

NON-INVASIVE SCREENING - THE PROBABILITY OF EVENTS

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Abstract- Congenital anomalies are the cause of 20.0-25.0% of cases of perinatal death, while 3.0% of children are born with malformations of varying size. We examined the predictive values and defined the credibility ratio of the combined test results. Sensitivity of the test is 94.0%, and specificity is 99.0%. The positive likelihood ratio [likelihood ratio test (LR+)] is 94.00; a negative likelihood ratio [likelihood ratio test (LR-)] is 0.06. The pretest probability that pregnant women carry a fetus with chromosomal abnormality is 1:250. Posttest odds after the combined test to discover this abnormality are 0.3760, and probability of the same case is 0.2732 if it happens that the test result is positive. The result of our study confirms the justification of combined test usage in routine clinical practice, since the posttest odds rate in the case of a positive screening increases several times over (almost 90 times); the probability of detecting a chromosomal abnormality was about 70 times.

Key words: Predictive value, combined test, ultrasonography, biochemical markers

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INTRODUCTION

Congenital anomalies cause 20.0-25.0% of perinatal deaths, while 3.0% of children are borne with malformations of varying size (Ebrahaim and al., 2000). Analytical immunochemiluminescence assays and an automated analyzer IMMULITE 2000 [Diagnostics Product Corporation (DPC), Los Angeles, California, USA] were used. Usable values of the combined test were estimated on the basis of its sensitivity, specificity and possibility of the disease in case the result was positive. By combining the values of pregnancy-associated plasma protein A (PAPP-A) and free β -subunit of choriogonadotropin (free β -HCG) in serum with nuchal translucency (NT) diameter (combined test), the possibility of detecting trisomy

21 rises up to 90.0% with 5.0% false-positive findings (Alfirevic and Neilson, 2004). The testing was done between 11 and 13⁺⁶ weeks of gestation. If the result was positive, some invasive methods of prenatal diagnosis were suggested to the pregnant woman. A limit value of the combined test was 1: 250. A special problem was the test result interpretation. According to the literature, 32.0% of pregnant women did not know what the term "high risk" meant after getting the results and talking to the doctor (Spencer, 2001). The research objective was: 1) to examine the sensitivity and specificity of ultrasonographic (NT) and biochemical (free β -HCG and PAPP-A) markers as parameters of the combined test, and amniocentesis in diagnostics of congenital fetal anomalies; and 2) to set the credibility ratio of the combined test results.

MATERIALS AND METHODS

The prospective observational study was conducted at the Gynecology and Obstetrics Clinic at the Clinical Center Kragujevac (GOC, CC Kragujevac, Kragujevac, Serbia) during 2008 and 2009 on singleton intrauterine pregnancies in the first trimester of pregnancy. The clinical and experimental model of study was used throughout the research. The Ethics Committee at the CC Kragujevac confirmed the correctness of this study and authorized its conduct. Three hundred and seventeen pregnant women were included in the examination and observed by the Board of Genetic Counseling at GOC, CC Kragujevac. An informed consent form was signed by all participants. All ultrasound examinations were conducted on the Aloka Prosound 3500 Ultrasound Machine Package with UST-9124 (Aloka, Seoul, South Korea) at GOC, CC Kragujevac. Pregnancy was 11-13⁺⁶ weeks of gestation. The crown-rump length (CRL) of the fetus was between 45 and 84 mm. While measuring fetal nuchal translucency (NT), we used an ultrasound apparatus of high resolution with the option "cine loop" so that the image could be turned by calipers that allow measurements of one decimal point. The screen image on which NT was measured encompassed only the head and the upper part of fetal rib cage. We used the maximum enlargement, so that small movements of the caliper altered the diameter by only 0.1 mm. The NT was measured by transvaginal approach of color Doppler technique with the fetus in a neutral position. We measured the maximal thickness of subcutaneous illumination between the skin and soft tissue located above the cervical part of the fetal spine. We set the calipers on lines that define the fold, so that they were barely visible on the white limit line of the accumulation behind the neck. During the examinations we conducted more measurements and took into consideration only the highest thickness of the NT. We carried out the quantitative settings of free β -HCG and PAPP-A from the venous blood of patients by applying commercial tests from DPC. Tests were based on the analytical immunochemiluminescence assay and were realized by using the automated analyzer IMMULITE 2000 (DPC).

