Chromosomal Abnormalities in 238 Couples with Recurrent Miscarriages in Morocco

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Abstract

Purpose: A proportion of cases with repeated abortion are caused by chromosomal abnormality in one of the parents. The purpose of this study was to assess the frequency and nature of chromosomal aberrations that contribute to the occurrence of recurrent miscarriages. Several studies have been done to determine the role of chromosomal abnormalities in couples with recurrent spontaneous abortion in various countries. None of these studies was done in Morocco. Material and Methods: Cytogenetic study was done for 238 Moroccan couples who presented with repeated abortion at the Institut Pasteur, Casablanca, Morocco. Results: We found that the frequency of chromosomal abnormalities was not significantly different from that reported worldwide. Chromosomal abnormalities were detected in 13 (6.1%) of 238 couples. Twelve of chromosomal abnormalities were structural and one of them were numerical. Conclusion: This study highlights the importance of constitutional cytogenetic exploration of couples with a history of repeated spontaneous abortion. Cytogenetic findings could provide valuable information for genetic counseling and allow monitoring of future pregnancies by prenatal diagnosis in couples with a history of recurrent miscarriage.

Keywords
Cytogenetics, Recurrent Abortions, Chromosomal Aberrations

1. Introduction

Miscarriage, defined as spontaneous pregnancy loss at <20 - 28 weeks gestation,
is a common clinical problem. Early pregnancy loss in the first trimester is the most common complication, approximately 15% - 20% of clinically recognizable pregnancies end in spontaneous abortion [1] [2].

Recurrent miscarriage, defined as three or more consecutive miscarriages, affects up to 3% of couples trying to conceive [3] [4]. Some investigators feel that even two spontaneous abortions constitute recurrent miscarriage and deserve evaluation [5].

In almost 50% of cases the etiology is unknown. The causes of RM are heterogeneous and include endocrine dysfunction, autoimmune disorders, genetic abnormalities, maternal and paternal age, infectious diseases, environmental toxins and congenital or structural uterine anomalies etc. [6].

Repeated pregnancy losses during the first trimester are usually due to fetal genetic defects. Pregnancies lost in late gestations also have a high rate of chromosomal anomalies, about 30% in the second trimester and 5% in the third trimester [7].

The incidence of chromosomal abnormalities in those abortions is as high as 50% [8]. Chromosomal abnormalities, mainly balanced rearrangements, are common in couples with reproductive disorders including recurrent abortions [9]. In couples with two or more miscarriages the incidence of these abnormalities varies between 3% and 6% [10] [11]; when one parent carries a Chromosome rearrangement the chance of miscarriage is usually 25% - 50% [12]. This results from the production of gametes and embryos with unbalanced chromosome sets [13] [14].

The clinical consequences of such abnormal gametes include sterility, repeated abortions, and giving birth to malformed children [15] [16].

Several studies have been done in various countries to determine the contribution of chromosome abnormalities in parents with fetal wastage [17].

Cytogenetic studies have been reported to determine the contribution of chromosomal abnormalities in parents with reproductive failure from various other countries.

To our knowledge, no such studies have been done with sample size of 476 cases in Morocco.

The aim of this study was to assess the frequency and nature of chromosomal aberrations that contribute to the occurrence of repeated abortions in Morocco.

2. Materials and Methods

A retrospective study was done in couples with RM from the period between January 2010 to May 2015. This study included 238 (476 cases both partner) Moroccan couples with repeated abortions who were referred for cytogenetic studies at Institut Pasteur, Casablanca, Morocco. All cases were ascertained to have had two or more spontaneous abortions. The obstetric history of couples was either recorded on the request form or retrieved from the files of patients.
All of them had diagnosis more than one spontaneous abortion as we mentioned before. Standard method of 72 hours cultivation of peripheral blood lymphocytes has been applied. Cytogenetic analysis has been performed according to instructions and rules given by International System of Human Chromosomal Nomenclature (ISCN). At least 20 GTG (Gbanding using Trypsin and Giemsa) mitotic cells have been analyzed, if there was doubt of mosaic karyotype, 50 up to 100 cells have been analyzed. C-band has been applied for confirmation of chromosomal heteromorphy.

3. Result

A total of 238 couples (476 subjects) with history of recurrent abortion were examined. Their ages ranged from 19 to 60 years, with a mean of 34.22 years. The age range of women was 19 to 46 years, with a mean of 30.14 years and the age range of men was 19 - 60 years, with mean of 38.31 years.

The number of previous abortion varied from 2 to 12 abortions/couple with a mean of 3.69.

In this study, the 238 couples (476 individuals) studied were classified according to the number of previous miscarriages. In group 1, couples had two miscarriages, in group 2, they had three, in group 3 they had four and more miscarriages. The highest number of patients was seen in group 3 (42.85%) (Table 1).

