



Ultrasound-based gestational-age estimation in late pregnancy

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ABSTRACT

Objective Accurate gestational-age (GA) estimation, preferably by ultrasound measurement of fetal crown–rump length before 14 weeks' gestation, is an important component of high-quality antenatal care. The objective of this study was to determine how GA can best be estimated by fetal ultrasound for women who present for the first time late in pregnancy with uncertain or unknown menstrual dates.

Methods INTERGROWTH-21st was a large, prospective, multicenter, population-based project performed in eight geographically defined urban populations. One of its principal components, the Fetal Growth Longitudinal Study, aimed to develop international fetal growth standards. Each participant had their certain menstrual dates confirmed by first-trimester ultrasound examination. Fetal head circumference (HC), biparietal diameter (BPD), occipitofrontal diameter (OFD), abdominal circumference (AC) and femur length (FL) were measured every 5 weeks from 14 weeks' gestation until delivery. For each participant, a single, randomly selected ultrasound examination was used to explore all candidate biometric variables and permutations to build models to predict GA. Regression equations were ranked based

upon minimization of the mean prediction error, goodness of fit and model complexity. An automated machine learning algorithm, the Genetic Algorithm, was adapted to evaluate > 64 000 potential polynomial equations as predictors.

Results Of the 4607 eligible women, 4321 (94%) had a pregnancy without major complications and delivered a live singleton without congenital malformations. After other exclusions (missing measurements in GA window and outliers), the final sample comprised 4229 women. Two skeletal measures, HC and FL, produced the best GA prediction, given by the equation $\log_e(GA) = 0.03243 \times (\log_e(HC))^2 + 0.001644 \times FL \times \log_e(HC) + 3.813$. When FL was not available, the best equation based on HC alone was $\log_e(GA) = 0.05970 \times (\log_e(HC))^2 + 0.00000006409 \times (HC)^3 + 3.3258$. The estimated uncertainty of GA prediction (half width 95% interval) was 6–7 days at 14 weeks' gestation, 12–14 days at 26 weeks' gestation and > 14 days in the third trimester. The addition of FL to the HC model led to improved prediction intervals compared with using HC alone, but no further improvement in prediction was afforded by adding AC, BPD or OFD. Equations that included other measurements (BPD, OFD and AC) did not perform better.

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Conclusions Among women initiating antenatal care late in pregnancy, a single set of ultrasound measurements combining HC and FL in the second trimester can be used to estimate GA with reasonable accuracy. We recommend this tool for underserved populations but considerable efforts should be implemented to improve early initiation of antenatal care worldwide. © 2016 The Authors. *Ultrasound in Obstetrics & Gynecology* published by John Wiley & Sons Ltd on behalf of the International Society of Ultrasound in Obstetrics and Gynecology.

INTRODUCTION

Reliable estimation of gestational age (GA) is essential as it allows appropriate scheduling of a woman's antenatal care, informs obstetric management decisions and facilitates the correct interpretation of fetal growth assessment¹. Abnormal fetal growth patterns such as growth restriction or macrosomia may be missed or diagnosed incorrectly if GA is unknown or incorrect. Reliable GA estimation is also important at a population level to calculate rates of preterm delivery and small-for-gestational-age neonates at delivery. The lack of accurate GA estimation, particularly in geographical regions at greatest risk of these conditions, means that preterm delivery and small-for-gestational-age rates are mere approximations in many parts of the world^{2,3}.

Traditionally, GA is estimated using the first day of the last menstrual period (LMP), which assumes that ovulation occurs on day 14 of the menstrual cycle. Irregular menses, unknown or uncertain dates, oral contraceptive use or recent pregnancy or breastfeeding, issues that occur in a large proportion of women, may all influence the accuracy of this method^{4–6}. In such cases, early (< 14 weeks' gestation) ultrasound measurement of fetal crown–rump length (CRL) is recommended^{7,8}. First-trimester GA assessment is more accurate than is dating in late pregnancy because, with advancing gestation, fetal ultrasound measurements have a larger absolute error⁹ and growth disturbances become more noticeable, resulting in potential underestimation of GA for an abnormally small fetus and overestimation for a macrosomic fetus.

