

Prevalence of Chromosomal Abnormalities in Spontaneous Abortions or Fetal Deaths



Estudo da Prevalência de Anomalias Cromossômicas em Abortamentos Espontâneos ou Mortes Fetais

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ABSTRACT

Introduction: Spontaneous abortion is one of the most frequent problems of pregnancy, estimated to affect, at least, one in every four women who tries to conceive. The main purpose of this work was to study the prevalence of chromosomal abnormalities in gestational losses, evaluating their relation with maternal age, gestational age and previous abortion history.

Material and Methods: Retrospective study of 401 pregnancy loss cases that have gone through cytogenetic and anatomopathologic analysis between January 2008 and June 2012, in Centro Hospitalar de S. João, Oporto.

Results: Of the 401 cases sent to cytogenetic study, it was possible to obtain information about the chromosomal complement in 333 cases, of which 72.7% showed normal karyotype, and 27.3% abnormal karyotype. Aneuploidies represented 92.3% of the identified chromosomopathies, with trisomies being the most frequent, related with an advanced maternal age and an early gestational age. There was no agreement between the results of the cytogenetic and the anatomopathologic analysis.

Discussion/Conclusion: The prevalence of chromosomal abnormalities, during the first trimester, is similar between sporadic and recurrent miscarriages. With increased maternal age, trisomies, the most frequent type of aneuploidy, are more likely to occur, with a mean increment in probability of 7.4% per year. A significant karyotype-pathological correlation was not established. Maternal contamination is the main obstacle to the accurate determination of the prevalence of chromosomal abnormalities. The molecular cytogenetic techniques already available can overcome the limitations of the conventional technique.

Keywords: Abortion, Spontaneous/genetics; Chromosome Aberrations; Chromosome Disorders; Infertility, Female/genetics; Portugal.

RESUMO

Introdução: O abortamento espontâneo é um dos problemas mais frequentes da gravidez, estimando-se que afete, pelo menos, 25% das mulheres que tentam engravidar. O objetivo principal deste trabalho foi o estudo da prevalência das anomalias cromossômicas em perdas gestacionais, analisando a sua relação com a idade materna, idade gestacional e história de abortamentos prévios.

Material e Métodos: Realizou-se um estudo retrospectivo em 401 casos de perdas de gravidez que efetuaram análise citogenética e anátomo-patológica, entre janeiro de 2008 e junho de 2012, no Centro Hospitalar de S. João.

Resultados: Dos 401 casos enviados para estudo citogenético foi possível obter informação sobre o complemento cromossômico em 333 casos, dos quais 72,7% apresentaram cariótipo normal e 27,3% cariótipo anormal. As aneuploidias representaram 92,3% das cromossomopatias identificadas, sendo as trissomias as mais frequentes, associadas a uma idade materna avançada e a uma idade gestacional precoce. Não se verificou concordância entre os resultados da análise anátomo-patológica e citogenética.

Discussão e Conclusão: A prevalência de anomalias cromossômicas, no primeiro trimestre, foi semelhante entre os casos de abortamento esporádico e recorrente. Com o aumento da idade materna observou-se um aumento das trissomias, com um incremento médio de 7,4% no risco de ocorrência por ano. Não foi estabelecida uma correlação cariótipo-patológica significativa. A contaminação materna da amostra foi o principal entrave à determinação exata da prevalência de anomalias cromossômicas. As técnicas de citogenética molecular já disponíveis podem colmatar esta e outras limitações da técnica convencional.

Palavras-chave: Abortamento Espontâneo/genética; Aberrações Cromossômicas; Anomalias Cromossômicas; Infertilidade Feminina/genética; Portugal.

