1	Screening for Fetal Growth Restriction using Fetal Biometry Combined with Maternal
2	Biomarkers
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1 **Condensation:**

- 2 Combining fetal biometry and biomarkers reflective of placental insufficiency will help
- 3 developing screening tests able to differentiate between healthy and growth restricted SGA
- 4 fetuses
- 5
- 6 **Short title:**
- 7 Screening for fetal growth disorders

1 Abstract

2 Fetal growth restriction (FGR) is a major determinant of perinatal morbidity and mortality. 3 Screening for FGR is a key element of prenatal care but it is recognized to be problematic. 4 Screening using clinical risk assessment and targeting ultrasound to high risk women is the 5 standard of care in the USA and UK, but the approach is known to have low sensitivity. 6 Systematic reviews of randomized controlled trials do not demonstrate any benefit from 7 universal ultrasound screening for FGR in the third trimester, but the evidence base is not 8 strong. Implementation of universal ultrasound screening in low risk women in France failed 9 to reduce the risk of complications among small for gestational age (SGA) infants but did 10 appear to cause iatrogenic harm to false positives. One strategy to making progress is to 11 improve screening by developing more sensitive and specific tests with the key goal of 12 differentiating between healthy small fetuses and those which are small through FGR. As 13 abnormal placentation is thought to be the major cause of FGR, one approach is to combine 14 fetal biometry with an indicator of placental dysfunction. In the past, these indicators were 15 generally ultrasonic measurements, such as Doppler flow velocimetry of the utero-placental 16 circulation. However, another promising approach is to combine ultrasonic suspicion of an 17 SGA infant with a blood test indicating placental dysfunction. Thus far, much of the research 18 on maternal serum biomarkers for FGR has involved the secondary analysis of tests 19 performed for other indications, such as fetal aneuploidies. An exemplar of this is pregnancy 20 associated plasma protein A (PAPP-A). This blood test is performed primarily to assess the risk 21 of Down syndrome, but women with low first trimester levels are now serially scanned in later 22 pregnancy due to associations with placental causes of stillbirth, including FGR. The 23 development of "omic" technologies presents a huge opportunity to identify novel 24 biomarkers for FGR. The hope is that when such markers are measured alongside ultrasonic

fetal biometry, the combination would have strong predictive power for FGR and its related complications. However, a series of important methodological considerations in assessing the diagnostic effectiveness of new tests will have to be addressed. The challenge thereafter will be to identify novel disease-modifying interventions, which are the essential partner to an effective screening test in order to achieve clinically effective population based screening. Key words: Small for gestational age; Ultrasound; Placenta; Biomarker; Screening; Stillbirth; Review; fetal biometry; placental growth factor; PAPP-A; hCG; sFLT1; sEndoglin; PP13; Adam12; alpha fetoprotein; inhibin; human placental lactogen; SGA; models; prediction; fetal death; RCT; study design

1 Fetal growth restriction and small for gestational age: differences between two commonly

2 used terms and clinical implications

3 Fetal growth restriction (FGR) is defined as the failure of the fetus to reach its genetically 4 determined growth potential. FGR is a major determinant of perinatal and childhood 5 morbidity and mortality, and is also associated with the risk of chronic diseases in later life.¹⁻ 6 ³ An obstacle to the study of FGR is that there are no gold standard definition and diagnostic 7 criteria for this condition. The size of the fetus or newborn is quantified with reference to the normal range for gestational age and those with birthweight (BW) less than the 10th percentile 8 9 are called small for gestational age (SGA). Inaccurately, the small size of the baby often 10 becomes synonymous with FGR, and different thresholds for these measurements are used to "define" a FGR infant (e.g. <2500 g, <10th percentile, or <3th percentile). 11

12 Although SGA and FGR are sometimes used interchangeably, the two terms are distinct, as 13 many SGA infants are constitutionally small and healthy. Hence, clinical research on screening 14 for FGR has to address two main issues: (i) the sensitive and specific detection of SGA fetuses, 15 and (ii) the ability to discriminate between FGR and healthy SGA. The causes of FGR can be 16 broadly categorized into maternal (e.g. pregnancy-associated hypertensive diseases, autoimmune disease, poor nutrition, substance abuse and teratogen exposure),⁴⁻⁶ fetal (e.g. 17 multiple gestations, infections, genetic and structural disorders)^{7, 8} or placental. It is thought 18 that placental dysfunction accounts for the majority of FGR cases.⁹ Hence, one the most 19 20 promising approaches to screening for FGR is to combine ultrasonic fetal biometry with 21 measurement of biomarkers of abnormal placentation in the mother's blood.

22 *Current status of screening with fetal biometry*

1 In many countries, including the UK and USA, ultrasound scanning after the 20 week anomaly 2 scan is only performed on the basis of clinical indications as universal ultrasound is not supported by the most recent Cochrane review.¹⁰ It is worth noting that the evidence base 3 4 can be described as an absence of evidence rather than compelling high-quality evidence of 5 the absence of clinical effectiveness of screening. This is due to a number of problems with 6 the 13 studies analyzed in the systematic review, including limited statistical power and lack 7 of an effective interventional strategy.¹¹ Nevertheless, the current approach to screening for 8 FGR is to assess the women for pre-existing risk factors, acquired complications of pregnancy 9 and clinical examination (e.g. symphysis-fundal height [SFH] measurements) (Fig. 1). Women 10 identified as high risk using these methods are then selected for ultrasonographic assessment. Screening for FGR is just one element of the universal ultrasound.¹² Other 11 12 elements include macrosomia, late presentation of fetal anomalies, abnormalities of amniotic 13 fluid volume and diagnosis of undetected malpresentation.