Unfortunately, in many settings in which high-risk pregnancies are prevalent, women attend their first antenatal care visit late in pregnancy or even at the time of delivery. This makes it difficult to manage complications, evaluate fetal growth and implement evidence-based interventions, such as the administration of corticosteroids for fetal lung maturation in cases of threatened preterm labor.

The present analysis of the Fetal Growth Longitudinal Study (FGLS), one of the main components of the INTERGROWTH-21st Project, aimed to complement our previous work of early GA estimation by ultrasound measurement of CRL¹⁰. We explored a set of equations to estimate GA using fetal biometric measurements acquired during a single ultrasound scan performed between 14 and 34 weeks' gestation.

SUBJECTS AND METHODS

INTERGROWTH-21st was a multicenter, multiethnic, population-based project, conducted between 2009 and 2014 in eight countries¹¹. Its primary aim was to study growth, health, nutrition and neurodevelopment from < 14 weeks' gestation to 2 years of age, using the same conceptual framework as that of the WHO Multicentre Growth Reference Study^{12,13}.

Eight urban areas located at low altitude (≤ 1600 m) were chosen as study sites, within which we selected all institutions that provided pregnancy and intrapartum care and at which > 80% of deliveries occurred. Women receiving antenatal care had to plan to deliver in these institutions or in a similar hospital located in the same geographical area and there had to be an absence or low levels of major, known, non-microbiological contamination such as pollution, domestic smoke, radiation or any other toxic substances¹⁴.

Women from these populations with a singleton pregnancy that was conceived naturally and who met the individual inclusion criteria were recruited prospectively and consecutively into the FGLS. The study methods have been described in detail elsewhere^{11,15}.

The true GA (GA_{true}) was defined by the woman's LMP determined at the first visit at < 14 + 0 weeks' gestation, provided that: (i) the date was certain; (ii) she had a regular 24–32-day menstrual cycle; (iii) she had not been using hormonal contraception or breastfeeding in the preceding 2 months; and (iv) it was in agreement (within 7 days) with the measurement of fetal CRL at 9 + 0 to 13 + 6 weeks' gestation¹⁵.

The same type of ultrasound machine (HD-9; Philips Ultrasound, Bothell, WA, USA) with curvilinear abdominal transducers (C5-2, C6-3, V7-3) was used for all fetal measurements at $\geq 14 + 0$ weeks' gestation. To reduce expected value bias, the ultrasound machines were specially adapted so that the measurements were not visible on the screen. However, as women presented for their first visit at different clinics within the geographical area, for those ultrasound scans performed at < 14 + 0 weeks' gestation (CRL measurements only), it was considered acceptable to use other, locally available machines, provided that they were evaluated and approved by the study team. All sonographers ($n = 39$) at the eight study sites underwent rigorous training and standardization. In accordance with the protocol, CRL and other fetal biometry measures were assessed for quality; the former were also reviewed blindly by our collaborators at the Société Française pour l'Amélioration des Pratiques Echographiques^{16,17}.

Women were invited for follow-up ultrasound scans every 5 weeks (within 1 week either side) after the initial dating scan, so that the possible ranges after the dating scan were: 14–18, 19–23, 24–28, 29–33, 34–38 and 39–42 weeks' gestation. At each visit, fetal head circumference (HC), biparietal diameter (BPD), occipitofrontal diameter (OFD), abdominal circumference (AC) and femur length (FL) were measured three times

from three separately obtained ultrasound images of each structure¹⁸.

Head measurements were taken in an axial view at the level of the thalami, with an angle of insonation as close as possible to 90°. The head had to be oval in shape, symmetrical, centrally positioned and filling at least 30% of the monitor. The midline echo (representing the falx cerebri) had to be broken anteriorly, at one-third of its length, by the cavum septi pellucidi. The thalami had to be located symmetrically on either side of the midline. Calipers were then placed on the outer border ('outer to outer') of the parietal bones at the widest or longest part of the skull for the BPD and OFD, respectively; the HC was measured using the ellipse facility of the ultrasound machine on the outer border of the skull.

