First-trimester Combined Screening Test for Aneuploidies in Brazilian Unselected Pregnancies: Diagnostic Performance of Fetal Medicine Foundation Algorithm

Rastreio combinado do primeiro trimestre para aneuploidias em gestantes brasileiras não selecionadas: desempenho diagnóstico do algoritmo da Fetal Medicine Foundation

Laila Pinheiro Abi Abib1 Renato Augusto Moreira de Sá1 Fernando Maia Peixoto-Filho1

1 Department of Woman and Child Health, Universidade Federal Fluminense, Niterói, RJ, Brazil

Address for correspondence Renato Augusto Moreira de Sá, PhD, Departamento Materno-Infantil, Universidade Federal Fluminense, Avenida Marques de Paraná, 303 - 2° andar. Centro - Niterói 24033-900, Brazil (e-mail: rsa@cpdt.com.br).

Abstract

Objective The main objective of this study was to examine the diagnostic performance of the first-trimester combined test for aneuploidies in unselected pregnancies from Rio de Janeiro and compare it with the examples available in the literature.

Methods We investigated 3,639 patients submitted to aneuploidy screening from February 2009 to September 2015. The examination is composed of the Fetal Medicine Foundation risk evaluation based on nuchal translucency evaluation, mother’s age, presence of risk factors, presence of the nasal bone and Doppler of the ductus venous in addition to biochemical analysis of pregnancy-associated plasma protein A (PAPP-A) and beta-human chorionic gonadotropin (β-hCG) markers. The cut-off point for high risk for aneuploidies was defined as greater than 1:100, with intermediate risk defined between 1:100 and 1:1,000, and low risk defined as less than 1:1,000. The variable aneuploidy was considered as a result not only of trisomy of chromosome 21 but also trisomy of chromosomes 13 and 18.

Results Excluding the losses, the results of 2,748 patients were analyzed. The first-trimester combined test achieved 71.4% sensitivity with a 7.4% false-positive (FP) rate, specificity of 92.6%, positive predictive value (PPV) of 6.91% and negative predictive value (NPV) of 99.76%, when the cut-off point considered was greater than 1:1,000. Through a receiving operating characteristics (ROC) curve, the cut-off point that maximized the sensitivity and specificity for the diagnosis of aneuploidies was defined as 1:1,860. When we adjusted the false-positive (FP) rate to 5%, the detection rate for this analysis is 72.7%, with a cut-off point of 1:610.

Conclusion The combined test of aneuploidy screening showed a detection rate inferior to those described in the literature for a higher FP rate.
Introduction

Chromosomal diseases are the leading cause of perinatal mortality and developmental abnormalities. In 1866, Langdon Down described for the first time the syndrome that today bears his name, in reference to individuals affected by the trisomy of chromosome 21, the most common chromosomal aneuploidy in humans (0.12% or 1 in 800 births). The diagnosis of aneuploidies depends on invasive procedures that are associated with risks of gestational loss. The total fetal loss for chorionic villus sampling and amniocentesis ranges from 1.5 to 2.0%. In an attempt to indicate these additional ultrasound markers of the presence of nasal bone and Doppler of the venous duct, the cut-off point for high-risk patients will be 1:100. Patients with risk between 1:100 and 1:1,000 are considered to have an intermediate risk.

First-trimester screening allows early diagnosis of aneuploidy. There are many strategies that are available for chromosomal abnormality screening. The first-trimester combined test was introduced by Wald and Hackshaw (1997) and is one of the most popular and useful strategies. In this screening strategy, the risk is calculated based on the sonographic findings and maternal serum levels of free beta-human chorionic gonadotrophin (β-hCG) and pregnancy-associated plasma protein A (PAPP-A). The screening performance of the combined test has been reported to range from 82 to 95% detection rate, with a 5 to 7% false positive (FP) rate. However, it may vary between different ethnicities as well as by age group.

According to the Fetal Medicine Foundation (FMF) proposed screening, patients with a risk lower than 1:1,000 are classified as low risk and are reassured. If we include the additional ultrasound marker of the presence of nasal bone and Doppler of the venous duct, the cut-off point for high-risk patients will be 1:100. Patients with risk between 1:100 and 1:1,000 are considered to have an intermediate risk.

In Brazil, the first-trimester combined test has been widely used but little information is available on the performance of this screening method in the Brazilian population. The aim of this study was to examine the performance of the first-trimester combined test in unselected pregnancies from Rio de Janeiro by analyzing the detection rates (sensitivity), specificity, positive and negative predictive values, percentage of FPs and relative risks for the occurrence of aneuploidy, as well as to estimate the sensitivity and specificity for various risk cut-off points to construct a receiving operator characteristics (ROC) curve.

