

ISUOG VIRTUAL INTERNATIONAL SYMPOSIUM 2021

State-of-the-art Ultrasound Imaging in Obstetrics and Gynecology

17-18 April 2021

REGISTER NOW ►

Using ultrasound together with other technologies to improve the lifelong health of women and babies

- Two streams of scientific content over two days, delivered through the ISUOG virtual platform which will exceed your expectations
- A mixture of lectures and practical, interactive training, including scan demonstrations, pattern recognition sessions and case report discussion
- Leading international and local experts in obstetrics, gynecology and imaging
- Live program delivered from 7:30 - 18:30 Calgary, Canada time (Mountain Daylight Time)
- Content available on Demand, at a time, pace and location to suit you until 17 May 2021
- All non-member registration fees include a 12-month ISUOG basic membership

Provisional program

Sessions will run simultaneously, providing two streams of content both days.

Highlights include:

- Obstetrics: the first trimester, beyond the routine mid-trimester fetal ultrasound scan, screening to improve pregnancy outcomes, fetal growth and health, ultrasound in labor, and more
- Gynecology: ectopic pregnancy, miscarriage, endometriosis, menopause, ovarian tumors, tubal and uterine pathology, and more
- Special sessions include advanced imaging/MRI, fetal therapy and COVID

The symposium will be co-chaired by:

*Jo-Ann Johnson (Canada),
Denise Pugash (Canada)*

Symposium Advisory Group

*Shabnam Bobdiwala (UK)
George Condous (Australia)
Karen Fung-Kee-Fung (Canada)
Jon Hyett (Australia)
Simon Meagher (Australia)
Liona Poon (Hong Kong)
Angela Ranzini (USA)
Magdalena Sanz Cortes (USA)*

Who should attend?

This interactive course is designed for Maternal Fetal Medicine (MFMs), OB-GYNs, Radiologists, Sonographers, Geneticists, Researchers, Trainees/Residents and other maternity care providers. The program will appeal to a wide global audience, with a focus on North American educational needs.

See you ONLINE in 2021! For more information, please visit:
isuog.org/event/17th-isuog-international-symposium.html



Opinion

Genome-wide cfDNA testing of maternal blood

J. C. JANI^{1*}, M. M. GIL² , A. BENACHI³,
F. PREFUMO⁴, K. O. KAGAN⁵, A. TABOR⁶,
C. M. BILARDO⁷, G. C. DI RENZO⁸ and
K. H. NICOLAIDES⁹

¹Department of Obstetrics and Gynecology, University Hospital Brugmann, Université Libre de Bruxelles, Brussels, Belgium; ²Obstetrics and Gynecology Department, Hospital Universitario de Torrejón, Torrejón de Ardoz, Universidad Francisco de Vitoria, Madrid, Spain; ³Department of Obstetrics and Gynecology, Antoine-Béclère Hospital, Assistance Publique-Hôpitaux de Paris (AP-HP), Université Paris-Sud, Clamart, France; ⁴Division of Obstetrics and Gynecology, Department of Clinical and Experimental Sciences, University of Brescia, Brescia, Italy; ⁵Tuebingen University Hospital, Obstetrics and Gynaecology, Tuebingen, Germany; ⁶Center of Fetal Medicine, Department of Obstetrics, Copenhagen University Hospital Rigshospitalet, Copenhagen, Denmark; ⁷Department of Obstetrics & Gynaecology, Amsterdam University Medical Centers, Amsterdam, The Netherlands; ⁸Department of Obstetrics and Gynecology, Centre for Perinatal and Reproductive Medicine, University Hospital, University of Perugia, Perugia, Italy; ⁹Fetal Medicine Research Institute, King's College Hospital, London, UK
*Correspondence. (e-mail: jackjani@hotmail.com)

Measurement of cell-free(cf) DNA in maternal blood has been shown to provide effective prediction of fetal trisomy 21 and, to a lesser extent, of trisomies 18 and 13, both in singleton and in twin pregnancy^{1,2}. This has led to clinical implementation of the test in several countries, usually in women identified through prior screening by the first-trimester combined test to have a high or moderate risk for trisomy 21. In Belgium and The Netherlands, however, cfDNA testing is being offered to all pregnant women, as an alternative to the first-trimester combined test, and the test is based on genome-wide (GW) analysis rather than being confined to screening for the three major trisomies. The rationale for such a policy is that GW testing has the potential to diagnose clinically significant rare autosomal trisomies (RATs) and rare additional fetal segmental imbalances (SIs).