The measurements of the fetal abdomen were taken in a cross-sectional view (as close as possible to a circle), with the umbilical vein in the anterior third of the abdomen (at the level of the portal sinus) and the stomach bubble visible. The operator was instructed to avoid applying too much pressure with the transducer as this can distort the circular shape of the fetal abdomen. The abdomen had to fill at least 30% of the monitor screen; the spine preferably had to be positioned at either 3 or 9 o'clock to avoid internal shadowing; and the kidneys and bladder did not have to be visible. For the measurements, the contour of the ellipse was placed on the outer border of the abdomen.

Finally, the FL was measured using a longitudinal view of the fetal thigh closest to the probe and with the femur as close as possible to the horizontal plane. The angle of insonation of the ultrasound beam was approximately 90° with the full length of the bone visualized, unobscured by shadowing from adjacent bony parts, and the femur had to fill at least 30% of the monitor screen. The intersection of the calipers was placed on the outer borders of the edges of the femoral diaphysis (outer to outer) ensuring clear femoral edges.

Detailed measurement protocols, standardization procedures and quality-control methods employed across all sites are described in detail elsewhere^{15,19–21}.

Statistical analysis

For each woman included in the study, a single ultrasound scan between 14 + 0 and 40 + 0 weeks' gestation was selected randomly using the 'sample' function in Stata (version 13). At each scan, the routinely measured fetal biometric variables were recorded. To overcome the problem of data truncation at the lower end of gestation (<14 + 0 weeks), we followed the approach described previously and applied to CRL data²². Using the international fetal growth equations for HC, AC, FL, BPD and OFD¹⁸, we simulated 20 observations for each day between 12 + 0 and 13 + 6 weeks' gestation ($n = 280$), which is approximately the same number of observations for each day of GA in the untruncated dataset. After simulation, we restricted the data based on HC by excluding values < 85 mm or > 330 mm and

visually inspecting a plot of the data to assess whether the truncation problem had been overcome. Using the augmented dataset, fractional polynomial regression analyses were employed, using the Xrfigls function in Stata, to model the mean and SD of GA for each biometric variable²².

In order to establish the relationship of fetal biometric variables and GA we used an automated machine learning 'Genetic Algorithm' (Appendix S1). This method was chosen because a more traditional fractional polynomial approach, which is well-suited to modeling a single variable, has limited scope when used with multiple biometric variables that are highly correlated. By virtue of the automated approach, the Genetic Algorithm is able to evaluate large numbers (in this case > 64 000) of potential combinations of biometric variables that are used to build polynomial equations as predictors of GA, which would not be feasible using conventional approaches. By specifying a mathematical definition of optimal performance, based upon minimization of the mean prediction error (root mean squared error (RMSE)), the first stage of model development was entirely automated with the capacity to assemble, evaluate and modify equations. We were therefore able to use the data themselves to generate preliminary models in an entirely objective manner.

Briefly, a large number of preliminary candidate equations were developed using combinations of all candidate biometric variables (including powers (0.5, 1, 2 and 3), their logs and their products). Each of the candidate equations was used to obtain for each fetus a predicted GA ($GA_{\text{predicted}}$) as an estimate of their GA_{true} . After preliminary analysis it was clear that the $GA_{\text{predicted}}$ values were not normally distributed; this was addressed by predicting the natural logarithm of $GA_{\text{predicted}}$. The equations were then ranked to assess which had the lowest uncertainty based on the 95% prediction interval.

We used a four-step approach to determine our final equation (see Appendix S1 for more detailed explanation).