INTRODUCTION

Spontaneous miscarriage is a common problem during pregnancy and it is estimated to affect 25 to 50% of women trying to get pregnant, although only 15% of the cases are clinically detected.^{1,2} It is known that most miscarriages occur over the first weeks of pregnancy, especially until the 14th week.¹ Although most miscarriages are sporadic, 1-3% of the couples present recurrent miscarriage, defined as consecutive loss of three or more pregnancies, with a significant psycho-social impact.^{1,3,4}

Several illnesses have been considered as a cause for sporadic miscarriage, with chromosomal conception abnormalities representing approximately 50-60% of the

cases.^{2,5,6} Most of these abnormalities are numerical (95%) and trisomy is the most prevalent (60%), more frequently trisomy 16, followed by polyploidy and X monosomy.^{1,2} As regards recurrent miscarriage, the literature is not so consensual as to the frequency of chromosomal abnormalities, some stating that they occur with a similar frequency as sporadic miscarriage⁷⁻¹⁰ while others argue for a lower frequency.^{11,12}

Maternal age increase is related with an increase in chromosomal conception abnormalities and consequently with spontaneous miscarriages.^{2,10,13,14} Over the years and worldwide a progressive maternity delay with a significant

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increase in the age of the first pregnancy is becoming evident. In Portugal, during a 10 year period (1998-2008), an increase in the number of deliveries above 40 (from 1.8% to 3.2%), as well as a change in the age group in which most cases occurred (from 25-29 to 30-34) have been observed.¹⁵ This increase in maternal age is related with a higher spontaneous miscarriage and foetal loss rate, following a J-curve with a steep rise at the age of 35.¹⁶ This fact represents an increase in the proportion of abnormal foetal karyotypes: 51-54% in women aged below 35 and 74-76% in older women.^{10,17}

Taking into account the chances at the time of fertilisation, male and female embryos would be expected in equal numbers. Nevertheless, there is a distortion at the time of delivery, with a higher number of male foetus in an approximate relation of 1.05^{18,19} (1.053 in Portugal, during 2011).²⁰

Therefore, taking into account the huge impact of chromosomal abnormalities in the success of pregnancy, our study aimed mainly to identify the most common chromosomal abnormalities related with spontaneous miscarriages or foetal losses and to assess their relation to gestational age, maternal age and previous miscarriages. In addition, we have also assessed the relative number of miscarriages per gender and found a relation of these results with cytogenetic and histopathological examinations.

MATERIAL AND METHODS

Our study involved a retrospective analysis of the results obtained through cytogenetic and histopathological examination of spontaneous miscarriage or foetal loss products, at the *Centro Hospitalar de S. João* between January 2008 and June 2012.

Clinical records were consulted and information obtained on mother's age and past obstetric history, gestational age at the time of pregnancy loss and results of genetic and histopathological examination of miscarriage products. This study was approved by the Ethics Committee of the *Centro Hospitalar de São João/Faculdade de Medicina da*

Universidade do Porto (FMUP).

The criteria for cytogenetic examination were the following: termination of pregnancy; foetal loss and late pregnancy loss; repeated early pregnancy loss (more than 2) and miscarriage in a couple with a known chromosomal abnormality. For the study of spontaneous miscarriages, the terminations of pregnancy were excluded, in order to avoid a possible bias in the results. Beyond those meeting the previously described criteria, our group of patients also included spontaneous miscarriages occurring in primigravidae, as part of a previous research project.

Taking into account the gestational age at the time of miscarriage, the products that were examined were assigned according to the gestational trimester: the first until the 12th week, the second between 13th and 24th and the third from the 24th week of pregnancy. Three age groups were also defined for the assessment of the impact of maternal age on the results: younger than 35, between 35 and 39 and above 40 years of age.

The decision was taken to establish six groups taking into account the results suggested by the histopathological examination, in order to establish a correlation between the histopathological and cytogenetic information (normal or abnormal karyotype: (1) probable chromosomal abnormality; (2) without an apparently adequate implantation; (3) apparent growth disorganization of foetus/foetal tissue; (4) probable infection; (5) signs of foetal growth restriction; (6) other indications.