All the pregnant women included in this study underwent amniocentesis by a standard transabdominal procedure with ultrasound control using the "free hand" technique at 16 to 17⁺⁶ weeks of gestation. We did the puncture by needles with mandrene of 20-22G thickness and aspirated 15-20 mL of amniotic fluid into a syringe without a rubber seal. The amniotic fluid sample was delivered to the Cytogenetics Laboratory at the GOC, CC Kragujevac, and all results received were deposited into the unique database with the required logistic control.

RESULTS

After conducting the combined test in the total sample of pregnant women, we found the following individual values of the examined parameters (see Table 1): the statistically significant difference in values of free β -HCG and NT in the examined group of pregnant women was $p < 0.05$. Parameter PAPP-A does not show any statistically significant difference in the examined group of pregnant women. We also found the same characteristics of the examined parameters in the receiver operating characteristic (ROC) curve analyses in the examination of the predictive characteristics of the specified parameters (Figs. 1 and 2).

Analysis of the value distribution of the NT thickness measurement showed that the distribution was regular and that measurements were being set regularly around the median (44.0% below and 56.0% above the median), which was in accordance with the criteria for quality control established by the Fetal Medicine Foundation (London, UK) and was supposed to be 40.0-60.0% above the median. The distribution of fetal NT for given CRL in examination was no different from the established distribution of the Fetal Medicine Foundation used as a standard. On the basis of that, our measurements of NT thickness can be considered to be regularly conducted and usable in further examination.

The diameter of the NT did significantly statistically differ in the examined group of pregnant women ($p < 0.05$). Crown-rump length and gestational age were not different statistically ($p > 0.05$). Using the

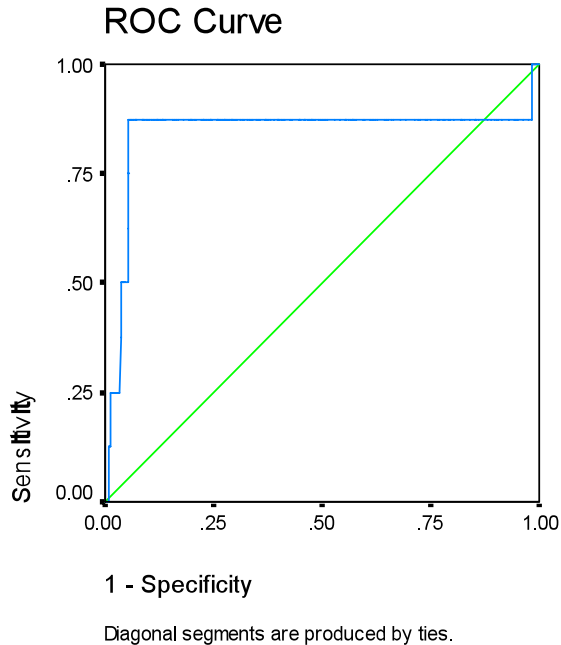


Fig. 1. Probability of the predictive value of free fraction β -HCG [field below ROC curve 0.846, 95% confidence interval (95% CI) 0.628-1.064].