- (1) Equation discovery using the Genetic Algorithm. The automated Genetic Algorithm was used to determine the equation providing the best prediction of GA using combinations of fetal biometric variables. Briefly, the model initiated itself by assigning polynomial equations linking fetal biometry within the dataset and GA_{true} . Model terms, coefficients and powers were selected randomly within specified limits. Once defined, the individual equations were each used to predict GA_{true} using the observed fetal biometry data. The performance of individual equations was measured by calculating the RMSE between the GA_{true} and $GA_{\text{predicted}}$ at each iteration of the Genetic Algorithm. For each combination of biometric variables, the equation with the lowest RMSE was selected automatically and modified by methods that mimic the genetic principles of mutation and crossover. Thus, a second generation of equations

was developed with the ‘positive predictive qualities’ of the first generation preserved in their structure. Furthermore, random variation was introduced as ‘mutations’ into the best-performing equations in order to assess whether such mutations conferred an advantage in prediction, evaluated using the RMSE. By repeating the process over many iterations, the structure of equations was continuously refined until there was convergence upon the equation, or series of equations, that most accurately predicted GA_{true} . All data processing at this stage was performed using the ‘GAPolyfitm’ function in MATLAB version R2014b.

- (2) Goodness of fit. Visual inspection of scatter plots was used to compare GA_{true} with $GA_{\text{predicted}}$ for each candidate equation obtained from the Genetic Algorithm. Quantile–quantile (QQ) plots were used to compare the distributions. Well-fitting models were identified by a QQ plot with minimal deviation from the line of equality. For each equation, absolute residuals between $GA_{\text{true}} - GA_{\text{predicted}}$ were regressed on GA using fractional polynomial methods (powers $\pm 0.5, 1, 2$ and 3) to provide an equation that approximated the SD, and multiplied by a constant to estimate the 2.5th and 97.5th centiles using the Xrigls function in Stata²³. The goodness of fit of these estimated SDs was assessed by calculating the proportion of $GA_{\text{predicted}}$ that were outside the 95% prediction intervals (± 1.96 SD), which should be 5%.
- (3) Evaluation of model complexity. To facilitate a suitable balance between parsimony and model performance, estimates of Akaike’s information criterion (AIC)²⁴ were calculated and compared. The AIC combines an estimate of the goodness of fit of a model with a penalty for increasing model complexity. In addition, candidate equations with similar indices were compared in terms of number of terms and complexity, defined as the sum of the powers of each variable. Where two equations demonstrated similar performance, the equation with a less complex structure was preferred.
- (4) Postproduction model refinement. After examining the model complexity, it appeared that most of the contribution to the prediction of GA was based on HC. Therefore, simplified models were constructed, restricted to biometry of the fetal head (HC, OFD and BPD).

The INTERGROWTH-21st Project was approved by the Oxfordshire Research Ethics Committee ‘C’ (ref: 08/H0606/139), the research ethics committees of the individual participating institutions, as well as the corresponding regional health authorities in which the project was implemented.

RESULTS

Of the 4607 women recruited into the FGLS, 4321 delivered live singletons without congenital malformations (Figure 1). Women with missing fetal

