Early prediction of live birth for assisted reproductive technology patients: a convenient and practical prediction model

Hong Gao1,2, Dong-e Liu3, Yumei Li3, Xinrui Wu2 & Hongzhuan Tan2*

Live birth is the most important concern for assisted reproductive technology (ART) patients. Therefore, in the medical reproductive centre, obstetricians often need to answer the following question: “What are the chances that I will have a healthy baby after ART treatment?” To date, our obstetricians have no reference on which to base the answer to this question. Our research aimed to solve this problem by establishing prediction models of live birth for ART patients. Between January 1, 2010, and May 1, 2017, we conducted a retrospective cohort study of women undergoing ART treatment at the Reproductive Medicine Centre, Xiangya Hospital of Central South University, Hunan, China. The birth of at least one live-born baby per initiated cycle or embryo transfer procedure was defined as a live birth, and all other pregnancy outcomes were classified as no live birth. A live birth prediction model was established by stepwise multivariate logistic regression. All eligible subjects were randomly allocated to two groups: group 1 (80% of subjects) for the establishment of the prediction models and group 2 (20% of subjects) for the validation of the established prediction models. The sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV) of each prediction model at different cut-off values were calculated. The prediction model of live birth included nine variables. The area under the ROC curve was 0.743 in the validation group. The sensitivity, specificity, PPV, and NPV of the established model ranged from 97.9–24.8%, 7.2–96.3%, 44.8–83.8% and 81.7–62.5%, respectively, at different cut-off values. A stable, reliable, convenient, and satisfactory prediction model for live birth by ART patients was established and validated, and this model could be a useful tool for obstetricians to predict the live rate of ART patients. Meanwhile, it is also a reference for obstetricians to create good conditions for infertility patients in preparation for pregnancy.

Live birth is the most important concern for assisted reproductive technology (ART) patients; additionally, it is the only criterion used to determine whether ART treatment is successful. In recent years, ART in humans has developed very quickly, but the live birth rate is still less than optimal1. Fundamentally, numerous key factors play important roles in live birth outcomes, including embryo parameters2,3, reproductive hormone levels4–6, and the patient’s age7,8, among others; however, the degrees to which these factors affect birth outcomes are not clear. Consequently, it is very significant to investigate how these factors affect live birth outcomes, determine their degrees of influence and then establish live birth prediction models.

Previous studies have used various strategies to predict live birth for different sub-groups of ART patients, including in vitro fertilization (IVF) patients, intracytoplasmic sperm injection (ICSI) patients, ICSI patients with uncompromised ovarian reserve, and poor ovarian responders. Different strategies have included risk scoring systems, deep phenotyping, granulosa cell biomarkers, and clinical characteristics (female age, testicular sperm extraction cycle, male and female reproductive hormones, spermatozoa parameters, infertility diagnosis, and oocyte parameters). The sensitivity and specificity of these live birth prediction models exhibit large variations. Prajna Banerjee et al. used 52 clinical characteristics to establish a model. The area under the receiver operating characteristic (ROC) curve was up to 0.80 for IVF patients in their first cycles, but in their

1School of Nursing, University of South China, Hengyang 421001, Hunan, China. 2Department of Epidemiology and Health Statistics, XiangYa School of Public Health, Central South University, Changsha 410008, Hunan, China. 3Reproductive Medicine Centre, Xiangya Hospital of Central South University, Changsha 410008, Hunan, China. *email: tanhz99@qq.com
subsequent treatments, this value decreased to less than 0.68. Although many live birth prediction models have been established, they are rarely used in clinical practice. The main reasons may include the following: (1) they cannot be applied to all ART patients because the model is based on only 1–2 types of ART patients; (2) some predictors need more complicated and expensive laboratory tests; (3) the use of these models is not sufficiently convenient, and (4) some models are less accurate than others for predicting live birth.

We aimed to establish a convenient and practical live birth prediction model that has higher predictive value and that can be applied to all ART patients.

**Results**

There were 15,717 ART treatments performed from 2012 to 2017. Of these, 1891 subjects who had missing information on their live birth outcomes were excluded, leaving 13,826 subjects for analysis. Among them, 80% of the subjects (11,071) were allocated to group 1 (establishment model), and 20% of them (2755) were allocated to group 2 (validation model).