14

15 Ultrasonic markers of FGR

16 Fetal biometry and Doppler flow velocimetry are the primary methods used currently to 17 diagnose FGR. The use of ultrasound markers of FGR is discussed in detail elsewhere in this 18 issue, and will be only briefly summarized here. An estimated fetal weight (EFW) is derived 19 from ultrasonic measurements of head size, abdominal circumference and femur length, and 20 an EFW centile is calculated using a reference standard.^{13, 14} While a single measurement of 21 fetal size and the EFW<10th centile cut-off appears to be insufficient to discriminate growth 22 restricted and healthy small fetuses, serial fetal biometry reveals the growth trajectory of the fetus, and this helps differentiate between healthy SGA and FGR.^{15, 16} Doppler flow 23

velocimetry provides information on the resistance to blood flow in the feto-placental unit and it features in several proposed FGR definitions.¹⁷ High resistance patterns of flow in the uterine and umbilical arteries in early and mid-pregnancy have been associated with an increased risk of preeclampsia, FGR and stillbirth.¹⁸⁻²² Other measurements associated with adverse pregnancy outcomes are middle cerebral artery (MCA) and ductus venosus flow resistance, and cerebroplacental ratio (CPR) (reviewed elsewhere).^{17, 18, 23}

7

8 Biochemical biomarkers for FGR

9 Abnormal placentation leads to inadequate remodeling of maternal spiral arteries, altered 10 uteroplacental blood perfusion and impaired materno-fetal exchange of nutrients, gases and 11 waste products. These defects, collectively referred as placental insufficiency, are thought to 12 be underlying mechanisms of placentally-related complications including FGR, preeclampsia 13 and stillbirth. Hence, biochemical markers reflective of placental insufficiency become 14 attractive tools to identify women at risk of these adverse pregnancy outcomes (Figure 1, 15 Table 1).

16 First trimester screening

17 It is increasingly recognized that placental dysfunction leading to disease in the second half 18 of pregnancy has its origins in the first trimester of pregnancy.²⁴ Studies of associations have 19 been facilitated by the secondary analysis of first trimester biomarkers derived from the 20 placenta, which were evaluated in screening studies for aneuploidies.

21 Down syndrome markers: PAPP-A and fb-hCG

Low maternal circulating levels of pregnancy associated plasma protein A (PAPP-A) and high
 concentrations of the free beta subunit of human chorionic gonadotropin (fβ-hCG) are both

1 associated with the risk of Down syndrome.²⁵ PAPP-A determines the availability of the 2 insulin-like growth factors (IGFs), key pregnancy growth hormones, as it is a protease that 3 acts on IGF binding proteins. A causal role for PAPP-A in controlling fetal growth has been established in the PAPP-A knock out mouse.²⁶ In women, low serum concentrations of PAPP-4 5 A in the first trimester are associated with an increased risk of FGR, preterm delivery, 6 preeclampsia and stillbirth.²⁷⁻³³ The last of these associations is particularly strong for stillbirth associated with placental dysfunction (preeclampsia, FGR and abruption).³⁴ In the UK, and 7 8 many other countries, low first trimester PAPP-A levels are an indication for late pregnancy ultrasonic assessment of fetal growth.^{35, 36} Human chorionic gonadotrophin (hCG) is 9 predominantly produced by the placental syncytiotrophoblast cells.³⁷ hCG is a glycoprotein 10 11 composed of an α -subunit (common to luteinizing hormone, follicle stimulating hormone and 12 thyroid stimulating hormone) and a β -subunit (unique to hCG). In first trimester Down syndrome screening the f β -hCG is measured.²⁵ In general, extremes of f β -hCG in the first 13 trimester are less strongly associated with adverse outcome than low PAPP-A.^{27, 30-33} 14

15 Other placental markers: PIGF, sFLT1, sENG, PP-13 and ADAM-12

16 While many of the largest studies of first trimester markers have focused on secondary 17 analysis of Down syndrome screening research, other investigators have focused on 18 measuring proteins on the basis of a known role in placentation. Angiogenic factors play a key 19 role in the extensive vasculature remodeling of the uterus during pregnancy. The placenta 20 itself produces several factors with pro- or anti-angiogenic activity and regulation of their 21 expression and secretion is necessary for optimal placentation, maternal adaptation to 22 pregnancy and, consequently, fetal development and growth. Preeclampsia-like changes 23 were induced in pregnant rats by adenoviral mediated expression of soluble fms-like tyrosine

