deletion syndrome)



**Genomic Technologies**  
Whole genome sequencing  
-vs-  
Exome sequencing

**Figure 1.** Diagnostic capability of prenatal genetic tests. (Reprinted from Hardisty EE, Vora NL. Advances in genetic prenatal diagnosis and screening. *Curr Opin Pediatr* 2014;26:334–8.) ←

COMMITTEE OPINION Number 682 (Reaffirmed 2020) Committee on Genetics Society for Maternal-Fetal Medicine Microarrays and Next-Generation Sequencing Technology: The Use of Advanced Genetic Diagnostic Tools in Obstetrics and Gynecology

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## Genetics and Pregnancy – Chromosomal Abnormalities

**Table 1. Chromosomal Abnormalities in Second-Trimester Pregnancies Based on Maternal Age at Term**

	Trisomy 21	Trisomy 18	Trisomy 13	Sex Chromosome Aneuploidy (XXX, XY, XYY, 45, X)	Microarray or Rare Chromosomal Abnormality	All Chromosomal Abnormalities
Age 20	8 per 10,000 1 in 1,250	2 per 10,000 1 in 5,000	1 per 10,000 1 in 10,000	34 per 10,000 1 in 294	37 per 10,000 1 in 270	82 per 10,000 1 in 122
Age 25	10 per 10,000 1 in 1,000	2 per 10,000 1 in 5,000	1 per 10,000 1 in 10,000	34 per 10,000 1 in 294	37 per 10,000 1 in 270	84 per 10,000 1 in 119
Age 30	14 per 10,000 1 in 714	4 per 10,000 1 in 2,500	2 per 10,000 1 in 5,000	34 per 10,000 1 in 294	37 per 10,000 1 in 270	91 per 10,000 1 in 110
Age 35	34 per 10,000 1 in 294	9 per 10,000 1 in 1,111	4 per 10,000 1 in 2,500	35 per 10,000 1 in 285	37 per 10,000 1 in 270	119 per 10,000 1 in 84
Age 40	116 per 10,000 1 in 86	30 per 10,000 1 in 333	14 per 10,000 1 in 714	51 per 10,000 1 in 196	37 per 10,000 1 in 270	248 per 10,000 1 in 40

ACOG Practice Bulletin Screening for Fetal Chromosomal Abnormalities. 2018

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## Microdeletions, Duplications and other Variants

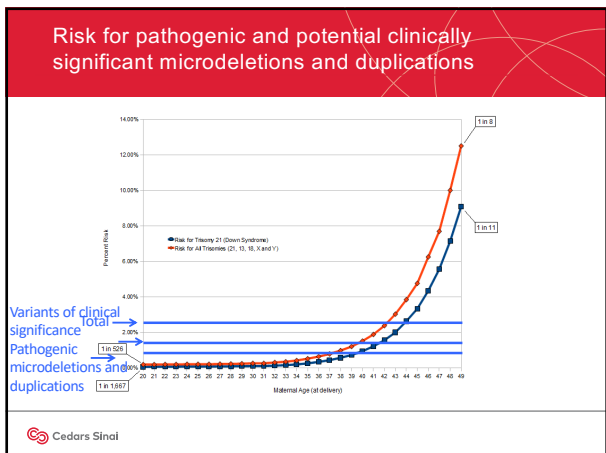
**Table 3. Frequency and Clinical Interpretation of Microdeletions and Duplications on Chromosomal Microarray in the 3822 Samples with a Normal Karyotype, According to Indication for Prenatal Testing.**

Indication for Prenatal Diagnosis	Normal Karyotype no.	Common Benign no. (%)	Uncertain Clinical Significance (N=130)		Total Known Pathogenic and Potential for Clinical Significance <sup>a</sup> no. (%) [95% CI] <sup>b</sup>	
			Pathogenic no. (%)	Likely to Be Benign no. (%)		
						Potential for Clinical Significance
Any	3822	1234 (32.3)	45 (0.9)	69 (1.8)§	61 (1.6)	96 (2.5) [1.1–3.1]
Advanced maternal age	1966	628 (31.9)	9 (0.5)	37 (1.9)	25 (1.3)	34 (1.7) [1.2–2.4]
Positive on Down's syndrome screening	729	247 (33.9)	3 (0.4)	13 (1.8)	9 (1.2)	12 (1.6) [0.9–2.9]
Anomaly on ultrasonography	755	247 (32.7)	21 (2.8)	16 (2.1)	24 (3.2)	45 (6.0) [4.5–7.9]
Other <sup>c</sup>	372	112 (30.1)	2 (0.5)	3 (0.8)	3 (0.8)	5 (1.3) [0.6–3.1]