**Univariate analysis results.** We analysed the relationships between the variables and live birth outcomes by univariate analysis. Twenty-two variables were found to be significantly associated with live birth outcomes ($p$-value < 0.05, Tables 1, 2).

**Logistic regression analysis and prediction model establishment.** Based on our univariate analysis results, we found that maternal age, body mass index, number of previous ART treatments, female infertility duration, number of previous pregnancies, number of abortions, basal FSH, sperm concentration, endometrial thickness before embryo transfer, number of antral follicles, total number of oocytes, sperm viability, sperm progressive motility, type of embryo transfer, quality of transferred embryos, maternal education, infertility diagnosis, uterine volume, artificial insemination technology used, stimulation protocol, total number of transferred embryos, and total dose of gonadotropin were significantly associated with live birth. We used these variables as independent variables to perform multiple logistic regression analysis. Each step of this model is shown in Supplementary Table S1. The area under the ROC curve was 0.722 (95% CI, 0.709–0.735). The model is as follows: 

$$\text{Logit} \ P = -1.857 + 0.199X_1 + 0.150X_2 + 0.276X_3 + 0.077X_5 - 0.149X_8 + 1.205X_9 + 0.690X_{12} + 0.770X_{13} + 0.534X_{19}.$$ 

**Verification prediction model.** We performed a validation of the prediction model of live birth by using the model in the validation group [with 20% of the subjects (2755)]. The area under the ROC curve was 0.743 (95% CI, 0.719–0.768) (Fig. 1). Table 5 shows the practical predictive value for live birth with different cut-off values. The sensitivity (SE) and specificity (SP) ranged from 97.9–24.8% and 7.2–96.3%, respectively, at different cut-off values.

Table 1. Comparison of the characteristics between the live birth and no live birth groups comprising 13,826 ART patients from 2010–2017 (continuous variables). (Mean ± SD). *FSH, follicle-stimulating hormone.*
**Discussion**

We established a prediction model of live birth by multivariable logistic regression analysis, and this model included 9 common variables. In the establishment model group, the ROC value was 0.722, and there was good calibration. In the validation model group set, the ROC value was 0.743, and the model was calibrated well. Our model has several obvious advantages. Our model is a convenient and practical prediction model because information on all the variables included in the model is generally available in the clinic, and there is no need for any special test. Some predictors in our model, such as endometrial thickness, stimulation protocol, and embryo parameters, can be the focus of interventions. Therefore, the model has a certain predictive value and instructional clinical value in early treatments. The model can predict a live birth in the early pregnancy stage because information on almost all the variables included in the model is available at the beginning of pregnancy. Moreover, our model has acceptable clinical predictive value, and the area under the ROC curve reached 0.743 in the validation group, which is higher than the values of most of the previously reported models that were similar to our model\(^1,10,13,16\). Although the ROC values of some models are larger than ours, the variables in these models, such as gene or granulosa cell biomarkers, may be inconvenient to assess\(^1,17\). Moreover, a variable, such as gene expression, may be unchangeable and have no preventive value\(^1\). Furthermore, information on variables, such as HCG and progesterone may only be attainable after pregnancy is achieved and cannot be used for the prediction of a live birth\(^9,18,19\). Many previous prediction models of live birth are not applicable to all ART patients but are instead only applicable for a specific infertility subgroup\(^9,20–22\); however, our model is suitable for different artificial insemination technologies and all ART patients.

We have established a highly discriminatory, well-calibrated, robust, and practical prediction model that can use available clinical data to predict the live birth rate and may be transferred to corresponding computer