Methods

The current study was based on a cohort of singleton gestation women who booked a combined test for first-trimester risk assessment at our center from February 2009 to September 2015. Women with pregnancies resulting from ovum donation,
multiple gestation or without postpartum follow-up were excluded.

The fetal crown rump length (CRL) was measured, and if it was between 45 and 84 mm, we evaluated the following fetal ultrasound parameters: nuchal translucency (NT), nasal bone (NB), and ductus venosus (DV) flow. After that, the PAPP-A and free β-hCG levels in the maternal serum were determined. The sample was analyzed by means of a fluoroimmunometric assay using an automated AutoDelfia system (Perkin Elmer, Wallac, Turku, Finland). Analysis of NT thickness, PAPP-A and free β-hCG was performed using the algorithm provided by the FMF, in London, UK, and was calculated using the Astra software (Astra software gmbh, Munich, Germany).

The ultrasound parameters were evaluated only by experienced sonographers who had been certified by the FMF for 11–13 weeks’ scan. The measurements were taken using a transabdominal transducer (5 MHz curvilinear transducer, Voluson E6 [GE Healthcare, Milwaukee, WI, USA]).

First-trimester risk assessment was provided for trisomy 21, trisomy 18 and trisomy 13. The risk was calculated using a previously described algorithm.20 The cut-off point for high risk for aneuploidies was defined as greater than 1:100, intermediate risk was defined to be between 1:100 and 1:1,000 and low risk was defined as less than 1:1,000.4 The variable aneuploidy was considered as a result not only of trisomy of chromosome 21 but also the trisomy of chromosomes 13 and 18. Patients classified as high or intermediate risk were referred for fetal medicine counseling. Chorionic villus sampling or amniocentesis was performed for karyotype analysis in women who expressed their wish and signed an informed consent.

Information about patient characteristics, chromosomal abnormalities and the pregnancy outcome was obtained by the personnel, hospital registry or the postpartum routine follow-up registry.

The statistical analysis was performed using the SPSS software version 20.0 (IBM Corp., Armonk, NY, USA). A descriptive analysis of the population was performed in the form of mean and median with standard deviation (SD) for quantitative variables and the proportions, percentages and ratios by calculating the 95% confidence intervals (95% CIs) for categorical variables.

We calculated detection rates (sensitivity), specificity, positive and negative predictive values and percentage of FP for aneuploidies. **P**-values < 0.05 were considered statistically significant. For quantitative variables, the Mann-Whitney test was used for comparison between two independent groups, and for categorical variables, we used the Chi-square test. A ROC curve was constructed to estimate the sensitivity and specificity of aneuploidy screening for various risk cut-off points. The Spearman coefficient was calculated to evaluate the correlation of four factors (age, NT, β-hCG and PAPP-A) with the risk for aneuploidies in the first trimester.

## Results

Among 3,639 pregnant women who underwent a combined test in the first-trimester risk assessment at the perinatal group, a total of 2,748 hospital registries were analyzed. Among the patients that were excluded, there were 775 that were lost to follow-up after delivery or the pregnancy resulted from ovum donation, and 116 cases of multiple gestation. The ethnic origin of the pregnant women was almost all Latin American.

The median CRL was 62.5 mm (with a range from 45.0 to 84.0 mm). The median maternal age was 33 years (with a range from 18 to 46 years). A total of 1,142 (41.6%) of the women were aged 35 years or older, and 1,606 (58.4%) women were aged between 18 and 34 years old. A total of 173 (6.3%) of the women were aged 41 years or older.

Considering the cut-off point for high risk for aneuploidies (risk greater than 1:100), 62 (2.3%) women were classified as high risk, 155 (5.6%) as intermediate risk (between 1:100 and 1:1,000), and 2,531 (92.1%) as low risk (less than 1:1,000). In the population that was screened, there were 21 pregnancies (0.76%) detected for aneuploidy.

The Spearman coefficient was calculated in our population for four algorithm composition factors, age, TN, β-hCG and PAPP-A, to analyze the influence of these factors in the risk assessment. Among the factors that influence the risk for aneuploidies, age is the one with the greatest weight (Table 1). Also, if we analyze these four variables separately, we can see that all of them influence the risk classification with a statistically significant **p**-value (Table 2). In the groups aged under 34 years, between 35 and 40 years and over 41 years, 71 (4.4%), 77 (7.9%) and 69 (39.9%) patients, respectively, were classified as intermediate and high risk and were referred for fetal medicine counseling (Table 3). When stratifying by age group, we found that the chance of being classified as high/medium risk is 1.8 times higher in the age group between 35 and 40 years and up to 9 times higher in the age group over 41 years, in relation to the age group up to 34 years (Table 3).