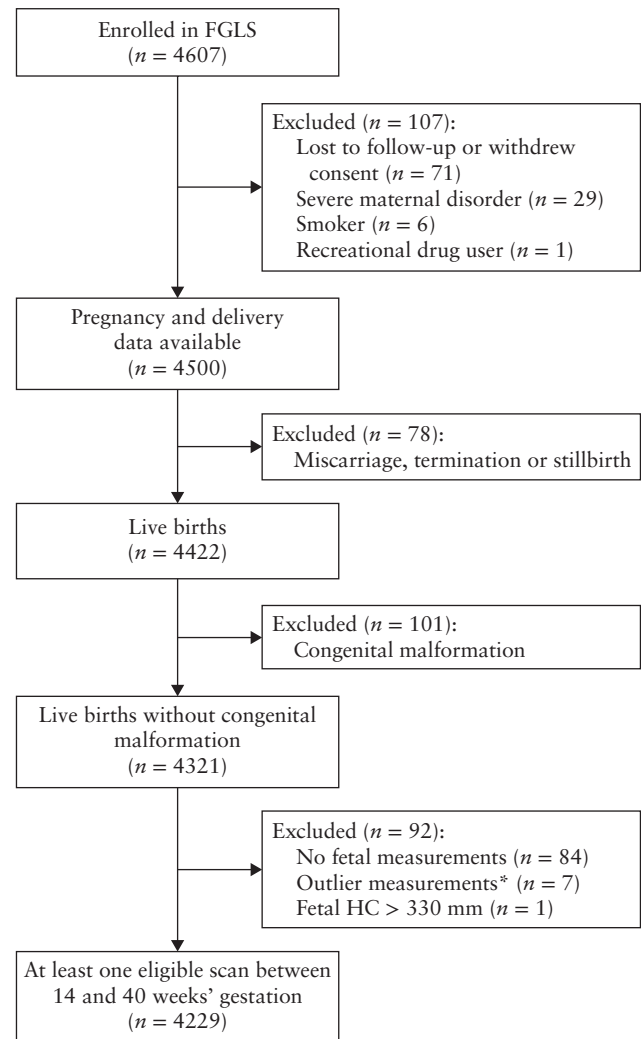


Figure 1 Flowchart of recruitment of women with singleton pregnancy to the Fetal Growth Longitudinal Study (FGLS). * > 5 SD above or below mean fetal measurement. HC, head circumference.

Table 1 Maternal and pregnancy characteristics of 4229 women enrolled in Fetal Growth Longitudinal Study of the INTERGROWTH-21st Project who had a singleton live delivery

Characteristic	Value
Maternal age (years)	27.8 \pm 3.8
Maternal height (cm)	162.2 \pm 5.8
Maternal weight (kg)	61.5 \pm 9.2
Maternal BMI (kg/m ²)	23.2 \pm 3.0
GA at first visit (weeks)	11.3 \pm 1.4
Nulliparous	2815 (66.6)
Pre-eclampsia	31 (< 1)
Preterm delivery (< 37 weeks' gestation)	189 (4.5)
Birth weight (kg)*	3.3 \pm 0.4
Birth weight < 2500 g*	127 (3.1)
Newborn sex male	2101 (49.7)

Data are given as mean \pm SD or n (%). Maternal baseline characteristics were measured at < 14 weeks' gestation. * ≥ 37 weeks' gestation only. BMI, body mass index; GA, gestational age.

Table 2 Equations for estimating gestational age (GA) and its SD in late pregnancy, derived from biometric data of 4229 singleton pregnancies

Equation	Variables in equation (mm)	Equation to estimate $\log_e GA_{\text{predicted}}$ (days)	Equation to estimate SD of $GA_{\text{predicted}}$ (days)
1	HC	$0.05970 \times (\log_e(\text{HC}))^2 + 0.000000006409 \times (\text{HC})^3 + 3.3258$	$0.6492 \times (\text{GA} \times 0.01)^3 + 2.991$
2	HC, FL	$0.03243 \times (\log_e(\text{HC}))^2 + 0.001644 \times \text{FL} \times \log_e(\text{HC}) + 3.813$	$0.04009 \times \text{GA} - 1.149$

FL, femur length (in mm); HC, head circumference (in mm); \log_e , natural logarithm.

measurements ($n=84$), and outliers defined as fetal measurements >5 SD above/below the mean ($n=7$) were excluded. We further restricted the actual data at the top end by excluding HC values >330 mm ($n=1$), resulting in a total of 4229 women who contributed a single, randomly selected ultrasound scan between 14+0 and 40+0 weeks' gestation (Figure 1). Of the 280 observations simulated and added to actual data, we similarly excluded HC values <85 mm ($n=148$) and obtained a final analysis sample of 4361 observations. The baseline characteristics and perinatal events of the study population ($n=4229$ excluding simulated observations) are shown in Table 1.

The equations that best estimated GA_{true} based on lowest RMSE, best fit and optimal AIC are shown in Table 2. The equations selected were based on HC alone and a combination of HC and FL; despite including multiple measures (HC, BPD, OFD, AC and FL) in the models, only HC and FL were retained after the selection process.

As an example, to calculate GA using Equation 1, if HC is 250 mm, median $GA = \exp[0.05970 \times (\log_e(250))^2 + 0.000000006409 \times (250)^3 + 3.3258] = \exp[5.245986506] = 189.8$ days (equivalent to 27.1 weeks). To calculate GA using Equation 2, if HC is 250 mm and FL is 55 mm, median $GA = \exp[0.03243 \times (\log_e(250))^2 + 0.001644 \times 55 \times \log_e(250) + 3.813] = \exp[5.300929] = 200.5$ days (equivalent to 28.6 weeks).