| Variables                        | N   | Live birth (n) | Live birth rate (%) | p-value |
|----------------------------------|-----|----------------|---------------------|---------|
| **Pre-ART factors**             |     |                |                     |         |
| Maternal education              |     |                |                     | <0.001  |
| Under 6 years                   | 1086| 392            | 36.1                |         |
| 6–9 years                       | 6186| 2723           | 44.0                |         |
| 10–12 years                     | 1737| 834            | 48.0                |         |
| 13 years and over               | 4341| 2047           | 47.2                |         |
| Infertility diagnosis           |     |                |                     | <0.001  |
| Male factors                    | 1890| 818            | 43.3                |         |
| Ovulation dysfunction           | 58  | 25             | 43.1                |         |
| Decreased ovarian reserve       | 150 | 36             | 24.0                |         |
| Tubal factors                   | 9568| 4164           | 43.5                |         |
| Uterine factors                 | 142 | 54             | 38.0                |         |
| Chromosome abnormality          | 14  | 5              | 35.7                |         |
| Unexplained factors             | 44  | 15             | 34.1                |         |
| Male + female factors           | 1700| 774            | 45.5                |         |
| **Protocol and treatment factors** |   |                |                     |         |
| Stimulation protocol            |     |                |                     | <0.001  |
| Long protocol                   | 5648| 3024           | 53.5                |         |
| Short protocol                  | 1752| 521            | 29.7                |         |
| Other protocol                  | 2703| 752            | 27.8                |         |
| Artificial insemination technology |   |                |                     | 0.02    |
| IVF                              | 9820| 4344           | 44.2                |         |
| ICSI                             | 2871| 1215           | 42.3                |         |
| IVF + ICSI                      | 884 | 358            | 40.5                |         |
| IUIa                             | 77  | 27             | 35.1                |         |
| Type of embryo transfer         |     |                |                     | <0.001  |
| Fresh embryo                    | 7431| 3544           | 47.7                |         |
| Frozen-thawed embryo            | 6309| 2442           | 38.7                |         |
| Quality of the transferred embryosb |   |                |                     | <0.001  |
| I                               | 11,585| 5482       | 47.3                |         |
| II                              | 1112 | 334           | 30.0                |         |
| III                             | 443  | 62            | 14.0                |         |

Table 2. Comparison of the live birth rates of the different sub-groups of the 13,826 ART patients from 2010–2017 (categorical variables). \(^1\)IUI, intrauterine insemination. \(^b\)Quality of the transferred embryos: I is the best-quality embryo, followed by II and III.
software for easy operation. Clinicians and public health workers can easily use this model to identify high-risk populations for the management by ART.

As we all know, breeding a new life is a very complex process, which will be affected by many known and unknown factors. Especially, for infertility patients, the condition will be more complex and changeable. Medical technology level of different hospitals and doctors quite naturally plays a fundamental role. To date, it is hard to predict live birth rates before embryo transfer. In the early stage of infertility treatment, patients and doctors are most concerned about "How many normal oocytes are there?" , "How many normal sperm are there?" and "How many embryos can be transferred?" Therefore, the prediction of live birth rate is at least based on the successful embryo transfer. However, this finding can be used as a guidance to try to create good conditions for infertility patients in preparation for pregnancy.

Table 3. Variable assignment in the multivariate logistic regression analysis. *X2, X4, X7, X9, X10, X13, X22 were entered into the models as dummy variables, and the group with the highest live birth rate was selected as the reference group, which was assigned "0".

| Variables | Variable names | Assignment statement |
|-----------|----------------|----------------------|
| Maternal age (year) | X1 | 1 = < 25, 2 = 25~, 3 = 30~, 4 = ≥ 35 |
| Maternal education (year)* | X2 | 1 = <10, 0 = ≥ 10 |
| Number of previous ART treatments | X3 | 1 = 1, 2 = 2, 3 = ≥ 3 |
| Uterine volume (mL)* | X4 | 1 = <30, 0 = 30~, 2 = 50~, 3 = ≥ 70 |
| No. of abortions | X5 | 1 = 0, 2 = 1, 3 = 2, 4 = ≥ 3 |
| Sperm concentration (million/mL) | X6 | 1 = < 43, 2 = 43~ , 3 = 85~ , 4 = ≥ 144 |
| Infertility diagnosis* | X7 | 1 = male factors, 2 = ovulation dysfunction, 3 = decreased ovarian reserve, 4 = tubal factors, 5 = uterine factors, 6 = chromosome abnormality, 7 = unexplained, 0 = male + female factors |
| Endometrial thickness before embryo transfer (mm) | X8 | 1 = < 8, 2 = 8~ , 3 = 10~ , 4 = ≥ 12 |
| Total no. of transferred embryos* | X9 | 1 = 1, 0 = ≥ 2 |
| Total dose of gonadotropin (IU)* | X10 | 1 = 1500, 0 = 1500 ~ , 2 = 2000 ~ , 3 = ≥ 2500 |
| Total no. of oocytes | X11 | 1 = 8, 2 = 8~ , 3 = ≥ 13 |
| Quality of the transferred embryos | X12 | 1 = 1, 2 = II, 3 = III |
| Stimulation protocol* | X13 | 0 = long protocol, 1 = short protocol, 1 = other protocol |
| Sperm viability (%) | X14 | 0 = <40, 1 = 40~ , 2 = 50~ , 3 = ≥ 70 |
| Sperm progressive motility (%) | X15 | 0 = <32, 1 = 32~ , 2 = 50~ , 3 = ≥ 70 |
| BMI (kg/m²) | X16 | 1 = <18, 2 = 18~ , 3 = 24~ , 4 = ≥ 28 |
| Female infertility duration (yr) | X17 | 1 = < 3, 2 = 3~, 3 = 5~, 4 = ≥ 7 |
| No. of previous pregnancies | X18 | 1 = 0, 2 = 1, 3 = 2, 4 = 3, 5 = ≥ 4 |
| Basal FSH (mIU/mL) | X19 | 0 = <10, 1 = ≥ 10 |
| No. of antral follicles | X20 | 0 = <5, 1 = ≥ 5 |
| Type of embryo transfer | X21 | 0 = fresh embryo, 1 = frozen-thawed embryo |
| Artificial insemination technology* | X22 | 0 = IVF, 1 = ICSI, 2 = ICSI + IVF, 3 = IUI |
| Live birth outcome | Y | 1 = live birth, 0 = no live birth |