Among these 238 couples, 13 (5.46%) were found to be carriers of different chromosomal abnormalities, nine females (69.23%) and 4 males (30.77%) (Table 2). Twelve of chromosomal abnormalities were structural and one of them were numerical. These abnormalities included eight balanced reciprocal translocations, one Robertsonian translocation, and four pericentric inversion of chromosome 9 (inv [9] [p11q13]) (Table 2).

Chromosome abnormalities were found in (3/57) = 5.26% of the couples with

<table>
<thead>
<tr>
<th>Table 1. Couples grouped according to the number of miscarriages.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of abortion</td>
</tr>
<tr>
<td>---------------------</td>
</tr>
<tr>
<td>Group 1</td>
</tr>
<tr>
<td>Group 2</td>
</tr>
<tr>
<td>Group 3 ≥4</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Table 2. Sex distribution of abnormal karyotype.</th>
</tr>
</thead>
<tbody>
<tr>
<td>sex</td>
</tr>
<tr>
<td>--------------</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>male</td>
</tr>
<tr>
<td>female</td>
</tr>
<tr>
<td>total</td>
</tr>
</tbody>
</table>

Rcp: reciprocal translocation, Rob: Robertsonian translocation, Inv: inversion.
a history of two abortions, in (4/79) = 5.06% with three abortions, and in (6/102) = 5.88% with four or more abortions (Table 1). Among cases with abnormal karyotype, the mean maternal age was 29.8 and the mean number of abortions was 3.61 per couple. In this study, in couples who reported chromosomal abnormalities only one partner in each couple was affected and the affected individuals showed only one anomaly (Table 3).

4. Discussion

Recurrent spontaneous abortion (RSA), is a multifactorial disorder that challenges both patients and clinicians technically and emotionally [18]. The American Society for Reproductive Medicine (ASRM) recommends that clinical evaluation for RSA may proceed after two first trimester pregnancy losses, which increases the prevalence of this problem to 5% of all couples attempting to conceive [19].

Since cytogenetic finding is considered as a "gold standard" it may give valuable clues [4] for the medical evaluation of patients presenting with recurrent miscarriages.

Several studies have been carried out to determine the prevalence of chromosomal aberrations among couples with recurrent abortion. Schmidt (1962) was the first to report the results of cytogenetics analysis in patients with a history of two or more spontaneous abortions. This was followed by a series of cytogenetic studies of couples with a history of repeated pregnancy loss [10].

The incidence of chromosomal abnormalities among those cases varied in different studies, from none [20] to as high as 21.4% [21].

The variations in the size of the sample, the criteria used for ascertainment of

<table>
<thead>
<tr>
<th>Type</th>
<th>Karyotype</th>
<th>Number Of Miscarriage</th>
<th>Age</th>
<th>Sex</th>
</tr>
</thead>
<tbody>
<tr>
<td>Rec</td>
<td>46XX, t (5, 10) (p15; q26)</td>
<td>7</td>
<td>30</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XY, t (13, 19) (p33; q11)</td>
<td>3</td>
<td>44</td>
<td>M</td>
</tr>
<tr>
<td></td>
<td>46XX, t (1, 7) (p16; q19)</td>
<td>5</td>
<td>31</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XY, t (8, 8) (p22; q23)</td>
<td>5</td>
<td>39</td>
<td>M</td>
</tr>
<tr>
<td></td>
<td>46XX, t (14; 15) (p11q23)</td>
<td>4</td>
<td>25</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XY, t (2, 8) (p22; q22)</td>
<td>2</td>
<td>38</td>
<td>M</td>
</tr>
<tr>
<td></td>
<td>46XX, t (2, 11) (p14q13)</td>
<td>4</td>
<td>26</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XX, t (6, 11) (p21; q21)</td>
<td>2</td>
<td>29</td>
<td>F</td>
</tr>
<tr>
<td>Rob 1</td>
<td>45XY, rob (13, 14) (p10q10)</td>
<td>3</td>
<td>37</td>
<td>M</td>
</tr>
<tr>
<td></td>
<td>46XX, inv (9) (p11; q13)</td>
<td>3</td>
<td>35</td>
<td>F</td>
</tr>
<tr>
<td>Inv</td>
<td>46XX, inv (9) (p11; q13)</td>
<td>2</td>
<td>38</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XX, inv (9) (p11; q13)</td>
<td>4</td>
<td>28</td>
<td>F</td>
</tr>
<tr>
<td></td>
<td>46XX, inv (9) (p11; q13)</td>
<td>3</td>
<td>27</td>
<td>F</td>
</tr>
</tbody>
</table>
cases, and the technique of cytogenetic study have contributed to these wide
differences between various studies [22]. It is also possible that different popul-
ations vary in the incidence of carriers of chromosomal aberrations.

In our study, we found that the incidence of chromosomal abnormalities
among couples with repeated abortions was 5.46%, which is not significantly
different from the global incidence.

We found that nine women and four men had chromosomal abnormalities,
which was a ratio of 2.25. A similar male to female ratio has been found in most
of the reported studies. This predominance of females appears to be due to the
fact that chromosomal abnormalities are compatible with fertility in females may
be associated with sterility in males [15] [10].