The genetic study involved carrying out the Multiplex Ligation-dependent Probe Amplification (MLPA) technique, using the commercial kit P095 (MRC Holland, Amsterdam, The Netherlands) and GTL-banding karyotyping, according to previously established protocols in the Department of Genetics at the FMUP, where this study was carried out.^{21,22}

Data statistical study used Microsoft Office Excel 2007 and IBM SPSS Statistics 20 software. We used the Student t-test to validate the sample regarding distribution and dispersion, as well as to test significance between variables, together with the chi-square test, as appropriate. A logistic

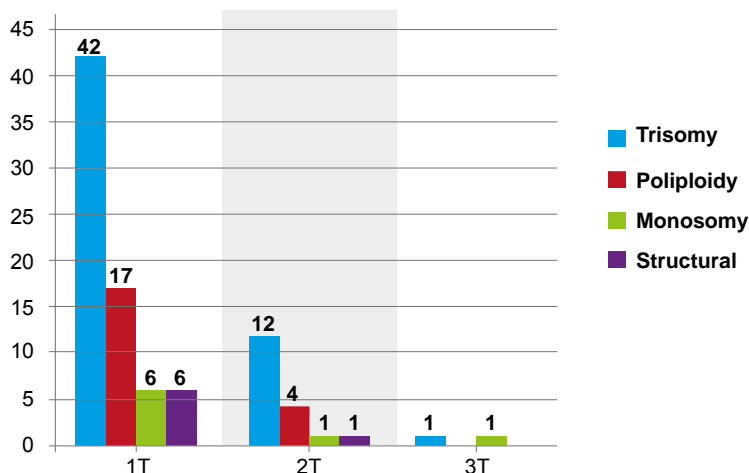


Figure 1 - Absolute number of chromosomal abnormalities identified in each pregnancy trimester

Table 1 - Description of the structural abnormalities found and their clinical information

Structural Abnormalities	Maternal age	Recurrence	Trimester
45,X[26]/46,X,del(X)(q)[22]/46,XX[2]	< 35	No	1 st
46,XX,der(8)t(6;8)(q13;p23,1)mat	< 35	Yes	1 st
46,XX,dup(15)(q12q13)mat	35 – 39	Yes	1 st
45,XX,dup(1q),der(13;14)	35 – 39	No	2 nd
46,XX, der(22)	35 – 39	No	1 st
46,XX,i(8)(q11,1)	< 35	No	1 st
47,XX,mar[3]/46,XX[27]	< 35	No	1 st

regression model was also used for the multivariable analysis and risk modelling. We considered as significant p values < 0.05.

RESULTS

According to the above criteria, a cytogenetic examination was carried out in 401 spontaneous miscarriage or foetal loss products, from which 251 occurred in the first, 118 in the second and only 32 occurred in the third trimester of pregnancy.

The chromosomal analysis was not conclusive, as cells failed to grow in culture in 20.0% (80/401) of the cases. This failure occurred in 21.5% (54/251) from the first trimester cases, 12.7% (15/118) from the second and 34.4% (11/32) from the third. In 12 of these 80 cases, the analysis using the MLPA technique allowed for identification of aneuploidies or male genetic material and these cases were included in our sample. Therefore, from the 401 products sent for genetic examination, we were able to obtain information on

the chromosomal complement in 333 (83.0%), from which 72.7% (242/333) presented a normal karyotype and 27.3% (91/333) an abnormal karyotype.

Considering the cases with a normal karyotype, numerical abnormalities were found in 92.3% (84/91). Trisomy was the most common chromosomal abnormality (55/84; 65.5%). Trisomy 16 ($n = 17$; 30.9%) was the most frequent abnormality, followed by trisomy 18 ($n = 12$; 21.8%) and trisomy 21 ($n = 8$; 14.5%). Twenty-one cases of polyploidy were also identified (25.0%) and seven cases with X-chromosome monosomy (8.3%) (Fig.1). In addition, seven structural abnormalities were found and are described in Table 1. The inherited abnormality was found to have a maternal origin in two of these cases and the remaining were *de novo* abnormalities.