1 kinase-1 (sFLT1)³⁸ or soluble endoglin (sENG) alone or in combination with sFLT1.³⁹ Placental 2 growth factor (PIGF) is a pro-angiogenic factor highly expressed in placenta throughout all 3 stages of pregnancy. It is readily detectable in maternal circulation where it may have direct 4 effects on endothelial maintenance and well-being. Consistent with this role, low first 5 trimester levels of this factor have been shown to be associated with an increased risk of later 6 adverse perinatal outcome, including preeclampsia and SGA.^{31, 40, 41} The results are variable 7 for anti-angiogenic factors. High maternal levels of sENG in the first trimester were associated with preeclampsia and SGA.⁴¹ However, results are less consistent for sFLT1.^{40, 42, 43} A large 8 scale study employing correction of analyte levels using multiples of the median actually 9 10 demonstrated that low sFLT1 levels were associated with an increased risk of SGA, preterm birth and stillbirth⁴⁰ whereas data in later pregnancy indicate the opposite association (see 11 12 below). These findings indicate that the commonly used sFLT1:PIGF ratio should be 13 interpreted cautiously in the first trimester. Finally, increased attention has been paid to longitudinal changes of these factors during pregnancy, but the results are inconclusive.^{41, 44,} 14 45 15

Data exist from a number of other proteins. Low maternal first trimester levels of placental protein 13 (PP-13), another protein regulating placental vascular development, have been reported in pregnancies complicated by SGA,^{31, 46} but results are, again, inconsistent.^{47, 48} Similarly, A-disintegrin and metalloprotease 12 (ADAM12), a protease with similar function to PAPP-A, was reduced between 11 and 14 weeks in mothers who subsequently delivered small infants (BW<5th or <10th centile).^{31, 49, 50}

22 Second and third trimester

The screening efficacy of tests performed in the second and third trimester was assessed for two main reasons: (i) the availability of data collected during screening studies for the identification of aneuploidies and birth defects during the second trimester; (ii) the idea that measurements performed in the third trimester may have better predictive ability due to being temporally closer to the onset of disease.⁵¹

6 Down syndrome and anomaly screening: AFP, total hCG, uE3 and inhibin A

The second trimester quadruple screening is performed at 15-22 weeks of gestation (wkGA)
and includes measurements of alpha fetoprotein (AFP), hCG (intact and/or its β subunit),
unconjugated estriol (uE3) and inhibin A.⁵² The factors measured may also provide
information on placental permeability (AFP) and endocrine activity (hCG, uE3 and inhibin A).
Hence many studies have addressed the ability of these proteins to predict placentally-related
pregnancy complications.

Elevated maternal serum levels of AFP are associated with SGA (BW<5th centile) with or 13 without preterm delivery^{53, 54} and stillbirth due to reduced BW (<5th centile).⁵⁵ The 14 15 combination of low PAPP-A in the first trimester and high AFP in the second trimester is particularly strongly predictive of severe FGR.^{56 57} In the FASTER trial, f β -hCG alone (\geq 2.0 16 17 MoM) was not associated with any adverse outcome studied. In contrast, maternal circulating 18 AFP (\geq 2.0 MoM), Inhibin A (\geq 2.0 MoM) and uE3 (\leq 0.5 MoM) were significantly associated 19 with an increased risk of delivering an SGA infant.⁵⁸ The risk was particularly high when 20 measurements were combined. In other studies, women with an elevation of hCG alone 21 (threshold starting at >2.5 MoM) or in combination with high AFP had an increased risk of 22 SGA.^{54, 59} Low levels of unconjugated estriol (uE3) alone or in combination with AFP and/or hCG were associated with SGA (BW<5th percentile).^{54, 60} 23

1 Other placental markers: sFLT1, PIGF, ENG, hPL, uE3

2 A number of large scale studies have analyzed biomarkers in the second and third trimester 3 with the aim of identifying useful candidates for FGR screening. The SCOPE study recruited 4 >5000 low risk women and has reported that maternal levels of sFLT1 and PAPP-A were 5 decreased in pregnancies with SGA without maternal hypertension, whereas PAPP-A and PIGF 6 were decreased in pregnancies with SGA and maternal hypertension.⁶¹ Multicenter studies 7 conducted by Prof Kypros Nicolaides (King's College Hospital, London, UK) have assessed the 8 predictive associations of a number of circulating biomarkers at different stages in the second 9 and third trimesters. At 19-24wkGA, maternal PIGF was lower and AFP were higher in women 10 who subsequently delivered an SGA infant preterm. Term SGA was associated with lower maternal levels of PIGF, sFLT1 and PAPP-A.⁵³ When measured at 30-34wkGA, low PIGF and 11 12 high sFLT1 were associated with delivery of an SGA infant⁶² and the associations were 13 stronger for preterm SGA. At 35–37wkGA, PIGF and sFLT1 concentrations in the lowest and highest 5th centile, respectively, were associated with SGA.⁶³ Moreover, extremes of the ratio 14 have also been associated with the subsequent risk of intra-uterine fetal death.⁶⁴ 15

Mothers with a reduced rate of decrease of the anti-angiogenic factor sENG between the first and second trimesters had a higher risk of adverse pregnancy outcome, including SGA (BW<10th centile).⁶⁵ Moreover, as mentioned above, maternal plasma sENG concentrations remained elevated throughout the second and third trimester in patients destined to deliver an SGA neonate.⁴¹ Maternal serum levels of the hormones human placental lactogen (hPL) and uE3 were shown to be reduced in pregnancies with SGA fetuses in the second and third trimesters.^{66, 67}