<sup>a</sup> Total includes those predetermined as known to be pathogenic and those classified by the clinical advisory committee as clinically relevant.  
<sup>b</sup> CI denotes confidence interval.  
<sup>c</sup> Includes 36 samples determined likely to be benign by the study geneticist and 33 determined by the independent clinical advisory committee on the basis of size, gene content, inheritance, the literature, and ultrasonography findings.  
<sup>d</sup> Other indications include family history, previous pregnancy with chromosomal abnormalities, and elective decision.

Wapner, et al. December 6, 2012N Engl J Med 2012; 367:2175-2184 DOI: 10.1056/NEJMe1203382 30

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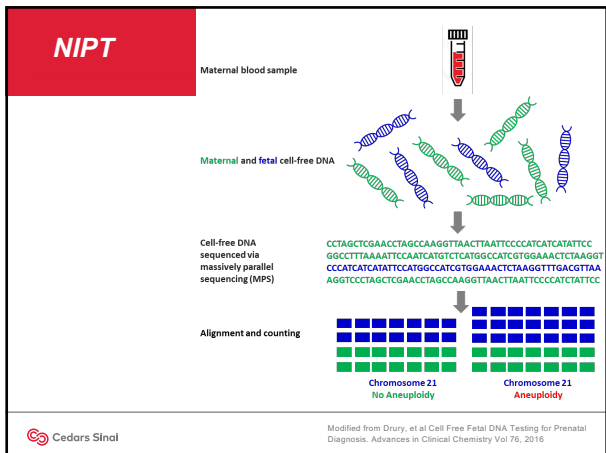
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### Pregnancy – NIPT high risk populations

Chromosomal Abnormality	Sensitivity (%)	95% CI (%)	Specificity (%)	95% CI (%)
<b>Trisomy 21</b>	<b>99.5</b>	<b>96.3-99.9</b>	<b>100</b>	<b>99.87-100</b>
<b>Trisomy 18</b>	<b>97.7</b>	<b>87.9-99.6</b>	<b>99.97</b>	<b>99.81-99.99</b>
<b>Trisomy 13</b>	<b>100</b>	<b>83.2-100</b>	<b>99.97</b>	<b>99.81- 99.99</b>

- High sensitivity and high specificity
- Not reportable or no call results – increased risk of chromosomal abnormality – diagnostic testing is recommended

**Source:** Geppert, et al Prenatal Diagnosis. 2020;40:454–462. ACOG Practice Bulletin Screening for Fetal Chromosomal Abnormalities. 2018. Cedars Sinai

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### Pregnancy – NIPT low risk population

- Low risk population
- 13,043 (73.1%) were considered low-risk for aneuploidy < 35
- 3,873 that were ≥35 but had a low-risk result on a blood screening test

Chromosomal abnormality	Sensitivity % (n)	Specificity % (n)	PPV % (n)	NPV % (n)
<b>Trisomy 21</b>	100 (18/18)	99.98 (12,815/12,818)	85.71 (18/21)	100 (12,815/12,815)
<b>Trisomy 18</b>	75 (3/4)	99.98 (12,829/12,832)	50 (3/6)	99.99 (12,829/12,830)
<b>Trisomy 13</b>	100 (5/5)	99.98 (12,828/12,831)	62.50 (5/8)	100 (12,828/12,828)

Cedars Sinai Dar, et al American Journal of Obstetrics and Gynecology (2022), doi: https://doi.org/10.1016/j.ajog.2022.01.019

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### Pregnancy – NIPT

**Table 3.** The Effect of Maternal Age on the Positive Predictive Value of Cell-Free DNA Screening for Trisomy 21, 18, and 13 at 10 Weeks Gestation