Using equations of the median and SD one can easily compute any desired centiles using the relation $P_{\text{th}} \text{ centile} = \text{median} + KSD$, where K is the normal equivalent deviate (z-score) corresponding to a particular centile, e.g. $K=1.88$ for the 97th centile and -1.88 for the 3rd centile. The SDs in this equation are the predicted estimates from the regression analysis. For example, the 3rd centile for $GA = \exp(0.05970 \times (\log_e(\text{HC}))^2 + 0.000000006409 \times (\text{HC})^3 + 3.3258) + (-1.88 \times (0.6492 \times (\text{median GA} \times 0.01)^3 + 2.991))$.

Overall, based on a model using HC only, the uncertainty of estimated GA gradually increased with advancing GA, from 6–7 days in either direction at 14+0 weeks' gestation to 15–20 days at 32+0 weeks' gestation (Table 3). Inclusion of FL led to an improvement in the accuracy of prediction throughout gestation of about 1–6 days. Inclusion of the other parameters led to no further improvement and therefore these were not included in the equations resulting from the Genetic Algorithm search.

Table 3 Characteristics of goodness of fit of an equation based on fetal head circumference (HC) (Equation 1) and an equation based on fetal HC and femur length (FL) (Equation 2) for predicting gestational age (GA) in late pregnancy

Characteristic	Equation	
	1	2
Variables in equation	HC	HC, FL
RMSE (log days)	0.0423	0.0352
R^2	0.98	0.99
Goodness of fit (%)*	5.21	6.20
AIC	16.33	14.69
Variation (days) around mean GA estimate† at:		
14 weeks	7.1	5.4
16 weeks	7.7	6.5
18 weeks	8.4	7.6
20 weeks	9.4	8.7
22 weeks	10.5	9.8
24 weeks	11.9	10.9
26 weeks	13.5	12.0
28 weeks	15.4	13.2
30 weeks	17.6	14.3
32 weeks	20.1	15.4
34 weeks	23.0	16.5

*Percentage of predicted estimates of GA outside 95% prediction interval across all GA. †Half width of 95% prediction interval. AIC, Akaike's information criterion; RMSE, root mean squared error.

The plots of $GA_{\text{predicted}}$ vs GA_{true} between 14+0 and 34+0 weeks' gestation demonstrated good model fitting (Figure 2 for the scatterplots and Figures S1 and S2 for the QQ plots).

Apart from estimating the most likely GA (by using $GA_{\text{predicted}}$ for a set of measurements), we also present the lower and upper bounds of the estimation of GA (Table S1). The lower bound can be used in clinical management for women who present in late pregnancy; it is an estimate of the likely 'least GA', e.g. they are 97.5% likely to be at least X weeks.

DISCUSSION

Main findings

We have shown that a single set of basic ultrasound measurements of HC and FL in the second trimester can be used to estimate GA with reasonable accuracy. The estimation is best at lower GAs where the 95% prediction interval is within 6 days, whereas it is just over 12 days at