Table 4. The association between live birth and the predictive variables included in the logistic predicting model.

| Variables | B3 | B1 | p | Collinearity Statistics |
|-----------|----|----|---|-------------------------|
| Maternal age (X1) | 0.199 | 0.035 | < 0.001 | 0.879 | 1.138 |
| Maternal education (X2) | 0.130 | 0.058 | 0.009 | 0.996 | 1.004 |
| Number of previous ART treatments (X3) | 0.276 | 0.047 | < 0.001 | 0.679 | 1.472 |
| No. of abortions (X5) | 0.077 | 0.034 | 0.023 | 0.905 | 1.105 |
| Endometrial thickness before embryo transfer (X8) | −0.149 | 0.033 | < 0.001 | 0.881 | 1.135 |
| Total no. of transferred embryos (X9) | 1.205 | 0.119 | < 0.001 | 0.949 | 1.053 |
| Quality of the transferred embryos (X12) | 0.690 | 0.071 | < 0.001 | 0.983 | 1.018 |
| Stimulation protocol (X13) | 0.770 | 0.071 | < 0.001 | 0.654 | 1.530 |
| Basal FSH (X19) | 0.534 | 0.123 | < 0.001 | 0.954 | 1.048 |
| Constant | −1.857 | 0.175 | < 0.001 | | |
Our live birth prediction models were further validated with a separate sample, allowing us the ability to evaluate the true predictive performance of the models when they are being used in other populations. We also examined the impacts of different cut-off values on sensitivity, specificity, PPV and NPV, to establish an appropriate reference range. Clinicians and public health workers could conveniently select different cut-off values in their live birth assessment process to obtain optional results.

There are several limitations to this study. Our data were obtained only from a large reproductive medicine centre, and the ROC values of our model are not the largest among the reported models. Therefore, we do not think the model is unchangeable. With the development of medical technology, new variables will be added to our model, and our prediction model will be continuously optimized. In addition, the applicability of the model in other clinics needs to be further verified, which will be our next research work.

In conclusion, a prediction model for live birth by ART patients was established and validated. The model is stable, reliable, convenient, and satisfactory; furthermore, this model could be a useful tool for early-gestation predictions by obstetricians of whether or not a ART patient has a high probability of a live birth and for trying to create good conditions for ART patients in preparation for pregnancy.