In the present report, the detection rate for aneuploidy or sensitivity was 42.9%, specificity was 98.1%, for a FP rate of 1.9%, positive predictive value (PPV) of 14.52% and negative predictive value (NPV) of 99.55% for a 1:100 cut-off point. The first-trimester combined test achieved 71.4% sensitivity with a 7.4% FP rate, specificity of 92.6%, PPV of 6.91% and NPV of 99.76%, when the cut-off point was adjusted to greater than 1:1,000 (Table 4).

### Table 1 Influence of algorithm composition factors in the studied population’s first-trimester risk assessment

<table>
<thead>
<tr>
<th>Algorithm Factor</th>
<th>Risk for aneuploidies correlation</th>
<th>Spearman</th>
<th><strong>p</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td>Maternal Age</td>
<td>-0.770</td>
<td>&lt; 0.001</td>
<td></td>
</tr>
<tr>
<td>NT</td>
<td>-0.111</td>
<td>&lt; 0.001</td>
<td></td>
</tr>
<tr>
<td>Free β-hCG</td>
<td>-0.122</td>
<td>&lt; 0.001</td>
<td></td>
</tr>
<tr>
<td>PAPP-A</td>
<td>0.163</td>
<td>&lt; 0.001</td>
<td></td>
</tr>
</tbody>
</table>

Abbreviations: β-hCG, beta-human chorionic gonadotropin; NT, nuchal translucency; PAPP-A, pregnancy-associated plasma protein A.
When we plotted the ROC curve for our population, the cut-off point that maximized the sensitivity and specificity for the diagnosis of aneuploidies was 1:1,860. If we adjust the FP rate to 5%, the detection rate for this analysis is 72.7% with a cut-off point of 1:610 (**Table 5**).

### Discussion

Many authors have reported that a combined screening test for aneuploidy makes sense because it results in a sensibility of 80% and higher.\(^{12,21-26}\) However, many critics of the methodology used by these authors cast doubt on these results.\(^{14,27,28}\) Therefore, it is important to analyze the combined screening test performance in our own population.

In our study, the FP rate for the first-trimester combined test for chromosomal abnormalities altogether, considering the cut-off of 1:1,000, was 7.4%, which was similar to the range of 5 to 7% found in other studies. It was somewhat higher than the usual set value of 5%. The detection rate was 71.4%, which is less than the values of 75.9 to 95% produced by other studies.\(^{6-8,14,28}\)

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**Table 2** Correlation between four variables (age, nuchal translucency, \(\beta\)-hCG and PAPP-A) and the risk classification of aneuploidy

<table>
<thead>
<tr>
<th>Variables and risk for aneuploidy</th>
<th>n</th>
<th>Average</th>
<th>SD</th>
<th>Minimum</th>
<th>Median</th>
<th>Maximum</th>
<th>P-value*</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Age (years)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>High/intermediate risk</td>
<td>217</td>
<td>37.1</td>
<td>5.5</td>
<td>22.0</td>
<td>38.0</td>
<td>50.0</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>Low</td>
<td>2,531</td>
<td>33.3</td>
<td>4.2</td>
<td>18.0</td>
<td>33.0</td>
<td>49.0</td>
<td></td>
</tr>
<tr>
<td><strong>Nuchal translucency (mm)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>High/intermediate risk</td>
<td>217</td>
<td>2.04</td>
<td>1.13</td>
<td>0.70</td>
<td>1.80</td>
<td>7.20</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>Low</td>
<td>2,531</td>
<td>1.50</td>
<td>0.47</td>
<td>0.50</td>
<td>1.50</td>
<td>9.00</td>
<td></td>
</tr>
<tr>
<td><strong>(\beta)-hCG (MoM)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>High/intermediate risk</td>
<td>216</td>
<td>1.71</td>
<td>1.40</td>
<td>0.09</td>
<td>1.25</td>
<td>8.34</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>Low</td>
<td>2,529</td>
<td>2.06</td>
<td>24.42</td>
<td>0.01</td>
<td>1.01</td>
<td>933.00</td>
<td></td>
</tr>
<tr>
<td><strong>PAPP-A (MoM)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>High/intermediate risk</td>
<td>215</td>
<td>0.915</td>
<td>0.702</td>
<td>0.029</td>
<td>0.680</td>
<td>3.964</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>Low</td>
<td>2,499</td>
<td>1.437</td>
<td>0.960</td>
<td>0.108</td>
<td>1.228</td>
<td>9.987</td>
<td></td>
</tr>
</tbody>
</table>

Abbreviations: \(\beta\)-hCG, beta-human chorionic gonadotropin; MoM, Multiple of the Median; PAPP-A, pregnancy-associated plasma protein A; SD, standard deviation.