23 Strength of prediction

1 A systematic review evaluated the strength of association between a wide range of 2 biomarkers and utero-placental Doppler and different causes of stillbirth. The review 3 concluded that only high resistance patterns of uterine Doppler in the second trimester and 4 low PAPP-A in the first trimester had associations which were in the clinically useful range, with positive likelihood ratios of 5 to 15.68 Moreover, these strong associations were only 5 6 observed for the sub-types of stillbirth attributed to placental dysfunction. However, the 7 review did not report the combination of ultrasound and biomarkers. The same authors also 8 evaluated 53 studies and 37 potential biomarkers for FGR screening, and suggested that none of the proposed maternal circulating factors had a high predictive accuracy.⁶⁹ Their conclusion 9 10 was that combining biomarkers with biophysical measurements and maternal characteristics 11 could be a more effective strategy. Table 2 summarizes the strongest associations identified 12 in the 2 meta-analyses.

13 Combined assessment using ultrasonic biometry and biomarkers

14 An international, prospective, multicenter observational study determined the prediction of 15 SGA achieved by the combination of ultrasonic fetal biometry and maternal serum PIGF in 16 women who were identified as clinically small for dates between 24-37wkGA. While the study demonstrated that median PIGF concentrations were lower in SGA pregnancies⁷⁰ these 17 18 authors reported that the combination of EFW and of PIGF had only modest test performance. 19 However, it should be borne in mind that many SGA infants are healthy, i.e. they are 20 constitutionally small. It would not be expected that PIGF, a test for pathology, would be 21 strongly associated with the delivery of a healthy small infant. One of the studies from the 22 King's group reported that placental biomarkers measured at 35-37 weeks performed poorly as a screening test for perinatal morbidity.⁶³ An important feature of that study was that the 23

results of ultrasound scans, performed at the same time as the biomarker, were reported and
would have influenced clinical care. Both of these studies raise important questions about the
methodological approach to future studies attempting to develop novel screening tests for
FGR, and these are discussed below.

5 We performed a prospective study of unselected nulliparous women, the Pregnancy Outcome 6 Prediction (POP) study, which combined the use of ultrasound and biochemical markers, 7 where the results of both were blinded.^{71, 72} The analysis of this study is on-going. However, 8 we have published a preliminary report that the combination of ultrasonic EFW<10th 9 percentile and an elevated sFLT1:PIGF ratio at 36wkGA was strongly predictive for late FGR 10 (BW<10th centile plus perinatal morbidity and/or preeclampsia), with a positive likelihood 11 ratio of 17.5 and a sensitivity and specificity of 38% and 98%, respectively.⁷³

12

13 **Future directions**

14 Despite years of research, screening for FGR remains clinical. Implementation of ad hoc screening using ultrasound appears to cause more harm than good.¹¹ This probably reflects 15 16 the fact that, currently, the primary intervention to manage FGR is delivery of the infant. In 17 the event of a false positive diagnosis in the preterm or early term weeks of gestational age, 18 the effect will be to cause harm through the associations between earlier delivery and 19 neonatal morbidity. The lack of progress could lead to the perception that the task being 20 attempted is futile. However, there are a number of issues about the approach to this 21 problem which have been relatively neglected. We believe that addressing some of the issues 22 outlined below may help accelerate research into clinically useful tools for screening and 23 intervention (Figure 3).

1

2 Identifying populations to screen

3 One of the most important screening parameters is the positive predictive value (PPV) of the 4 test. For the individual woman, this might, indeed, be her primary interest: how likely is it that 5 she will experience an adverse event. The PPV is determined by two factors: the prior odds of 6 disease and the positive likelihood ratio of the test. Much of the research on screening has 7 focused on the latter. However, if a woman has a very low prior risk of an outcome, even a 8 highly predictive test may result in a low absolute risk that she experiences disease. This issue 9 suggests that screening efforts might initially focus on women who are high risk and 10 nulliparous women. In the latter group the key marker of risk in pregnancy – previous 11 pregnancy outcome – is necessarily absent. Screening studies that include a high proportion 12 of parous women with previous normal pregnancies will tend to yield results with low PPVs.

13

14 High quality studies of diagnostic effectiveness

15 With the increasing awareness of the importance of Evidence-Based Medicine, it is universally 16 recognized that new interventions, such as novel drugs, must be evaluated, wherever 17 possible, by a double blind, randomized controlled trial (RCT). This is due to the biases that 18 result from patients and their caregivers being aware of allocation to the novel treatment. 19 The equivalent approach in studies of diagnostic effectiveness is to blind the results of the 20 new test. If a research study of screening reveals the test result, it ceases to be purely a 21 research study and becomes an *ad hoc* screening program. Such an approach may arise from 22 a perception that it would be unethical to conceal the result of a screening test and is based 23 on the assumption that revealing the result of the test would produce a net benefit. This is 24 true for some situations, such as diagnosis of major placenta praevia by ultrasound scan.

However, given that screening for SGA in France actually seemed to result in net harm¹¹ and 1 2 given that routine ultrasound screening has not been shown to be safe and effective, ¹⁰ it could 3 equally be argued that revealing the result of an unproven screening test and intervening on 4 the basis of the result is unethical. Similarly, there are several studies evaluating the 5 combination of ultrasound and biomarkers where the ultrasound scan result is revealed but 6 the biochemical tests are not. In these cases, the ability of the biochemical test to predict the 7 adverse outcome may be underestimated due to interventions initiated by an abnormal 8 ultrasound result. These issues underline the value of studies where all new elements of the 9 approach to screening are conducted blind, wherever possible. These examples indicate that 10 blinding of test results in studies diagnostic effectiveness is justified. Otherwise, it is very hard 11 to see how progress on screening using new diagnostic tests in pregnancy can be achieved.