We found that 78.0% (71/91) of chromosomal abnormalities occurred in the first trimester, 19.8% (18/91) in the second and 2.2% (2/91) in the third. A positive correlation was found between foetal maturation and the

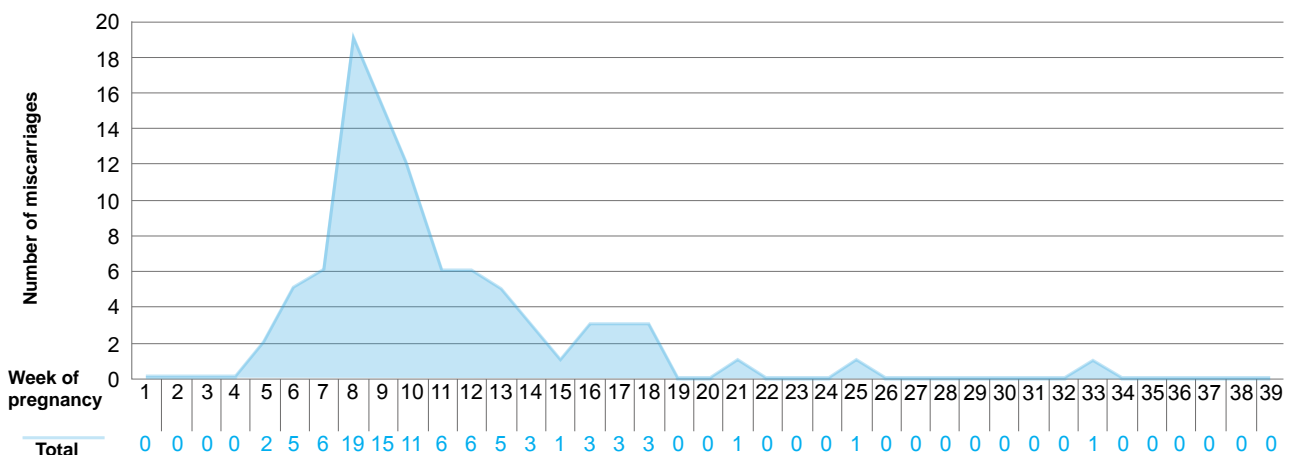


Figure 2 - Distribution of the total number of spontaneous miscarriages or foetal losses per week of pregnancy

presence of chromosomal abnormalities ($p < 0.001$) and half of these occurred (45/91) between the 8th and the 10th week of pregnancy (Fig. 2).

From the 333 cases in our study, 71 were recurrent and 262 were from sporadic miscarriage products. We did not find any significant difference in chromosomal 1st trimester abnormality prevalence between these two groups (36.2% vs. 33.8%, respectively; $p > 0.05$). Nevertheless, maternal age was 32.4 in the group of sporadic miscarriage and 34.8 in the group of recurrent miscarriage ($p = 0.002$).

Maternal age was on average 32.9 (standard deviation: 5.8; 99% Confidence Interval (CI) [32.1 – 33.8]); 185 (55.6%) mothers were aged below 35, 106 (31.8%) between 35 and 39 and 42 (12.6%) above 40 years of age. The younger age group presented 22.2% (41/185) of chromosomal abnormalities, 31.1% (33/106) in the 35-39 age group and 40.5% (17/42) in the group above 40. A positive correlation was identified between maternal age increase and the presence of an abnormal karyotype ($p = 0.02$), with a mean increase of 7.4% (95% CI: [2.6 – 12.4]; $p = 0.002$) in the risk of occurrence of these abnormalities for every additional year of age in the mother. Nevertheless, when considering the diverse chromosomal abnormalities detected, this association was only found in the presence of trisomy ($p < 0.01$).