	Maternal Age	Age Related Risk <sup>1</sup>	Positive Predictive Value <sup>2</sup>
Trisomy 21	20	1:804 or 12 per 10,000	38–80%
	35	1:187 or 53 per 10,000	73–95%
	40	1:51 or 196 per 10,000	91–99%
Trisomy 18	20	1:1,993 or 5 per 10,000	11–41%
	35	1:465 or 22 per 10,000	34–75%
	40	1:126 or 79 per 10,000	66–92%
Trisomy 13	20	1:6,347 or 16 per 10,000	5–13%
	35	1:481 or 7 per 10,000	17–40%
	40	1:401 or 24 per 10,000	43–71%

<sup>1</sup>Sensitivity and specificity approximately 99%  
<sup>2</sup>Age related risk of aneuploidy per 10,000 pregnancies at 10 weeks gestation based on maternal age at term  
<sup>3</sup>Percent varies by laboratory

Adapted from University of North Carolina at Chapel Hill. Positive predictive value of cell free DNA calculator. Available at: <https://www.medunc.edu/mm/nips-calc>. Retrieved February 24, 2020.

- Low positive predictive value means many false positive test results

Cedars Sinai ACOG Practice Bulletin Screening for fetal Chromosomal Abnormalities 2018

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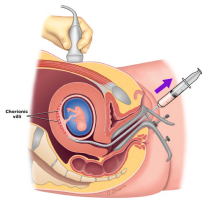
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### Pregnancy – Diagnostic Testing

- Chorionic villus sampling
- Karyotype and microarray
- Detects 99.8% of trisomies, pathogenic microdeletions/duplications, clinically significant variants, point mutations, chromosomal rearrangements and de novo mutations
- Performed between 10-13 weeks
- Miscarriage rate overall - 0.5-3.0%
- Procedure-related risk of miscarriage 0.22% = 1/500



Cedars Sinai ACOG Committee Opinion Preimplantation Genetic Testing 2020 Akolekar, et al Ultrasound Obstet Gynecol 2015; 45: 16–26

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### Pregnancy – Diagnostic Testing

- Amniocentesis
- Karyotype preferred for balanced translocations and triploidy
- Performed between 15-20 weeks
- Miscarriage rate overall - 0.5-1.0%
- Procedure-related risk of miscarriage 0.11% = 1/900

ACOG Committee Opinion Preimplantation Genetic Testing 2020  
Akolekar, et al Ultrasound Obstet Gynecol 2015; 45: 16-26  
Image courtesy -UCLA MFM website

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### Pregnancy – Prenatal testing

- Testing for chromosomal abnormalities should be an informed patient choice based on adequate and accurate information
- All patients should be offered both screening and diagnostic tests, and all patients have the right to accept or decline testing after counseling
- Due to the background rate of pathogenic microdeletions/duplications and clinically significant variants (2.5%) - chromosomal microarray analysis through diagnostic testing should be offered to all women regardless of age
- Diagnostic testing/chromosomal microarray is recommended for a fetus with a structural abnormality on ultrasound
- Procedure related risk of loss (0.11-0.22%) should be addressed with the patient
- At this time, NIPT is a screening test best suited ONLY for identification of aneuploidies (Trisomy 21, 18, and 13?) in high-risk populations

COMMITTEE OPINION Number 682 Committee on Genetics Society for Maternal-Fetal Medicine Microarrays and Next-Generation Sequencing Technology: The Use of Advanced Genetic Diagnostic Tools in Obstetrics and Gynecology 2020

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### Preimplantation Genetic Testing – Now I am pregnant, what's next?