**Table 5.** The validity of the prediction model with different cut-off values in different groups (%). *PPV: positive predictive value. *NPV: negative predictive value.

| Cut-off value | Group 1 | | | | Group 2 | | | |
|---------------|---------|---|---|---|---------|---|---|---|
| 0.25          | 99.7    | 1.7 | 43.9 | 88.0 | 97.9    | 7.2 | 44.8 | 81.7 |
| 0.30          | 95.5    | 14.3 | 46.2 | 80.5 | 93.8    | 20.2 | 47.5 | 80.9 |
| 0.35          | 86.7    | 35.1 | 50.7 | 77.4 | 89.0    | 34.7 | 51.2 | 80.4 |
| 0.40          | 78.7    | 0.50 | 54.8 | 75.3 | 84.2    | 43.9 | 53.6 | 78.3 |
| 0.45          | 72.4    | 58.9 | 57.6 | 73.5 | 77.6    | 57.3 | 58.3 | 76.9 |
| 0.50          | 67.8    | 66.3 | 60.8 | 72.8 | 68.3    | 65.9 | 60.7 | 73.0 |
| 0.55          | 62.2    | 72.5 | 63.5 | 71.4 | 65.1    | 71.3 | 63.6 | 72.6 |
| 0.60          | 55.3    | 78.9 | 66.9 | 69.6 | 57.5    | 78.7 | 67.5 | 70.6 |
| 0.65          | 44.5    | 84.9 | 69.4 | 66.5 | 50.3    | 84.7 | 71.7 | 68.9 |
| 0.70          | 34.4    | 90.8 | 74.2 | 64.3 | 43.1    | 89.4 | 75.8 | 67.1 |
| 0.75          | 25.4    | 94.3 | 77.4 | 62.1 | 32.4    | 93.4 | 79.1 | 64.2 |
| 0.80          | 18.3    | 96.5 | 80.1 | 60.5 | 24.8    | 96.3 | 83.8 | 62.5 |
Methods

Study data. Between January 1 2010 and May 1 2017, we conducted a retrospective cohort study of women undergoing ART treatments at the Reproductive Medicine Centre, Xiangya Hospital of Central South University, Hunan, China. All data of the subjects were retrieved from the electronic medical records (Haitai, Nanjing, China) of Xiangya Hospital Central South University. The inclusion criteria for data analysis were as follows: (1) completion of the basic medical examination; (2) completion of the entire ART treatment cycle; (3) complete record of the couple’s basic information; and (4) thorough documentation of the live birth outcome.

Outcome measures. In our study, we focused on live birth. The birth of at least one live-born baby per initiated cycle or embryo transfer procedure was defined as a live birth, and all the other adverse pregnancy outcomes were classified as no live birth.

Statistical analysis. A live birth prediction model was established by stepwise multivariate logistic regression (α = 0.05 for entry, and α = 0.10 for removal). When establishing the model, the criteria for selecting predictive variables were as follows: first, p value was less than 0.05; second, it contributed to improving the area under the ROC curve. All eligible subjects were randomly allocated to two groups: group 1 (80% of subjects) for the establishment of the prediction models and group 2 (20% of subjects) for the validation of the established prediction models. Subjects of the two groups were independent without repetition. The areas under the ROC curve generated by the logistic regression model were applied to evaluate the effectiveness of the prediction models. We further calculated the sensitivity, specificity, positive predictive value, and negative predictive value of the prediction models with different cut-off values.

All data were managed and analysed using the statistical package for social sciences (SPSS) software version 17.0 (Chicago, IL, SPSS Inc. 2008) and Excel (Microsoft Corp., Redmond, WA, USA). Measurement data are described as the mean ± standard deviation (SD), and enumeration data are described as numbers (percentages). All p values corresponded to two-sided tests, and p < 0.05 was considered statistically significant.

Ethical approval and consent to participate. The study was approved by the Ethics Committee of Xiangya Hospital of Central South University, mainly including the use of data from various clinical examination and laboratory tests of patients. All infertility patients presenting to the Reproductive Medicine Centre, Xiangya Hospital of Central South University, Hunan, China, who were planned for ART treatments and who signed the informed consent were enrolled in the study from January 1 2010 and May 1 2017. In addition, we confirmed that all methods were performed in accordance with the assisted reproductive technology relevant guidelines and regulations.

Data availability

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

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Author contributions

H.G., D.E.L. and H.Z.T. formulated the study questions and directed their implementation. H.G., D.E.L. and H.Z.T. contributed to the study design. H.G. performed the statistical analysis. X.R.W. and H.Z.T. provided guidance on the statistical analysis. D.E.L. and Y.M.L. provided clinical data. H.G. and H.Z.T. drafted the article. All authors were involved in revising the article and have approved this final version.

Competing interests

The authors declare no competing interests.

Additional information

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Correspondence and requests for materials should be addressed to H.T.

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