There was no association between maternal age increase and a higher risk of miscarriage in the presence of polyploidy ($p = 0.939$) or X-monosomy ($p = 0.568$). Taking into account the most common trisomies, we found a higher mean maternal age in the presence of trisomy 21 (38.3; $p < 0.005$) than in the presence of trisomy 16 (33.1; $p < 0.005$) or 18 (34.3; $p < 0.005$).

Considering the results, we found that 219/333 (65.8%) were female and 114/333 (34.2%) were male. Taking into account the gestational age, this female predominance was

only found in the first trimester: 163 (79.1%) were female and 43 (20.9%) were male. In the second trimester, female/male ratio was 41.0% vs. 59.0% and in the third trimester it was 59.1% vs. 40.9%.

We did not find any differences on chromosomal abnormality prevalence between male and female foetus (28.9% vs. 26.5%; $p = 0.19$) except when the results were stratified per trimester: a higher number of abnormal karyotypes were found in male cases in the first trimester of pregnancy (55.8% vs. 28.8%; $p = 0.01$). The comparison between the groups of sporadic and recurrent miscarriage regarding the relation between male and female foetus is presented in Table 2.

The histopathological examination of miscarriage products suggested the presence of a chromosomal abnormality in 89/333 (26.7%) cases. From these, karyotype chromosomal abnormalities were only found in 39 (43.8%) cases, with no agreement between the results of both examinations ($p = 0.608$). We also found 74 cases with signs suggesting an abnormal foetal implantation, 42 cases of infection, 29 cases of disorganized growth and 15 of foetal growth restriction. The group that included a lower proportion of chromosomal abnormalities (4/42; 9.5%) was the one where miscarriage was possibly due to an infectious cause.

DISCUSSION

The identification of the cause in a pregnancy loss is challenging. Aneuploidies must be examined, as they are usually a cause of sporadic miscarriage, namely in the first trimester.^{21,23} Although most chromosomal abnormalities are sporadic and therefore stand a small chance of re-emergence, their identification is reassuring for the couple and has a prognostic value regarding the success of the next pregnancy. Several studies suggested a better

Table 2 - Distribution by trimester of foetal gender and corresponding chromosomal abnormality frequency in pregnancy losses

Karyotype	General						
	1 st Trimester		2 nd Trimester		3 rd Trimester		
	Male	Female	Male	Female	Male	Female	
Sporadic Miscarriage	Normal	13	85	47	30	8	11
	Abnormal	17	33	7	9	0	2
Recurrent Miscarriage	Normal	6	31	6	4	1	0
	Abnormal	7	14	2	0	0	0
Total	43	163	62	43	9	13	
Abnormality frequency	55.8%	28.8%	14.5%	20.9%	0%	15.4%	

prognosis in a subsequent pregnancy when a miscarriage is due to aneuploidy than in the presence of a normal karyotype.^{11,23,24} In addition, when there is a structural abnormality as a cause of spontaneous miscarriage, the presence of possible chromosomal abnormalities in the parents must be ruled out, in order to provide the best possible genetic/reproductive counselling.²⁵

In our study, karyotype was not obtained in a high percentage of cases (20.0%). This fact is described in literature as one constraint related to cell-culture technique and foetal karyotype determination, with reported culture cell failure rates up to 40%.²² This high cell-failure rate may be due to factors like prolonged *in utero* retention time, bacterial or fungal contamination during sample collection or even sample contamination with maternal cells.²⁶ Therefore, the higher failure percentage obtained in the third trimester in our group of patients may be due to cell failure related to a long time interval elapsed between foetal loss and the reception of material for cytogenetic analysis.