12

13 Classification of SGA

14 SGA is defined on the basis of an arbitrary threshold of birth weight percentile. It follows that 15 many infants born SGA were healthy. Using unqualified SGA as an outcome may, therefore, 16 lead to weaker associations with an effective screening test than would have been obtained 17 if the analysis was focused on cases of SGA where the baby was small due to FGR. A number 18 of studies of candidate biomarkers have shown stronger associations with preterm compared 19 to term SGA. These results might indicate that FGR is more common at preterm gestational 20 ages. However, another interpretation is that the population of SGA infants delivered preterm 21 is enriched with cases of true FGR compared to SGA births at term. Similarly, the association 22 between a true test of placental dysfunction and SGA is likely to be stronger when the 23 outcome of SGA is combined with an indicator of maternal or perinatal morbidity, such as 24 preeclampsia or asphyxia, respectively. Therefore, studies which simply report the screening statistics for all SGA without reference to whether there was evidence supporting a
 pathological cause may underestimate the screening performance of an informative test.

3 Assessment of new biomarkers may also be facilitated by phenotyping SGA using prenatal 4 ultrasound. We have recently shown an association between SGA and low maternal plasma 5 levels of the non-canonical NOTCH1 ligand delta-like 1 homolog (DLK1).⁷⁴ Using genetically 6 modified mice, it was demonstrated that during pregnancy maternal circulating DLK1 is 7 mostly of fetal origin. This protein appears to provide a link between fetal demand and 8 maternal metabolic adaptation to pregnancy. Altered maternal levels of DLK1 were recently 9 measured in pregnancies complicated with preeclampsia.⁷⁵ In our cohort, low DLK1 levels 10 were associated with SGA only in presence of one or more ultrasonic indicators of FGR, and 11 levels were not different in SGA without such markers (Figure 2). Interestingly, in every case 12 of SGA with a high resistance pattern of umbilical artery Doppler, maternal serum DLK1 was 13 lower than in the matched control. This also suggests that biomarkers might help define 14 subgroups of causes of FGR. Recent studies in human placenta have correlations between 15 methylation in the DLK1 domain and birth weight.⁷⁶

16

17 Developing multi-parameter models

18 It is possible in the future that a single highly informative marker for FGR, or a subtype of FGR, 19 might be identified on the basis of mechanistic understanding of the cause of the disease. 20 However, in the meantime, it is more likely that screening tests for FGR will include multiple 21 measurements, which are derived from both imaging procedures and measurement of 22 biomarkers. Several studies adopting this approach have been described above.^{61-63, 73} 23 Development of screening models based on a multi-parameter assessment raises several 24 challenges. These include measuring and scaling the given parameter to generate consistent

1 associations in different centers, combining multiple measures into a single predictive model, 2 and accounting for interactions between parameters. Approaching this task has a number of 3 common pitfalls, such as over-fitting statistical models leading to over optimistic prediction. These issues necessitate the use of statistical methods that account for optimism⁷⁷ and 4 5 underline the value of studies which assess the development and validation of the novel test in separate groups of patients.^{78, 79} These issues are particularly important when using "omic" 6 7 methods which might yield hundreds or thousands of data points per patient. These methods 8 carry a high risk of generating over-optimistic prediction and they require a rigorous 9 methodological approach. Nevertheless, next generation sequencing (NGS) allows for 10 unbiased interrogation of genomes or transcriptomes in clinical specimens, providing 11 unprecedented opportunities to accelerate the discovery of novel biomarkers in FGR 12 screening. A successful application of the NGS techniques to the biomarkers discovery field 13 has been maternal circulating cell-free fetal DNA (cff-DNA) measurement for Down syndrome and trisomy 18 screening.^{80, 81} Elevated maternal cff-DNA levels have also been measured in 14 pregnancies complicated by FGR compared to normal pregnancies,⁸² but more recent data 15 were not always consistent with these initial findings.^{83, 84} 16

Given the key role of the placenta in the etiology of FGR, an alternative approach is to study the differences in the placental transcriptome comparing cases of FGR and controls. This may allow identification of placental pathways altered in FGR and potential biomarkers for the condition although, so far, this approach has been extensively utilized in relation to preeclampsia.⁸⁵⁻⁸⁷ Development of biomarkers would follow either by measurement of differentially expressed RNA molecules (mRNAs, miRNAs and lncRNAs) or by measurement of the proteins encoded by the differentially expressed genes. The exemplar of this approach is