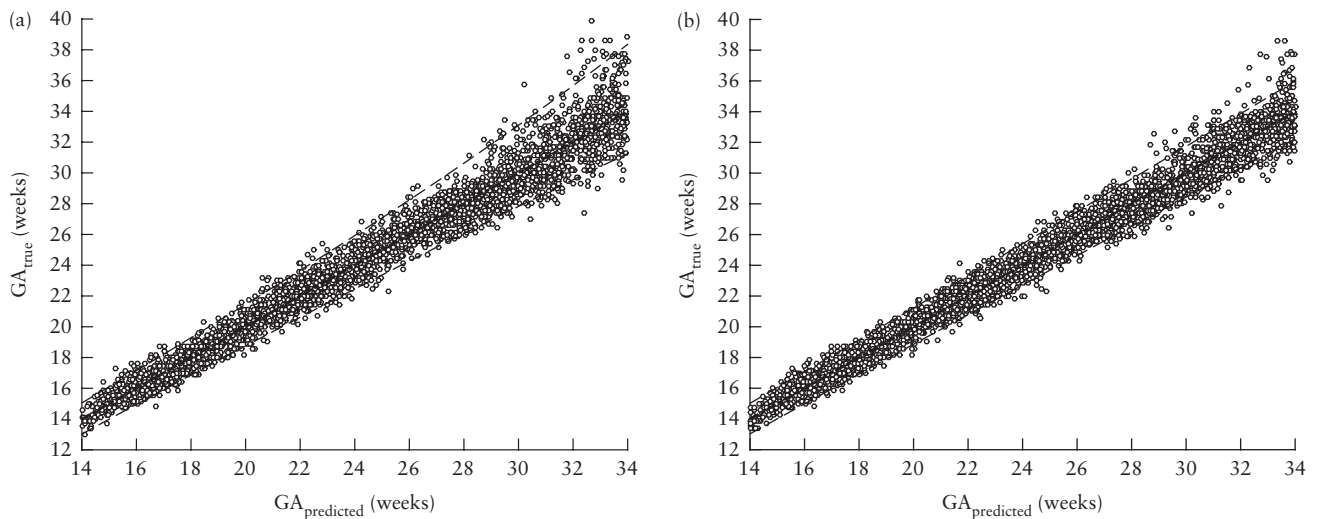


Figure 2 Scatterplots showing predicted gestational age (GA) based on fetal head circumference (a) and on fetal head circumference and femur length (b) at 14 to 34 weeks of true GA. Solid line is line of equality and dashed lines are ± 2 SD.

26 + 0 weeks' gestation. The addition of FL to HC leads to considerable improvement over just using HC, but no further improvement in prediction is afforded by adding AC, BPD or OFD.

Strengths and weaknesses

We have produced equations for GA assessment that are more precise than those used currently in routine clinical practice (Table 4). This may be due to the prospective nature of the study, a large sample size, accurately dated pregnancies, a clearly defined measurement protocol, quality control measures and a statistical approach that searched for the optimal combination of factors iteratively, rather than relying upon a user-controlled search. The multicenter, international setting of the study with measurements taken by a large group of sonographers provides external validity.

There is an intrinsic limitation when estimating GA by fetal anthropometric-based equations, i.e. that the measurement is of fetal size not GA. Fetal size may vary for reasons other than differences in GA, especially as factors conditioning abnormal fetal growth are more prevalent in the populations among which the equation is most likely to be used. In other words, it is important to take into account the impact of pathology (fetal growth restriction and overgrowth) on GA estimation. This is true for any equation estimating GA; the accuracy at an individual level will depend on the 'normality' of the fetal size and, at the population level, on the prevalence of abnormal growth patterns. Thus, efforts should focus on modifying health systems and referral pathways to prevent late presentation in pregnancy, rather than simply achieving technological advances in fetal size-based dating.

Interpretation

Ultrasound assessment of GA is performed assuming that fetal size can be used as a proxy for GA. This assumption

depends on: (i) the GA at which biometry is performed (at earlier GAs growth is more uniform and there is less measurable growth impairment); (ii) the choice of biometric variable; (iii) accuracy of the measurement, which is affected by technical aspects of imaging and operator skill; and (iv) absence of pathology that could affect growth.

The most accurate way to estimate GA is by measuring fetal CRL between 8 + 0 and 14 + 0 weeks' gestation, which is associated with a 95% prediction interval of 2.7 days^{10,25}. This method is the basis of recommended pregnancy-dating policies throughout much of the developed world²⁶. Beyond 14 + 0 weeks' gestation, fetal flexion limits the accuracy of CRL measurements for dating purposes and GA estimates are based on measurement of the HC, BPD, AC and FL or a combination of these^{8,27}.

Our results demonstrate the relative inaccuracy of late GA assessment, which is due to the increasing biological variability in fetal size as well as the increasing absolute error of fetal measurements with advancing GA⁹. Therefore, all information (clinical and imaging) should be considered when dating pregnancies and providing obstetric care, particularly after late pregnancy dating. Thus, we recommend that the following principles should be applied in clinical practice.