In this regard, currently there are new techniques allowing for alternative testing. In the absence of a karyotype determination due to culture failure or due to external contamination (conventional technique), other techniques may be requested from the Genetics Laboratory, such as array Comparative Genomic Hybridization (aCGH) and Quantitative Fluorescence PCR (QF-PCR) techniques. As long as a DNA sample is kept for subsequent studies, these may provide information regarding the chromosome complement of the foetal product.

Recently, Morgen *et al.*²⁷ suggested a diagnostic protocol which has proved to be effective. These authors proposed that the most suitable study protocol of foetal products related with miscarriages should comprise an initial QF-PCR and, in the case of a normal result, an aCGH technique should follow, in order to overcome the two major constraints related with cell-culture i.e. culture failure and maternal contamination. This protocol worked efficiently, allowing for the detection of a higher number of cases with a chromosomal abnormality and the recognition of most maternal contamination situations.

According with expectations, numerical errors were 92.3% of the identified abnormalities, with trisomies being the most prevalent (65.5%). The higher frequency of trisomy 16, followed by 18 and 21 is in line with the evidence available from the literature.^{24,28} We found a prevalence increase of these abnormalities with maternal age increase, with a mean 7.4% annual increment of the risk of occurrence. Taking into account its relevant contribution for spontaneous miscarriage, progressive maternity delay observed in Portugal as in the rest of the world has become an increasing concern. Regarding polyploidies and monosomies, also frequent causes of miscarriage, we did not find any correlation with maternal age, as found by Rolnik *et al.*²⁸

Regarding structural abnormalities, due to their high recurrence rate, is it extremely important to provide the couple with adequate information and genetic counselling

for future pregnancies. Therefore, karyotype of both parents must be obtained, in order to identify the hereditary or *de novo* nature of the abnormality found in the pregnancy loss.

The incidence of chromosomal abnormalities in the first trimester was lower than expected (50-60%)^{2,5,6} in sporadic (33.8%) when compared to recurrent miscarriage (36.2%). The presence of maternal contamination may have contributed to these values, as demonstrated by a female foetal predominance with a normal karyotype in the whole sample (161/242; 66.5%). This result is in line with what has been described: approximately 30 to 40% of 46,XX karyotype obtained by a conventional technique may have a maternal origin.^{29,30}

Therefore, the presence of a higher number of chromosomal abnormalities in male foetus may be related to excessive maternal contamination in our study, which may have masked the results. This result underlines the importance of carrying out a study to rule out maternal contamination every time early miscarriage products are sent for cytogenetic examination.

We did not identify any differences in the prevalence of chromosomal abnormalities from the first trimester, between sporadic and recurrent miscarriage groups. These results are in line with some previous studies^{9,10} and contradict others showing a lower abnormality rate in the case of recurrent miscarriage.^{11,12}

A reduction in chromosomal abnormality incidence was evident with pregnancy progression: 78.0% occurred in the first trimester, 19.8% in the second and only 2.2% in the third. Therefore, these abnormalities have shown to be inversely proportional to pregnancy age, with a lower odd's ratio as the end of pregnancy approaches.^{1,3,31,32} This could be interpreted as a process of early natural selection and other possible causes must be considered for miscarriages occurring after 14 weeks of pregnancy.

Kano *et al.*¹⁹ raise the possibility that, in recurrent miscarriage without any chromosomal abnormality or other known aetiology, foetal loss is predominantly female. However, in this study, the higher proportion of female foetus with a normal karyotype found in the first trimester, in sporadic as well as in recurrent miscarriage, may once more be explained by the presence of maternal contamination in the studied samples. The typical immaturity of miscarriage products in the first months of pregnancy may make it difficult to ensure the origin of the studied material.