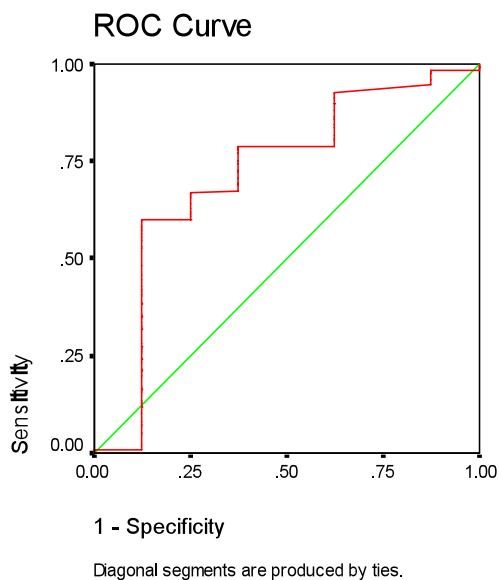


Fig. 2. Probability of the predictive value of PAPP-A [field below ROC curve 0.715 (95% CI 0.508–0.921)].

contingency table (Table 2), we set the predictive value of the combination of ultrasonographic and

biochemical markers after taking over the results of amniocentesis.

Estimation of probability that some disease is present before testing is called pretest probability (“a priori probability”). Pretest probability is received on the basis of available information about the patient, including testing previous to the actual one. Estimation of the probability of disease after testing is called posttest probability (“a posteriori probability”). Posttest probability is less or higher than pretest probability depending on the test results. Measures of diagnostic accuracy (sensitivity, specificity) cannot directly answer the following important clinical questions: 1) if the disease pretest probability is known, and the examinee is positive on the test, what is the probability that he/she really has the disease? 2) If the disease pretest probability is known, and the examinee is negative on the test, what is the probability that he/she really does not have the disease?

These questions can be answered by the application of the pretest odds of the disease and the credibility ratio. Disease odds ratio is the ratio of probability that the disease is present (p) and probability that is not present ($1-p$): $\text{odds} = p/1-p$. According to this, pretest disease odds are: $\text{pretest odds} = \text{pretest probability}/1-\text{pretest probability}$. Likelihood ratio (LR) is the probability ratio of a certain test result (+ or –) of the examinee who has the disease divided with the probability of the same result of the person who does not have the disease. Two types of likelihood can be calculated: 1) likelihood ratio of the positive test (LR+) is the ratio of sensitivity and false positive ratio ($1-\text{specificity}$): $\text{LR+} = \text{sensitivity}/1-\text{specificity}$; 2) likelihood ratio of the negative test (LR–) the ratio of sensitivity and false negative ratio ($1-\text{sensitivity}$) and specificity: $\text{LR-} = 1-\text{sensitivity}/\text{specificity}$. The likelihood ratio shows how the test result can alter the pretest disease probability. The LR+ shows how much the test result increases disease probability, LR– shows how much the test result decreases disease probability. The likelihood ratios are not under the influence of the disease prevalence.

Table 1. Difference between values of free β -HCG, PAPP-A and NT in 317 pregnant women (total sample).

Parameters	Free β -HCG (mL/mL)	PAPP-A (mL/mL)	NT (mm)
Pathological karyotype	114.00	1.36	2.55
Physiological karyotype	19.20	1.84	1.90
Mann-Whitney test (U)	704.50	2191.00	621.50
<i>p</i> Value	0.000	0.543	0.000

Table 2. Review of the middle values (MV) and standard deviations (SD) of the ultrasonographic parameters and gestational age in 317 pregnant women

Parameters	Pathological Karyotype: 16	Control Group: 301	<i>p</i> Value
Nuchal translucency (mm)	2.49 \pm 0,37	1.92 \pm 0,39	<0.05
Crown-rump-length (mm)	60.12 \pm 8.48	64.83 \pm 8.23	>0.05
Gestational age (days)	86.69 \pm 3.98	87.40 \pm 7.40	>0.05

Table 3. Contingency table created on the basis of data processing in the total sample of pregnant women after amniocentesis

Test Result	Disease Present	Disease Absent	Total
Positive	Really positive: 15	False-positive: 1	RP+FP: 16
Negative	False-negative: 3	Really negative: 298	FN+RN: 301
Total	18	299	317