- A normal or negative PGT result is not a guarantee of a newborn without genetic abnormalities.
- Traditional diagnostic testing or screening for aneuploidy should be offered to all patients who have PGT-A, in accordance to recommendations for all pregnant patients
- Confirmation of preimplantation genetic testing – monogenic results with CVS or amniocentesis should be offered
- PGT-SR to detect structural chromosomal abnormalities such as translocations - Confirmation of preimplantation genetic testing – and confirmation of unaffected or balanced translocation in offspring via CVS or amniocentesis should be offered,
- Limitations of PGT – do not detect microdeletions and microduplications, de novo variants, and imprinting disorders
- **PGT and NIPT remain only as screening tests!**

ACOG Committee Opinion Preimplantation Genetic Testing 2020

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**Genetic testing - Beyond**

Cedars Sinai

Gavril P, et al 1993 *Pediatr Pathol* 13:453-462  
 Zhang Y, et al 2008 *Proteomics* 8:4344-4356  
 Collier AC, et al 2009 *J Steroid Biochem Mol Biol* 116:21-28

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**Next-Generation Prenatal Diagnostics**

1. Target: Trophoblasts (TBs)

4. 4 to 8-mL blood per test

2. A new mechanism for rare-cell sorting: **NanoVelcro Chip with imprinted PLGA nanostructures**

3. Three types of downstream analyses: **FISH, microarray, and WGS**

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Crump *bioRxiv* **Medicine & Imaging**

Hou S., Pisarska, Zhu, Tseng. *ACS Nano*. 2017 Aug 22;11(8)

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**Circulating Trophoblast Cell Clusters for Early Detection of Placenta Accreta Spectrum Disorder**

**NanoVelcro Chips for Detecting cTBs and cTB clusters**

Serosa  
Myometrium  
Endometrium  
Accreta  
Increta  
Percreta

Pregnant woman with PAS

Blood

NanoVelcro Chips for capturing both single cTBs and clustered cTBs

PDMS chaotic mixer

Anti-EpCAM-coated SINWS

Confirmation of cTBs' trophoblast origin by detecting trophoblast-specific genes

Single cTBs and clustered cTBs are captured on SINWS

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Crump *bioRxiv* **Medicine & Imaging**

