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RESEARCH ARTICLE

POLYMORPHIC VARIATIONS OF Y CHROMOSOMES IN RECURRENT MISCARRIAGES

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ABSTRACT

Three or more consecutive miscarriages before 20 weeks of gestation is considered as recurrent miscarriage (RM). It is a challenge for obstetricians to handle these sensitive issues which involves not only wife but also the whole family. Though recurrent miscarriage is multifactorial genetic contribution plays important role in causing recurrent miscarriage. Numerical and structural aberrations in chromosomes become the causative factor in recurrent miscarriage. Polymorphic variations are seen to be associated with RM. To find out whether Y chromosomal polymorphic variations play role in recurrent miscarriages 339 couples (678 patients) of history recurrent miscarriages were analyzed by karyotyping. Out of 339 patients in 33 (8.2%) patients Y chromosomal polymorphic variations were noticed. Amongst 33 patients in 6 (1.7%) cases Yqh+, in 21 (6.7%) Yqh-, in 5 cases inversion Y (1.2%) was found.

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INTRODUCTION

Percy Malpas from Liverpool coined the term Habitual abortor for women with three or more consecutive abortions¹. Documented data reveals 15 to 20% of all pregnancies result in recurrent miscarriages. Clinician and geneticist point of view miscarriage is a cause of concern. Not only wife suffers but also whole family undergoes social and psychological trauma. Along with anatomical, hormonal, immunological factors, genetic factors also contribute in miscarriages. 6% of chromosomal abnormalities cause recurrent miscarriages. Parental chromosomal aberrations have proved to be the reason for recurrent miscarriages.

Not only chromosomal aberrations like inversion, deletion and translocation but also heteromorphisms in chromosomes are observed in recurrent miscarriages. Chromosomes like 1, 9 and 16 show heterochromatic variations² and they were observed to be associated with infertility³. Many studies indicate that chromosomal polymorphism may cause certain clinical effect such as infertility and spontaneous abortions^{4, 5}. Polymorphic variations were considered as normal for long time⁶. It was observed that heterochromatization has no effect on abortions and they are considered as normal variations^{6, 7}. Heterochromatin is formed by tandemly organized highly repeated sequences of satellite DNA that do not encode proteins. So chromosomal polymorphism are considered as normal Karyotypes⁸. Wide documentation is available regarding association of variants of

chromosomes in cases of recurrent abortions. Many chromosomal aberrations are involved in case of RM.

In human, Y chromosome has euchromatic region which is made up of short arm, long arm and centromere while heterochromatic region is made up of long arm Yq which varies in size. Y chromosome harbors 104 genes (21 genes on short arm, 40 on long arm, 4 on terminal region, and 39 gene position not identified). SRY gene is located on the short arm and three spermatogenesis loci are mapped on long arm known as azoospermia factors (AZFa, b, c). Out of three factors AZFc deletion is the most common cause of spermatogenic failure. It is well known that Y chromosome plays important role in sex determination and spermatogenesis¹. However its role in RM is conflicting and not proven completely. Also, whether polymorphic variations of Y chromosome are causative factors for RM is not yet confirmed. So, we attempted to focus on incidence of Y chromosomal polymorphic variations in RM. The present study was undertaken by keeping the objective in the mind of investigating the incidence of heterochromatic variations in Y chromosome in RM.

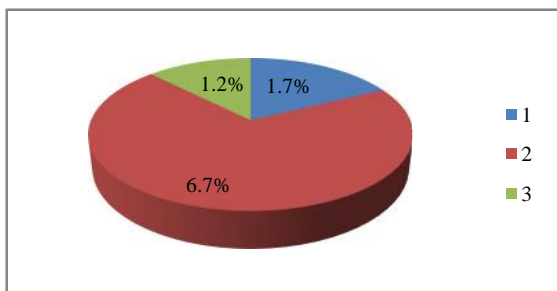
MATERIALS AND METHODS

OBSERVATIONS AND RESULTS

1. Out of 339 couples (678 patients) in 33 (8.2%) patients Y chromosomal polymorphic variations were noticed.
2. Yqh+ was observed in 6 (1.7%) cases.

3. Yqh- was seen in 21 (6.7%) cases.
4. Inversion Y was seen in 5 (1.2%) cases.

Following chart shows percentage of Y chromosomal polymorphic variations



Following diagram shows polymorphic variations of Y chromosome in present study.



First images shows Yqh+, 2nd image shows Yqh- and 3rd image shows inversion Y chromosome

DISCUSSION

Polymorphic variations and chromosomal heteromorphism are synonymus. Normal variant, heteromorphism and polymorphism are often used interchangeably. All three usually refer to variations in human karyotype or genome that are heritable. Heteromorphism show increased heterochromatin at chromosomal telomere, short arm of chromosome and the centromere^{6, 9}. Heterochromatin is defined as microscopically visible chromosome region that is variable in size, morphology and staining properties in different individuals. It is just one form of normal variation in human genome⁹. Heterochromatin located in centromere has an essential role in spindle attachment and chromosome movement, meiotic pairing¹⁰. When a chromatin variation occurs in these regions, it causes defects in centromere function and kinetochore assembly. This causes difficulty in homologous chromosome pairing and impacts on cell division. This could affect gamete formation⁶. Though there is a debate on the clinical significance of polymorphism of various chromosomes, cytogeneticist should not ignore their presence, occurrence and incidence. Such variants play important role in reproductive failure¹¹.