1 the identification of the role of sFLT1 in preeclampsia: microarray studies identified up-2 regulation of *FLT1* as one of the key changes in the preeclamptic placenta and this was 3 paralleled by elevated levels of sFLT1 protein in the maternal circulation.³⁸ Prof Stephen 4 Tong's group has used the alternative approach of measuring mRNAs in the mother's plasma 5 and they have identified a number of placental transcripts measured in maternal blood at 26-6 30wkGA which were associated with the subsequent risk of term FGR.^{88, 89} Proteomic and 7 metabolomic technologies offer theoretical advantages, because proteins and metabolites 8 are potentially more closely linked to the phenotype under investigation than mRNA. One 9 example of a validated proteomics study of FGR identified apolipoproteins CII and CIII in 10 maternal serum of mothers with FGR compared to gestational age matched controls.⁹⁰ 11 Despite the vast opportunities of "omic" research, there are still many challenges, including 12 the difficulty related to handling large and highly complex datasets. To date, most "omic" 13 studies assessing pregnancy disorders display several limitations: small sample sizes, lack of 14 predictive ability, and the absence of validation experiments. A set of guidelines published by the Institute of Medicine⁹¹ may help standardisation of future "omic" research. 15

16

17 Identifying novel biomarkers

18 It is apparent from the summary above that many biomarkers identified for FGR were 19 originally developed as predictive tests for other conditions, such as Down syndrome and 20 preeclampsia. A PubMed search yields approximately 3 times the number of citations for 21 genetic array studies of the placenta in preeclampsia compared with FGR. Moreover, given 22 the issues of phenotyping discussed above, application of "omic" methods to the placenta in 23 cases of SGA will require detailed phenotyping of the case. Hence, a further approach to 3

4 *Meaningful clinical outcomes*

5 An extension of the issues relating to phenotyping is the identification of clinically important 6 outcomes when studying FGR. Studies which focus on SGA have a limited potential as this 7 would not be a likely primary outcome in trials of screening and intervention until disease 8 modifying therapies are available. The most serious adverse outcome of FGR is intra-uterine 9 fetal death. However, this outcome affects about 4 per 1000 pregnancies. Even the highest estimates only indicate that 50% of stillbirths might be due to FGR.⁹² Other studies suggest 10 that the proportion may be lower.⁹³ In order to be powered to study stillbirth related to FGR, 11 12 tens of thousands of women or, in the case of a very strongly predictive test, many thousands 13 of women would need to be recruited. Given the expense involved, many studies use non-14 lethal proxies of stillbirth due to FGR. While it is likely that true FGR has common features 15 whether the baby survives or dies, the use of weak proxies will tend to obscure associations. 16 One commonly used proxy is caesarean section for fetal distress, which is particularly 17 problematic when it is employed in situations where the novel test result has been revealed. 18 If the attending clinician is aware that a scan has identified a baby as suspected FGR, this could 19 lead to an association with antepartum or intrapartum caesarean section for fetal distress 20 even where the new test is not informative. There are a number of possible approaches to 21 developing meaningful clinical outcomes. One might be to combine an assessment of the 22 infant's birth weight with anthropometric measures which support a diagnosis of FGR. This 23 would allow some degree of separation between healthy SGA and FGR. However, this 24 assumes that such measures are reproducible and will correlate with clinically meaningful 1 outcomes. One approach we have used is to combine SGA birth weight with clinical 2 complications which are consistent with FGR. These include signs of fetal asphyxia or 3 compromise, such as depressed 5 minute Apgar score, the presence of metabolic acidosis in cord blood gases obtained at delivery and admission to the neonatal intensive care unit.¹⁶ The 4 5 last of these could reflect asphyxia or it could reflect other complications of FGR, such as 6 hypoglycaemia. Finally, the co-existence of SGA and maternal preeclampsia is suggestive that 7 the baby's small size is more likely to be due to placental dysfunction and this could also have 8 utility in differentiating between healthy and pathological SGA infants.

9

10 The future for RCTs of screening and intervention

11 Given the capacity for screening to cause harm, any future program of screening and 12 intervention will need to be evaluated by an RCT. We have previously discussed some of the issues around the design of such studies.^{51, 94} If the primary outcome is clinically important it 13 14 is likely that any trial will have to be very large (>10,000 women). Sample size can be reduced 15 by randomizing screen positive women to revealing the result plus intervention versus 16 concealing the result. It should also be self-evident that a trial of screening will only improve 17 outcomes if the screening test is coupled to an effective intervention. At present, the primary 18 disease modifying intervention is to deliver the baby. Given the strong associations between 19 preterm birth and perinatal morbidity and mortality, we have suggested that this provides a 20 rationale for focusing initial efforts on screening for FGR at term. However, mechanistic 21 understanding of the causes of FGR could lead to the development of novel therapeutic 22 approaches, such as re-purposing of existing drugs⁹⁵ and gene therapy.⁹⁶

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Table 1. Maternal circulating biochemical markers for predicting FGR.