Table 4. Probability and predictive values of the parameters of the combined test in relation to result of early amniocentesis

Parameter	<i>p</i> Value	95% CI
Sensitivity	0.9375	0.6977-0.9984
Specificity	0.9900	0.9712-0.9979
Positive predictive value	0.8333	0.5858-0.9642
Negative predictive value	0.9967	0.9815-0.9999
Prevalence	0.0505	0.0291-0.0807
False-positive rate	0.0100	0.0046-0.245
False-negative rate	0.0625	0.0145-0.0998
Positive likelihood ratio (LR+)	94.0000	30.2937-292.0653
Negative likelihood ratio (LR-)	0.0631	0.01746-1.2712
Overall test accuracy	0.9873	0.07215-0.9981

Likelihood ratios can help measuring the posttest probability. How big the change from pretest to posttest probability depends considerably on the values of the likelihood ratio. It is desirable for (LR+) to have the highest values and (LR-) to have values closest to 0. For calculating the posttest disease probability,

posttest odds are first to be calculated. 1) For positive test result: posttest odds = pretest odds \times LR+; 2) for negative test result: posttest odds = pretest odds \times LR-. Posttest probability is obtained by the formula: posttest probability = posttest odds / (1 + posttest odds). According to the literature data, the diagnostic ac-

Table 5. Influence of the combined test on the pretest odds and probability of the outcome in relation to the likelihood ratio in case of positive and negative outcome

Parameter	p Value
Pretest odds/probability	0.0040
Posttest odds/positive test	0.3760
Posttest odds/negative test	0.0002
Likelihood ratio/LR+	94.000
Likelihood ratio/LR-	0.0600
Posttest probability/positive test	0.2732
Posttest probability/negative test	0.0002

curacy of the combined test, in relation to the result of the early amniocentesis (referral standard) is: sensitivity 0.88, specificity 0.90. In our sample sensitivity is 0.94 and specificity 0.99. Likelihood ratios are: $LR+ = 0.94/1-0.99 = 94.00$; $LR- = 1-0.94/0.99 = 0.06$. Pretest probability that the pregnant woman carries a fetus with chromosomal abnormality is $1:250 = 0.004$. Pretest odds $= 0.004/0.996 = 0.004$. If the test is positive: posttest odds $= \text{pretest odds} \times LR+ = 0.004 \times 94 = 0.3760$; posttest probability $= \text{posttest odds}/1 + \text{posttest odds} = 0.3760/1 + 0.3760 = 0.2732$. If the test is negative: posttest odds $= \text{pretest odds} \times LR- = 0.004 \times 0.06 = 0.00024$. Posttest probability $= \text{posttest odds}/1 + \text{posttest odds} = 0.00024/1 + 0.00024 = 0.00024$.

DISCUSSION

In our study, we had nine (2.84%) pregnant women with numeric aberrations in total and seven (2.21%) pregnant women with structural aberrations in fetuses which could be explained by the fact that the sample was preselected. All pregnant women were sent to Genetic Counseling at the GOC, CC Kragujevac (Table 2), for some suspected reason (positive personal and/or family case history, age of the pregnant woman, likelihood of giving birth to a child with chromosomal aberrations and/or fetal anomalies in previous pregnancies, *etc.*). Similar results were reported in the study conducted in Great Britain in 2000 which stated that the total incidence of Down's syndrome was 1 in 1,000 deliveries, which was 50.0%