**Table 3** Combined test risk classification for aneuploidies in the first trimester by maternal age group (\(n = 2,748\))

<table>
<thead>
<tr>
<th>Patient Age</th>
<th>Low risk for aneuploidy n (%)</th>
<th>Intermediate and high risk for aneuploidy n (%)</th>
<th>Total n (%)</th>
<th>Intermediate and high risk PR (CI95%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>≤ 34 years</td>
<td>1,535 (95.6)</td>
<td>71 (4.4)</td>
<td>1,606 (100)</td>
<td>1</td>
</tr>
<tr>
<td>35–40 years</td>
<td>892 (92.1)</td>
<td>77 (7.9)</td>
<td>969 (100)</td>
<td>1.8 (1.3–2.5)</td>
</tr>
<tr>
<td>≥ 41 years</td>
<td>104 (60.1)</td>
<td>69 (39.9)</td>
<td>173 (100)</td>
<td>9.0 (6.7–12.1)</td>
</tr>
<tr>
<td>Total</td>
<td>2,531 (92.1)</td>
<td>217 (7.9)</td>
<td>2,748 (100)</td>
<td></td>
</tr>
</tbody>
</table>

Abbreviations: CI, confidence interval; PR, prevalence ratio.

**Table 4** Detection rate or sensitivity, specificity, positive and negative predictive values for aneuploidy at different cut-off points

<table>
<thead>
<tr>
<th>Risk cut-off</th>
<th>Occurrence of aneuploidy</th>
<th>Predictive values</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>1/100</td>
<td>9</td>
<td>53</td>
</tr>
<tr>
<td>1/1,000</td>
<td>12</td>
<td>2,674</td>
</tr>
<tr>
<td>1/1,860</td>
<td>17</td>
<td>390</td>
</tr>
</tbody>
</table>

When we plotted the ROC curve for our population, the cut-off point that maximized the sensitivity and specificity for the diagnosis of aneuploidies was 1:1,860. If we adjust the FP rate to 5%, the detection rate for this analysis is 72.7% with a cut-off point of 1:610 (**Table 5**).
The detection rate was much lower (42.9%) if we consider the cut-off to be 1:100, for a FP rate of 1.9%. The best result that maximized the sensitivity and specificity for the diagnosis of aneuploidies in our population was 1:1,860. Most likely, our results were influenced by maternal age, because 41.6% of the women were aged 35 years or older. The influence of maternal age on combined screening test performance was also described in the Chinese population. Pan et al. demonstrated that the FP rate increases with increasing maternal age.

The limitation of this study is a small sample size in comparison with larger studies. Additionally, because this study was conducted in only one center, the results cannot fully represent the screening performance in all of the Brazilian population. In addition, the overall screening performance might be affected by the test timing. In this study, the first-trimester combined test was performed between 11 + 0 and 13 + 6 weeks of gestation. Both serum markers and NT are affected by the gestational age of the fetus. The discrimination of PAPP-A is greatest at 10 weeks and declines afterwards, whereas screening performance of free β-hCG improves with increasing gestational age until 13 weeks. Additionally, there might be a difference in screening power depending on the gestational age of serum marker measurements. The NT was measured by multiple observers; therefore, there is a possibility of error as a result of inter-observer variation. Finally, there were 775 women who were lost to follow-up at the center. There is a possibility that an error has occurred with these women who had no records on aneuploidy screening.

Conclusion
In conclusion, the combined test of aneuploidies screening showed a detection rate inferior to those described in the literature for a higher FP rate. It may suggest that we have to consider a different cut-off point (1:610) as a reference for the population assisted at our maternity center to achieve similar performance to the literature.

Contributions
Abib L. P. A., Sá R. A. M. and Peixoto-Filho F. M. contributed with project and interpretation of data, writing of the article, critical review of the intellectual content and final approval of the version to be published.

Conflicts of Interest
The authors declare that there are no potential conflicts of interest.

References