BIOMARKER	MAIN FUNCTION	Changes in maternal circulating levels frequently associated with FGR		
		1 st trimester	2 nd trimester	3 rd trimester
Down's syndrome biomarkers				
pregnancy associated plasma protein A, PAPP-A	 Protease activity towards insulin-like growth factor binding proteins (IGFBPs) Decrease of insulin-like growth factors bioavailability and signaling 	ţ	Ļ	
human chorionic gonadotrophin, hCG	Maintenance of progesterone secretion from the corpus luteum	ţ	1	
alpha fetoprotein, AFP	 Protein of fetal origin with similar function to albumin in the adult Carrier-molecule for several ligands (bilirubin, steroids, and fatty acids) 		1	
unconjugated estriol, uE3	Estrogen agonist		Ļ	Ļ
inhibin A	 Negative feedback on pituitary follicle-stimulating hormone secretion, Preventing ovulation during pregnancy 		1	
Angiogenic factors	•	-	-	-
placental growth factor, PIGF	 Member of the vascular endothelial growth factor (VEGF) family Pro-angiogenic factor 	Ļ	Ļ	Ļ
soluble fms-like tyrosine kinase-1 sFLT1	 Decrease of plgf and VEGF bioavailability and signaling 	1 ,	↓ , =	1
sFLT1:PIGF ratio		1	1	1
soluble Endoglin, sENG	• Decrease of TGF- β 1 and TGF- β 3 bioavailability and signaling	1	1	1
Placental protein 13 PP-13	Promoting trophoblast invasion and spiral artery remodeling	↓ , =		
Hormonal factors				
A-disintegrin and metalloprotease 12, ADAM12	 Protease activity towards insulin-like growth factor binding proteins (IGFBPs) Decrease of insulin-like growth factors bioavailability and signaling 	Ļ		
human placental lactogen, hPL	 Induction of maternal insulin resistance and lipolysis Induction of mammary glands development and milk production 		Ļ	Ļ
delta-like 1 homolog, DLK1	Adipose tissue homeostasisMaternal metabolic adaptation to pregnancy			Ļ

BIOMARKER	Outcome	Positive LR (95% CI)	Negative LR (95% CI)	Sensitivity (%) (95% Cl)	Specificity (%) (95% Cl)	Population (n)	
Down's syndrome biomarkers	Down's syndrome biomarkers						
	Stillbirth	3.3 (1.8–6.0)	0.9 (0.8–1.0)	15 (8–26)	95 (95–96)	21158 ^{97, 98}	
РАРР-А	Placental abruption and/or SGA-related stillbirth	14.1 (9.3–21.4)	0.3 (0.1–0.8)	70 (40–89)	95 (95–96)	7919 ³⁴	
hCG	Stillbirth	2.8 (1.9–4.3)	0.4 (0.2–1.0)	70 (40–89)	75 (73–77)	2406 ⁹⁹	
AFP	Stillbirth	4.0 (3.4–4.7)	0.9 (0.9–0.9)	9 (8–11)	98 (98–98)	186802 ^{60, 100-106}	
uE3	Stillbirth	4.0 (3.0–5.3)	0.9 (0.8–0.9)	15 (11–20)	96 (96–96)	58417 ^{58, 60, 107}	
inhibin A	Stillbirth	6.1 (4.0–9.3)	0.8 (0.8–0.9)	19 (12–27)	97 (97–97)	33145 ⁵⁸	
PAPP-A + mat. characteristics	Stillbirth	3.5 (2.8-4.4)	0.7 (0.6-0.8)	35 (28-43)	90 (90-90)	33452 ¹⁰⁸	
+ US markers	SGA-related stillbirth	4.4 (3.2–5.9)	0.6 (0.5–0.8)	44 (31–57)	90 (90–90)	33365 ¹⁰⁸	
inhibin A + mat. characteristics	Stillbirth	4.1 (2.8–6.0)	0.8 (0.8–0.9)	20 (14–29)	95 (95–95)	35253 ¹⁰⁹	
Angiogenic factors							
	FGR	1.3 (1.2–1.5)	0.9 (0.8–0.9)	38 (35–42)	71 (70–72)	5709 ^{45, 110-118}	
PIGF	FGR with BW<5 th centile and abnormal UT-Doppler	2.0 (1.3–3.0)	0.5 (0.3–1.0)	65 (43–82)	67 (58–76)	124 ^{119, 120}	
	FGR with placental pathology	19.8 (7.6–51.3)	0.0 (0.0–0.3)	100 (70–100)	95 (88–98)	88117	
sFLT1	FGR with BW<5 th centile and abnormal UT-Doppler	1.9 (1.3-2.6)	0.4 (0.2-0.9)	75 (53-89)	60 (50-69)	124 ^{119, 120}	
sFLT1:PIGF ratio	FGR with BW<5 th centile and abnormal UT-Doppler	1.7 (1.2-2.4)	0.4 (0.2-1.0)	75 (53-89)	57 (47-66)	124 ^{119, 120}	
sENG	FGR with BW<5 th centile	1.8 (1.4–2.3)	0.6 (0.5–0.7)	61 (52–69)	67 (60–7	355 ⁴⁵	
sENG slope	FGR with BW<10 th centile	2.4 (1.4–4.3)	0.9 (0.8–1.0)	19 (14–27)	92 (88–95)	34665	

Table 2. Predictive accuracy of maternal circulating biomarkers for stillbirth and fetal growth restriction.