Alshar, Zhu, Pisarska, Tseng. *Nat Commun* 2021

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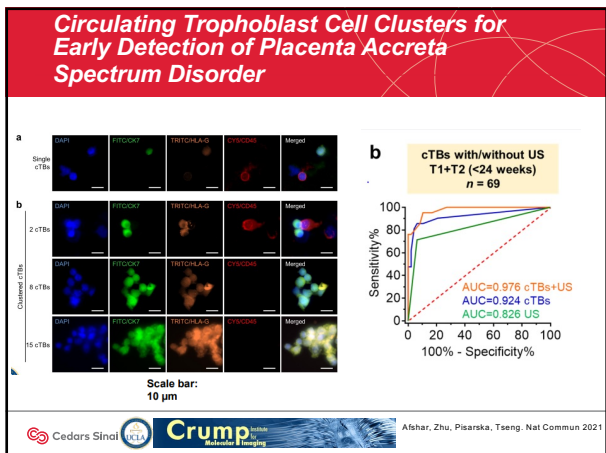
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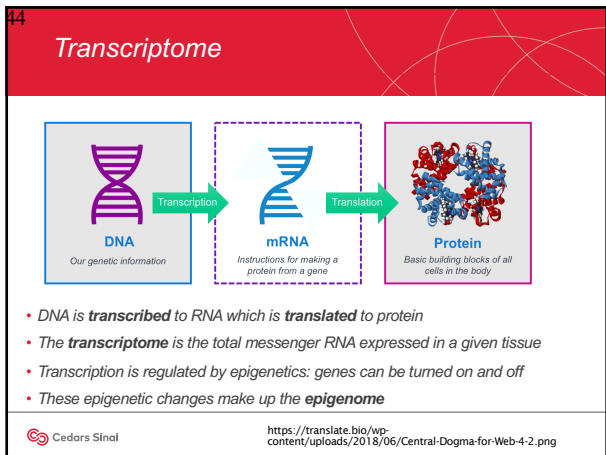
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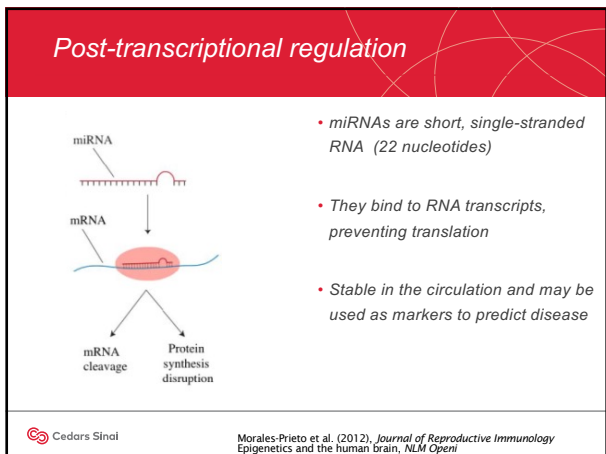
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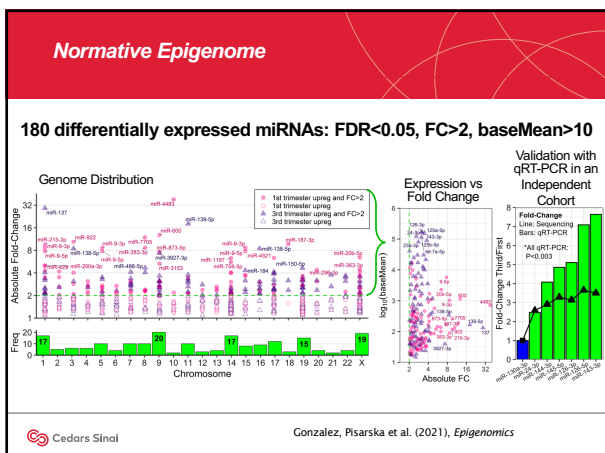
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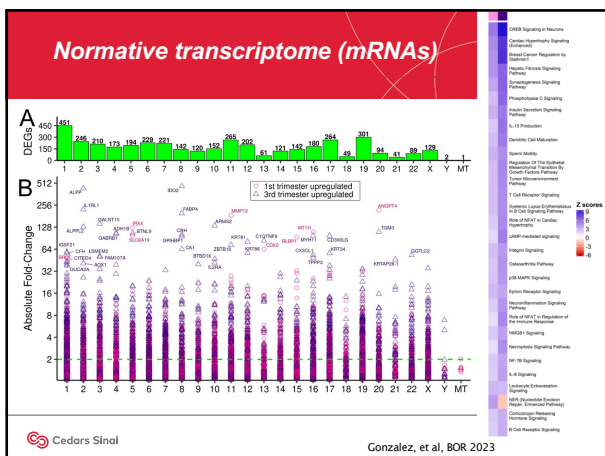
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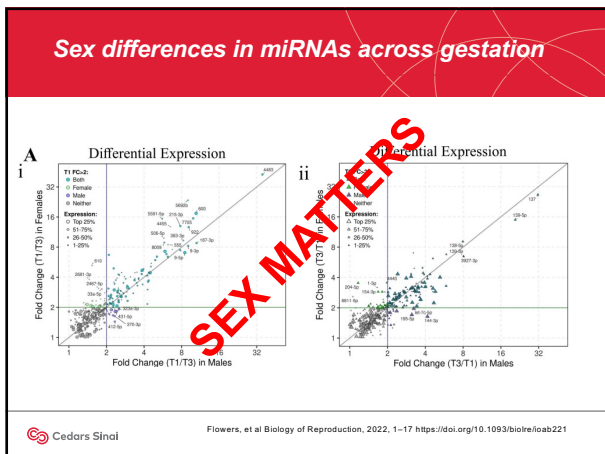
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## Conclusion

- **Preconception**
  - Current ACOG recommendations are limited – based on advances in NGS and recent recommendations by ACMG, carrier screening should screen a minimum of 74-112 genetic conditions
  - When utilizing commercially available genetic screening – the same panels should be performed for both genetic parents
- **Prenatal**
  - IVF/PGT testing does not replace prenatal genetic counseling with genetic screening and/or diagnostic testing
  - NIPT is currently only recommended for high-risk populations for aneuploidy screening (Trisomy 21, 18, and ?13)
  - Pathogenic microdeletions/duplications and clinically significant variants affect 2.5% of pregnancies regardless of maternal age
  - Diagnostic testing through CVS or amniocentesis should be offered to pregnant patients regardless of age and previous genetic screening
- **Future**

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## Acknowledgements

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  - Yalda Alshar, MD, PhD

**Our patients for participating in our studies to improve outcomes!**

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