Polymorphic variants usually occur in long arms of 1, 9, 16 and distal 2/3rd of long arm of Y chromosome^{6, 11}. Polymorphisms were more associated with chromosome 9 and Y¹².

Wide documentation of various forms of polymorphic variations of Y chromosome like long Y (Yqh+), small Y (Yqh-) and inversion Y (inv Y) reveals their incidence in RM. Yqh+ is the polymorphic variation of Y chromosome where the heterochromatin region of long arm is increased in the length. Too many DNA repeats at specific regions of Y chromosome may impact on the pairing and synapsis of X& Y chromosomes

during meiosis. Increase in the long arm of Y chromosome may inhibit the expression of genes of spermatogenesis by position effect variegation and decrease reproductive capacity³. This was supported by putting theory that Yqh+ heterochromatin may be related to clinical abnormalities, because large Y chromosome heterochromatin represents excessive duplication of DNA that may in turn be related to errors in mitosis, gene regulation or cell differentiation, ultimately leading to spontaneous abortions¹³.

Heterochromatic region of Y chromosome have been postulated to play a role in immune response during early embryo development. It is also suggested that Yqh+ may affect the viability of zygote but it does not affect fertility of its carrier¹. Increased length of Y chromosome may be an important cause of fetal loss¹⁴. Several studies of families with two or more abortions have also shown a suggestive increase in frequency of chromosome variant such as a long Y in father or a large satellite in acrocentric chromosomes¹⁵. Increase rate of spontaneous abortion associated with Yqh+⁹. A highly statistically significant increase in the frequency of Yqh+ was observed in men whose wives had a BOH³.

On the other side many authors deny that Yqh+ plays role in RM. Rather they say that Yqh+ in father may be unrelated to fetal loss¹⁶, Y chromosome length variability is unassociated with reproductive problems¹⁷, Yqh+ do not appear to have any functional or phenotypic effect¹⁸, increased length of Y chromosome is heritable and Yqh+ has not been associated with abnormal phenotype¹⁹.

The hypothesis explained was increased length of Y chromosome might be due to difference in the degree of contraction of chromosomes during cell division²⁰, Yqh+ is the result if an addition of chromosomal substance because it bears two secondary constrictions²¹ and difference in the size of chromosome is a morphological feature without any functional significance²².

Table no 1 Following table shows incidence of Yqh+ in different studies.

Name of Author	Incidence of Yqh+
S. Dubey ²³	0.13%
Abramsson L ²⁴	4.4%
Nagvenkar ²⁵	3.4%
De la fuente Cortes BF ²⁶	15.82%
Christofollini DM ²⁷	1.78%
Hou JW ²⁸	3.6%
Guluzar Aarhus ²⁹	0.42%
Caglayan AO ³⁰	1.82%
Ebru Onalan ³¹	4.6%
Present Study	1.7%

Small or deficient heterochromatic region in long arm of Y chromosome is Yqh-. Increased or decreased length of long arm of chromosome is decided after calculating Y/F index, which is calculated as total length of Y chromosome / average total length of F group chromosomes (19 & 20). Y chromosome belongs to G group but its average total length is equivalent to the average total length of F group chromosomes. Hence the Y/F index is calculated to compare to F group. The average Y/F index is 0.95 to 1.09¹⁶. Long Y (Yqh+) chromosome and short Y (Yqh-) chromosome are known to exit³².

Table no 2 Following table shows incidence of Yqh- in different studies.

Name of author	Yqh-
S Dubey ²³	4.76%
Nagvenkar ²⁵	27.3%
Christofollini DM ²⁷	0.2%
Retief AE ³³	1.6%
Guluzar ²⁹	0.85%
Caglayan AO ³⁰	0.9%
Ebru Onalan ³¹	1.9%
Present Study	6.7%

An inversion is a chromosome rearrangement in which a segment of a chromosome is reversed end to end. It does not involve a loss of genetic information but simply rearranges the linear gene sequence. Inversion can be pericentric where centromere is rotated and can be paracentric where breakage points are in one arm and centromere is not involved in it. Usually inversion does not cause any abnormality as long as rearrangement is balanced, however in heterozygous individuals there is an increased production of abnormal chromatids which leads to lowered fertility due to production of unbalanced gametes.

Pericentric inversion of Y was first documented by Jacobs in 1964³⁶. Association between inversion Y and recurrent miscarriage is noticed by many authors. Its relative high prevalence suggested that inversion was the most common chromosomal alteration associated with repeated pregnancy loss³⁵. We also found 1.2% of cases with pericentric inversion of Y chromosome. It is a rare chromosomal heteromorphism without clinical significance and does not affect sperm production^{16, 36}. Though number of cases with inversion Y in RM is scanty, but its presence in RM can not be ignored.

Table no 3 Following table shows incidence of inversion Y in different studies.

Name of the author	Inversion Y
Usha Dutta ¹²	0.08%
Tomomasa ³⁴	1 case report
Ebru Onalan ³¹	0.5%
Frenny Sheth ³⁵	2.2%
Zhou B ³⁷	22%
Present Study	1.2%

Heteromorphism do exist and show some association in RM. It is a tip of iceberg of which the base may be broad and deep. Combine efforts of geneticist and gynaecologist will be required to search exact cause of RM. Application of advanced molecular techniques will ease the understanding of molecular basis of heteromorphism.

CONCLUSION

1. Chromosomal analysis in recurrent miscarriages is must.
2. Any heteromorphism should not be ignored.

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