VEGF	FGR with BW≤2SD	4.4 (1.6–12.2)	0.5 (0.2–1.1)	56 (27–81)	88 (74–95)	49 ¹²¹	
Angiopoietin	FGR with BW<10 th centile and UA-Doppler	4.3 (1.9–9.4)	0.1 (0.0–0.7)	92 (67–99)	78 (58–90)	36 ¹²²	
Hormonal factors, endothelial stress markers & cytokines							
ADAM12	FGR with BW<5 th centile	2.2 (1.6–3.1)	0.9 (0.9–1.0)	12 (10–14)	95 (93–96)	1947 ^{50, 123, 124}	
IGFBP-1	FGR with BW<10 th centile	2.7 (1.1–6.5)	0.8 (0.7–1.0)	24 (12–43)	91 (85–95)	172 ¹²⁵	
PP-13	FGR with BW<5 th centile	3.6 (2.8–4.7)	0.7 (0.6–0.8)	34 (27–42)	91 (90–92)	3854 ^{46, 126, 127}	
Leptin	FGR with BW<10 th centile	2.2 (1.4–3.5)	0.5 (0.3–0.9)	63 (41–81)	72 (63–79)	139 ¹²⁸	
Fibronectin	FGR with BW<10 th centile	13.3 (5.0–35.0)	0.5 (0.2–0.8)	57 (33–79)	96 (90–98)	130129	
Homocysteine	FGR with BW<10 th centile	2.3 (1.7–3.1)	0.8 (0.8–0.9)	26 (19–33)	89 (87–90)	2088 ^{130, 131}	
sVCAM-1	FGR with BW<10 th centile	9.0 (3.9–20.7)	0.9 (0.7–1.0)	16 (7–30)	98 (97–99)	1404 ¹³²	
sICAM-1	FGR with BW<10 th centile	19.2 (11.5–32.1)	0.6 (0.5–0.8)	42 (28–58)	98 (97–99)	1404 ¹³²	
IFN-γ	FGR with BW<10 th centile	2.8 (1.4–5.6)	0.7 (0.6–0.9)	35 (24–48)	87 (78–93)	128 ¹³³	
IL-1Ra	FGR with BW<10 th centile	3.0 (1.7–5.1)	0.6 (0.4–0.8)	54 (42–67)	82 (71–89)	128 ¹³³	

Abbreviations: LR, likelihood ratio; Cl, confidence interval; SGA, small for gestational age; FGR, fetal growth restriction; BW, birthweight; UT, uterine artery; UA, umbilical artery; SD, standard deviation; PAPP-A, pregnancy associated plasma protein A; hCG, human chorionic gonadotropin; AFP, alpha fetoprotein uE3, unconjugated estriol; mat., maternal; PIGF, placenta growth factor; sFLT1, soluble fms-like tyrosine kinase-1; sENG, soluble endoglin; VEGF, vascular endothelial growth factor; ADAM12, A-disintegrin and metalloprotease 12; IGFBP, insulin-like growth factor binding protein 1; PP-13, placental protein 13; sVCAM, soluble vascular cell adhesion molecule; sICAM, soluble intercellular adhesion molecule; IFN-γ, interferon gamma; IL-1Ra, interleukin-1 receptor antagonist. Based on Conde-Agudelo et al (2013) and Conde-Agudelo et al (2015).^{68, 69}

Figure legends

Figure 1. Standard antenatal care in UK and biochemical markers measured throughout pregnancy. Biomarkers measured in clinical and research settings during pregnancy are plotted on a time scale representing the standard antenatal care for nulliparous women in the UK, which includes 10 routine midwife visits and additional visits for women delivering after 40wkGA.

Figure 2. Low maternal DLK1 levels are associated with FGR. Scatterplot of differences in maternal plasma concentrations of DLK1. DLK1 concentration was measured in maternal plasma from pregnancies with SGA (BW<10th centile) and normally grown infants (n=43 matched pairs; matching was based on maternal age, BMI, smoking status, fetal sex and mode of delivery). FGR indicators are: UT-PI in the 10th decile at 20wkGA (n=8 pairs); UA-PI in the 10th decile at 36wkGA (n=10 pairs); ACGV in the 1st decile at 20–36wkGA (n=12 pairs). BMI indicates body mass index; UT-PI indicates uterine artery pulsatility index; UA-PI indicates umbilical artery pulsatility index; ACGV indicates abdominal circumference growth velocity. Horizontal bars represent the means of the differences. Modified from Cleaton et al, Nature Genetics 2016⁷⁴.

Figure 3. Summary of key points to improve FGR screening.



Biochemical biomarkers

in UK Standard antenatal care

> Dating scan, maternal height/weight, blood pressure, test urine for proteinuria,

Figure 2



STUDIES OF DIAGNOSTIC EFFECTIVENESS

benefits:

- no changes in pregnancy management and outcome; test predictive ability can be evaluated
- higher prior odds of disease, hence higher positive \rightarrow predictive value for given positive likelihood ratio

stronger association between true tests of placental dysfunction and true FGR (as opposed to healthy SGA)

MULTI-PARAMETER **APPROACH**

key methodological consideration

- ✓ availability of assays/measurem
- ✓ combination of multiple measur into a single predictive model
- ✓ over-fitting statistical models (optimistic prediction)
- ✓ test development and validation different cohorts

feto-placental morbidity (e.g. abnormal ultrasound scan)

the results of the new test

t the population to screen:

nknown prior risk (nulliparous)

on SGA cases due to FGR:

h prior risk

erm SGA

aternal morbidity

.g. preeclampsia)

STUDIES OF CLINICAL EFFECTIVENESS (randomized controlled trials)

methodological considerations:

- arge cohort to address clinically important but rare outcomes
- andomize to (1) screen versus no-screen OR (2) randomize een positive to reveal/intervene versus conceal
- availability of a safe intervention