more than in the national reports (Wald et al., 1999). The importance of NT measurement in the screening for Down's syndrome during the first trimester of the pregnancy was recognized back in 1990. With a limit value of 3 mm NT thickness, the detection rate (DR) is 64.0% (Wald et al., 2003). Screening sensitivity of chromosomopathies in comparison to NT was 75.0% with a value of false-positive ratio of 2.1% (Stojiljkovic-Mikic and Rodeck, 2003). In our sample, 11 pregnant women in total from the group of 16 had a measured value of NT above the median for the given CRL in the group of pathological karyotypes that was 68, i.e. 75.0%. By the analysis of the total sample, we found that with 26 pregnant women we measured a NT of 2.55 mm above median for the given CRL and by invasive diagnostics we confirmed 16 cases of chromosomal fetal aberrations (61.54%) (Table 3). The methodology of the combined test (Table 1) indicated that the ultrasound screening was done first, and after that the level of free β -HCG and PAPP-A was set, where risks are calculated as the combination of the two data (Haddow et al., 1998). For a certain gestation time, the levels of free β -HCG and PAPP-A represent the factor of probability, which is multiplied by the initial risk in order to calculate a new risk (Wald et al., 1996). Differences in the concentration of free β -HCG between normal pregnancies and those with trisomy 21 increase, and differences in the level of PAPP-A decrease, with the age of the pregnancy. There is no significant connection between thicknesses of the fetal NT, level of free β -HCG or PAPP-A in maternal serum in pregnancies with trisomy 21 in relation to normal pregnancies, so ultrasound and biochemical markers can be combined in order to get more efficient screening results. Numerous studies have confirmed the connection between the low level of PAPP-A and trisomy 21 during the first trimester (Alfirevic and Neilson, 2004). In normal pregnancies, the level of PAPP-A in maternal blood increases with gestation time, and in pregnancies with trisomy 21 it decreases [multiple of median (MoM)] (<0.5). By setting the value of PAPP-A, it is possible to detect 52.0% of Down's syndrome cases with 5.0% false-positive results (Alfirevic and Neilson, 2004). In pregnancies with trisomy 21, the level of free β -HCG is increased between 8

and 14 weeks of gestation. The level of free β -HCG in maternal blood decreases normally with gestation time, and in pregnancies with trisomy 21, the level of free β -HCG increases (MoM >2.0) (Snijders et al., 1996; Cunningham et al., 2005). On the basis of free β -HCG level, DR amounts to 42.0% with 5.0% false-positive findings (Wellesley et al., 2002). The frequency of false-positive results, according to the available literature, is estimated at 5.0% (Loncar et al., 2010; Hallahan et al., 2000; Brigatti and Malone, 2004; Nicolaides, 2004). Our research has shown that the rate of false-positive findings is 1.0%, and that free β -HCG is a more sensitive predictor than PAPP-A. Other investigators have reported identical conclusions (Brigatti and Malone, 2004). The predictive value of the individual biochemical markers is represented in Charts 1 and 2 by setting the area below the ROC curve. On reviewing the combined test predictive value of our sample of pregnant women, we found the following results: sensitivity of the test is 94.0%, specificity is 99.0%. The positive predictive value of the test is 0.83, and negative predictive value of the test is 0.99 (99.0%). The positive likelihood ratio (LR+) is 94.00 and negative likelihood ratio (LR-) is 0.06 (Tables 4 and 5). We have already confirmed the published positive qualifications of this screening method (Spencer et al., 1999) and point to its justification in every day clinical practice (Krantz et al., 2000). Regarding posttest odds, the rate in the case of positive screening increases several times over (almost 90 times). In the available literature, we have not found any reports that have the calculation of the credibility of the combined test and prediction of posttest odds of this screening method. It is most important to tell the patients that this is a process of screening and not a final diagnosis. This can be given only on the basis of invasive intervention and defining of the fetal karyotype (Spencer, 2000).

CONCLUSIONS

By examining the sensitivity and specificity of the combined screening test in the period of 11.-13⁺⁶ weeks gestation, we found that the sensitivity of the test was 0.94 (94.0%), and its specificity 0.99 (99.0%). Pretest probability that a pregnant woman carries a

fetus with a chromosomal abnormality is 1:250 or 0.004. Posttest odds after the combined test to discover this abnormality were 0.3760, and the probability of the same case was 0.2732 if the test result was positive. Posttest odds after the combined test and the probability of the same case were identical if the test result was negative and amounted to 0.0002.

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