- (1) Assessment of fetal age should be based on the earliest available ultrasound measurement after 8 + 0 weeks' gestation, provided that the measurements are technically adequate. CRL should be used before 14 + 0 weeks' gestation and Equation 2 (HC and FL) after 14 + 0 but before 26 + 0 weeks' gestation. Equation 1 can be used in settings in which only HC is available.
- (2) If menstrual dates are reliable and within the prediction limits of the fetal measurement, ultrasound examination should merely confirm the GA assessed by LMP.

Table 4 Commonly used pregnancy-dating equations and their imprecision in estimating gestational age (GA) (half width of 95% prediction intervals)

Reference	R ²	Imprecision in GA estimation (days) for assessment at:			
		12–18 weeks	18–24 weeks	24–30 weeks	30–36 weeks
Hadlock <i>et al.</i> (1984) ²⁹					
BPD	0.967	8.3	12.1	15.3	21.6
HC	0.973	8.3	10.4	14.4	20.9
AC	0.969	11.6	14.4	15.3	20.7
FL	0.971	9.7	12.6	14.6	20.7
HC + BPD	0.974	7.6	10.4	13.9	20.0
HC + FL	0.976	8.4	10.6	13.9	18.8
HC + AC	0.98	7.6	9.4	13.0	17.6
HC + FL + BPD	0.981	7.3	9.5	12.7	17.6
HC + AC + FL	0.981	8.0	10.2	13.2	17.6
HC + BPD + AC + FL	0.981	7.6	9.8	12.6	17.1
Altman and Chitty (1997) ³⁰					
HC	NR	8.0	13.0	17.0	22.0

AC, abdominal circumference; BPD, biparietal diameter; FL, femur length; HC, head circumference; NR, not reported.

- (3) When menstrual dates are reliable and fall outside the prediction interval of ultrasound assessment there are two interpretations: the menstrual dates are in fact incorrect and GA should be based on ultrasound measurement, or the GA is correct as assessed by LMP and the fetus is an abnormal size for that GA (or both). Clinical features of growth restriction, e.g. reduced amniotic fluid or abnormal uterine or umbilical artery blood flow, should be taken into consideration, as should factors that may lead to overgrowth, e.g. maternal diabetes. An interval ultrasound scan should then be carried out to confirm GA.
- (4) When menstrual dates are unknown, GA estimation should be based on ultrasound examination, which has reasonable accuracy until 26 + 0 weeks' gestation, and a further ultrasound scan should be carried out.

Finally, when GA is estimated in the third trimester, the possible error is large and must be taken into account to ensure safe obstetric practice. The use of the concept of a 'minimum' GA, by using the lower limit of the prediction interval from the equation can be useful in this instance (Table S1). For example, if a woman presents with threatened preterm labor and the ultrasound examination suggests a median GA of 34 + 6 weeks based on HC and FL, it should be appreciated that the GA could be as low as 32 + 6 weeks, i.e. it is most likely that the GA is 34 + 6 weeks, but we are 95% certain that the GA is at least 32 + 6 weeks. However, if the fetus is growth restricted, the GA could be as much as 36 + 5 weeks. This analysis is very relevant to clinical decision making, e.g. when administering prophylactic corticosteroids or transferring a neonate to a higher level of care. In contrast, labor induction may be considered at 40 + 0 weeks' gestation based on late assessment, as the GA could be more advanced. Such a clinically cautious approach is particularly important as it is known that unreliable reporting of LMP and late

antenatal care are both associated with adverse pregnancy outcome^{5,28}.

Conclusion

We have shown that a single set of ultrasound measurements in the second trimester can be used to estimate GA with relative accuracy. We recommend these tools for the management of women who present late in pregnancy. However, we strongly encourage, as a priority, the promotion of early antenatal care in regions and subpopulations that are not yet benefiting from this practice.

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SUPPORTING INFORMATION ON THE INTERNET



Appendices S1 and S2, Figures S1 and S2 and Table S1 may be found in the online version of this article.