The results from the first year of GW-cfDNA testing in The Netherlands (TRIDENT-2 study) included 56 818 women who underwent GW-cfDNA testing, from an initial cohort of 73 239 women who had a cfDNA test; in 207 (0.4%) of these women, the test was positive for RATs ($n = 101$), SIs ($n = 95$) or complex abnormal profiles ($n = 11$)³. Among the 101 RATs, six were confirmed but only one of these was associated with an abnormal phenotype. Among the 95 SIs, 29 were confirmed but the number with abnormality not discoverable through

ultrasound was not defined. In another seven cases consistent with maternal malignancy or premalignancy, the benefit of the discovery was not demonstrated. An abnormal test result inevitably leads to anxiety and, in some cases, to termination, as well as the need for both fetal and maternal testing; however, even when the fetal karyotype is found to be normal after a positive RAT result, uncertainty persists as to whether there are true mosaicisms in crucial fetal tissues and organs. When an invasive procedure confirms a true fetal mosaicism after a positive RAT result, it is impossible to predict clinical outcome and, in case of confined placental mosaicism (CPM), except CPM for trisomy 16, there is evidence that the incidence of adverse pregnancy outcome in an unselected population is not different from that in pregnancies with normal karyotype at chorionic villus sampling (CVS)⁴. Therefore, TRIDENT-2 shows that, at present, the benefits of screening for all genetic imbalances do not seem to outweigh the potential harms and that clinical implementation, even in a research setting, may be questionable ethically.

A study in Belgium, involving 3373 women, reported that GW-cfDNA testing identified additional findings beyond the common trisomies in 28 (0.8%) cases; these included four sex-chromosome aneuploidies, six RATs and one rare autosomal monosomy, none of which was confirmed in the fetus or the neonate, as well as 17 large or sub-microscopic SIs, of which three were confirmed in amniocytes⁵. In all 28 cases, the clinical follow-up was normal. Benn *et al.* reviewed the types of RAT identified following CVS, as reported in 10 recently published cfDNA studies, and found that the clinical outcome of cases with cfDNA analysis positive for RATs mostly involved the birth of an apparently normal baby (40%) or a miscarriage/fetal loss (27%), for which screening tests are not recommended⁶. There was a weak association between RATs and pregnancy complications, such as fetal growth restriction and fetal abnormalities, in the tested population.

There are several points of concern that arise from GW-cfDNA testing.

- 1) Increase in the screen-positive rate of a test that was initially meant to reduce it, and increase in the rate of invasive testing for conditions of unknown clinical significance that remain of unknown significance even after an invasive procedure.
- 2) There is uncertainty as to the clinical significance of a heterogeneous set of chromosomal abnormalities and how best to manage a positive result. Consequently, no professional society currently recommends this test^{7–11}.
- 3) There is heterogeneity of home-brew massively parallel shotgun sequencing protocols.

- 4) There are ethical and legal challenges to overcome regarding how best to counsel parents before they give their informed consent, since accurate information is lacking. In fact, women are already undergoing GW-cfDNA screening without clear information about its limitations and drawbacks, and clinical decisions are already being made based on results of uncertain clinical significance^{12–14}. There are also ethical concerns regarding increased voluntary termination of pregnancy due to positive RAT results even after a normal karyotype and normal ultrasound scan.
- 5) The test violates World Health Organization screening principles¹⁵.

In conclusion, although research should always be encouraged, the benefits *vs* harms of implementation of GW-cfDNA screening must be weighed carefully. Healthcare providers and grant-awarding bodies have a responsibility to ensure that more robust data and management strategies are available before endorsing studies or strategies incorporating GW-cfDNA testing into nationally reimbursed screening programs.