With the histopathological examination, we found a suggestion of a chromosomal abnormality in 89 (26.7%) cases although only 39 (43.8%) of these were in line with the karyotype result, i.e. in the presence of a chromosomal abnormality. We found that histopathological examinations are poorly sensitive for the identification of chromosomal abnormalities and therefore its result should not be used as the rationale for diagnosis. Regarding the remaining causes of miscarriage suggested by histopathological examination, we found, as expected, a lower percentage of chromosomal abnormalities in the group where miscarriage was probably caused by infection (9.5%). As the specific characteristics

of inflammatory cells are easily identifiable, this probably explains a lower diagnostic error found in this group. On the contrary, the diagnostic precision in the remaining groups may be affected by the subjectivity of the criteria that we used. To the best of our knowledge, this is actually the first study comparing the results obtained with cytogenetic vs. histopathological examination.

We consider that the size of our sample (333 cases of spontaneous miscarriage or foetal loss) has shown to be a strong point, ensuring the reliability of the results obtained. We should also emphasize that, in the 12 cases in which we only obtained information regarding chromosome complement by MLPA, it is not possible to exclude other chromosomal abnormalities. Although this examination has the advantage of requiring only a small DNA sample and constitutes a rapid and inexpensive test for 13, 18, 21, X and Y chromosomal aneuploidy detection,²² it does not allow for analysis of other karyotype abnormalities, namely chromosome 16 trisomy, commonly involved as cause of spontaneous miscarriage. Even so, the use of the MLPA technique demonstrated its utility by confirming most of the results obtained by karyotype analysis, as well as allowing for the exclusion of some of the most common chromosomal abnormalities in the cases of culture failure.

The inclusion of sporadic and recurrent miscarriage products in the whole sample was very useful as it allowed for the simultaneous study of chromosomal abnormality prevalence in both situations.

CONCLUSIONS

Pregnancy loss, regardless of the time when it occurs, is an event with a very important impact on the couple attempting pregnancy. Apart from a possible prognostic value for a subsequent pregnancy, karyotype analysis may bring important psychological relief to the couple and avoid other investigations.²⁸ Even in the case of sporadic miscarriage, chromosomal abnormalities are commonly implicated most often until the 14th week of pregnancy, with a similar prevalence to the one observed in the case of recurrence. In the specific case of identification of a structural abnormality, it may have its origin in one of the parents and therefore it is crucial to provide all the information and genetic counselling to the couple. Karyotype analysis of every miscarriage product, recurrent or not, will give comfort to the couple and will help them to decide as regards a new pregnancy attempt.

We are aware that an advanced maternal age is also one of the major factors in pregnancy failure. Therefore,

with the current maternity delay and aneuploidy-related miscarriage probability increase, it is necessary for the doctor to previously inform patients of this progressive risk with advancing age.

We suggest a combination of cytogenetic analysis techniques allowing for the study of the whole karyotype or, at least, of the most frequently involved aneuploidies as a cause of spontaneous miscarriage, in order to obtain an efficient study of the chromosomal abnormalities, and in this way avoiding cell culture high failure rate.

Taking into account the high maternal contamination rate involved in this kind of study, maternal DNA additional analysis becomes crucial. In this way it will be possible to ensure the total exclusion of maternal cells from the sample, allowing for the precise prevalence of chromosomal abnormalities to be known and for the study of a possible difference between female and male rate of foetal loss.

We wish to emphasize that this study was based on retrospective data analysis and that, in order to solve the constraints of MLPA technique, the Genetics Department included QF-PCR technique in the study of first trimester miscarriages. Therefore, it was possible not only to exclude maternal contamination, but also to additionally study the number of copies for 15, 16 and 22 chromosomes. As in the latter method, the aCGH technique does not require a cell culture, and has the additional advantage of allowing for the total chromosomal complement analysis, with a much higher resolution than when conventional karyotype analysis is performed.

Therefore, the adoption of a new diagnostic protocol for the genetic study of pregnancy losses including QF-PCR and aCGH techniques to replace the conventional technique (karyotype) seems to be the most adequate methodology and in addition to perform with a more efficient cost-benefit ratio.

CONFLICTS OF INTEREST

The authors declare that there were no conflicts of interest in writing this manuscript.

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