The mean maternal age of women carrying chromosomal anomalies was 29.8
years. There was no positive correlation of advanced maternal age with the
number of abortions observed in these subjects indicating that the chromosomal
abnormalities could arise because of some reasons other than advanced maternal
age.

As reported in the literature [10], reciprocal translocations are the most fre-
quent balanced chromosomal anomalies that are detected in couples with recur-
rent miscarriage in our study.

Indeed, in our study, reciprocal translocations were detected in 3.36% of
couples, followed by inversions (1.68%). Robertsonian translocations were ob-
served only in 0.42% of couples. The strong prevalence of reciprocal transloca-
tions compared with Robertsonian translocations would be a result of the dif-
ference in the segregation modes of these anomalies.

In the present study, there was a frequent occurrence of inversion 9. Pericen-
tric inversion of chromosome 9 (inv [9] [p11q13]) is a frequently seen chromo-
somal alteration in humans due to its structural organization, making it more
prone to breakage. The incidence estimated is 1% - 3% of the general population
with the lowest among Asians around 0.25% [23]. Various reports on its associa-
tion with infertility, recurrent miscarriages, hydatidiform moles, azoospermia,
congenital anomalies, growth retardation, and rarely abnormal phenotype have
been published [24]. The largest study on inv [9] and have found a higher fre-
quency among females than in males, especially among those who suffer from
infertility [25].

This indicates the possibility of inversion having a role in the aetiology of re-
current miscarriage, which may be confirmed by molecular studies. An in-
creased tendency to early miscarriages in familial pericentric inversions has been
well documented [26].

Table 2 shows that the distribution of structural chromosomal rearrange-
ments in our study is similar to that reported worldwide Table 4 [27].

Numerical chromosomal aberrations are less frequently encountered among
couples with repeated abortions. Those aberrations are usually in the form of sex
chromosomal aneuploidy, and they occur in a low frequency (<0.15% of cases) [17].
Table 4. Worldwide studies of chromosomal rearrangements observed in couples with recurrent miscarriages.

<table>
<thead>
<tr>
<th></th>
<th>No. of studied couples</th>
<th>Structural aberration</th>
<th>Total (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Rob</td>
<td>Rcp</td>
</tr>
<tr>
<td>Belgium (Ghent)</td>
<td>96</td>
<td>2</td>
<td>6</td>
</tr>
<tr>
<td></td>
<td></td>
<td>8 (8.3%)</td>
<td></td>
</tr>
<tr>
<td>France (Paris)</td>
<td>315</td>
<td>5</td>
<td>7</td>
</tr>
<tr>
<td></td>
<td></td>
<td>16 (5.1%)</td>
<td></td>
</tr>
<tr>
<td>Italy (Padua)</td>
<td>145</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td></td>
<td></td>
<td>14 (9.6%)</td>
<td></td>
</tr>
<tr>
<td>Japan</td>
<td>639</td>
<td>9</td>
<td>19</td>
</tr>
<tr>
<td></td>
<td></td>
<td>29 (4.5%)</td>
<td></td>
</tr>
<tr>
<td>Netherland (Leiden)</td>
<td>67</td>
<td>3</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td>9 (13.4%)</td>
<td></td>
</tr>
<tr>
<td>Netherland (Rotterdam)</td>
<td>148</td>
<td>3</td>
<td>6</td>
</tr>
<tr>
<td></td>
<td></td>
<td>14 (9.6%)</td>
<td></td>
</tr>
<tr>
<td>Saudi Arabia (Riyadh)</td>
<td>193</td>
<td>1</td>
<td>10</td>
</tr>
<tr>
<td></td>
<td></td>
<td>13 (6.7%)</td>
<td></td>
</tr>
<tr>
<td>Oman</td>
<td>380</td>
<td>3</td>
<td>18</td>
</tr>
<tr>
<td></td>
<td></td>
<td>28 (5.5%)</td>
<td></td>
</tr>
</tbody>
</table>

This study has shown that the incidence and distribution of chromosomal abnormalities among Moroccan couples with repeated fetal loss is comparable to that reported worldwide.

This should assist physicians in Morocco by increasing their awareness of the frequency of cytogenetic abnormalities in cases with repeated abortions.

5. Conclusions

Cytogenetic analysis should be part of the investigation of any couple who have experienced at least two pregnancy losses of unknown origin.

Genetic counseling is essential in the management of couples who have had multiple pregnancy failures. A chromosomal anomaly finding in one of the two parents makes it possible to evaluate the prognosis of future pregnancies because the finding of translocation (reciprocal or Robertsonian) or an inversion in either parent is a strong indication for prenatal diagnosis (amniocentesis or chorionic villus biopsy) in making a precise reproductive decision regarding subsequent pregnancies.

Conflict of Interest

No conflict of interest was declared by the authors.

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