REFERENCES

1. Gil MM, Galeva S, Jani J, Konstantinidou L, Akolekar R, Plana MN, Nicolaides KH. Screening for trisomies by cfDNA testing of maternal blood in twin pregnancy: update of The Fetal Medicine Foundation results and meta-analysis. *Ultrasound Obstet Gynecol* 2019; 53: 734–742.
2. Gil MM, Accurti V, Santacruz B, Plana MN, Nicolaides KH. Analysis of cell-free DNA in maternal blood in screening for aneuploidies: updated meta-analysis. *Ultrasound Obstet Gynecol* 2017; 50: 302–314.
3. van der Meij KRM, Sistermans EA, Macville MVE *et al*. TRIDENT-2: National Implementation of Genome-Wide Non-Invasive Prenatal Testing as a First-Tier Screening Test in the Netherlands. *Am J Hum Genet* 2019; 105: 1091–1101.
4. Grati FR, Ferreira J, Benn P, Izzi C, Verdi F, Vercellotti E, Dalpiaz C, D'Ajello P, Filippi E, Volpe N, Malvestiti F, Maggi F, Simoni G, Frusca T, Cirelli G, Bracalente G, Re AL, Surico D, Ghi T, Prefumo F. Outcomes in pregnancies with a confined placental mosaicism and implications for prenatal screening using cell-free DNA. *Genet Med* 2019. DOI: 10.1038/s41436-019-0630-y.
5. de Wergifosse S, Bevilacqua E, Mezela I, El Haddad S, Gounongbe C, de Marchin J, Maggi V, Conotte S, Badr DA, Fils J-F, Guizani M, Jani JC. Cell-free DNA analysis in maternal blood: comparing genome-wide versus targeted approach as a first-line screening test. *J Matern Fetal Neonatal Med* 2019. DOI: 10.1080/14767058.2019.1686478.
6. Benn P, Malvestiti F, Grimi B, Maggi F, Simoni G, Grati FR. Rare autosomal trisomies: comparison of detection through cell-free DNA analysis and direct chromosome preparation of chorionic villus samples. *Ultrasound Obstet Gynecol* 2019; 54: 458–467.
7. Gregg AR, Skotko BG, Benkendorf JL, Monaghan KG, Bajaj K, Best RG, Klugman S, Watson MS. Noninvasive prenatal screening for fetal aneuploidy, 2016 update: a position statement of the American College of Medical Genetics and Genomics. *Genet Med* 2016; 18: 1056–1065.
8. Benn P, Borrell A, Chiu RW, Cuckle H, Dugoff L, Faas B, Gross S, Huang T, Johnson J, Maymon R, Norton M, Odibo A, Schielen P, Spencer K, Wright D, Yaron Y. Position statement from the Chromosome Abnormality Screening Committee on behalf of the Board of the International Society for Prenatal Diagnosis. *Prenat Diagn* 2015; 35: 725–734.
9. Dondorp W, de Wert G, Bombard Y, Bianchi DW, Bergmann C, Borry P, Chitty LS, Fellmann F, Forzano F, Hall A, Henneman L, Howard HC, Lucassen A, Ormond K, Peterlin B, Radojkovic D, Rogowski W, Soller M, Tibben A, Tranebjærg L, van El CG, Cornel MC; European Society of Human Genetics; American Society of Human Genetics. Non-invasive prenatal testing for aneuploidy and beyond: challenges of responsible innovation in prenatal screening. *Eur J Hum Genet* 2015; 23: 1438–1450.
10. Society for Maternal Fetal Medicine (SMFM) Publications Committee. Prenatal aneuploidy screening using cell-free DNA. Consult Series #36. *Am J Obstet Gynecol* 2015; 212: 711–716.
11. Society for Maternal-Fetal Medicine (SMFM), Norton ME, Biggio JR, Kuller JA, Blackwell SC. The role of ultrasound in women who undergo cell-free DNA screening. *Am J Obstet Gynecol* 2017; 216: B2–7.
12. Centrum Menselijke Erfelijkheid. Non-invasive prenatal testing (NIPT) request form. https://www.uzleuven.be/sites/default/files/NIPT_ENG_UZ_form.pdf.
13. Centrum voor Medische Genetica. <http://www.brusselsgenetics.be/default.aspx>.
14. Benn P, Grati FR, Ferreira J. Response to Sistermans *et al*. *Genet Med* 2019. DOI: 10.1038/s41436-019-0687-7.
15. Di Renzo GC, Bartha JL, Bilardo CM. Expanding the indications for cell-free DNA in the maternal circulation: clinical considerations and implications. *Am J Obstet Gynecol* 2019; 220: 537–542.