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Cytogenetic Analysis and Clinical Phenotype of Primary Amenorrhea in Indonesian Patients

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Article Info History: Received: 19 Apr 2018 Accepted: 06 July 2018 Available: 31 July 2018	 Abstract Background: Primary amenorrhea (PA) is a symptom that can be caused by different disorders such as gonadal, endocrinal, physiological and genetic disorders. Objectives: This study provided the clinical and cytogenetic profiles of Indonesian primary amenorrhea patients and matching the clinical criteria of those patients with their karyotype results by using a new scoring system. Methods: A retrospective descriptive study of 79 PA patients, whom referred to Cytogenetic and Molecular unit Center for Biomedical Research (CEBIOR), Faculty of Medicine Diponegoro University. We made a scoring system consisted of 4 scores, all patients had been distributed to match the scores according to their clinical criteria's and then confirmed with the karyotype results. Results: The karyotype results of 79 patients of PA revealed 55 (69.6%) patients with female karyotype 46,XX; 6 (7.6%) patients with male karyotype 46,XX; 3 (3.8%) patients with monosomy X; 3 (3.8%) patients with 45,X/46,XX; 3 (3.8%) patients with Isochromosome 45 X/46, X,i(Xq). Mosaicism with Y constitution 45,X/46,XY was seen in 2 (2.5%) patients; marker chromosome 45,X/46,X+mar (2%) in 1 patient (1.3%); and chromosome 1 and X translocation 46,XX,t(1;X)(p34;q25) detected in 1(1.3%) patient. Scoring system results showed that all patients with normal karyotype (46,XX/46,XY) were matched with score 1 and 2 while 17 patients with mosaic Turner syndrome 45,X(10%)/46,XX(90%) matched score 1. Conclusion: Turner syndrome was the most common cause of primary amenorrhea, which attests the importance of cytogenetic analysis for diagnosis of
	Conclusion: Turner syndrome was the most common cause of primary

Keywords: primary amenorrhea; karyotype; clinical phenotype; score system

INTRODUCTION

Primary amenorrhea defined as failure of menarche, associated with undeveloped secondary sexual signs by age 14, or failure of menarche with well-developed secondary sexual signs by age 16.¹It is a symptom that caused by different disorders.^{2,3} The worldwide incidence estimated to be 1%,^{4,5} and based on research results from different countries, there was

no evidence for higher frequency in a specific population or ethnic group.³

A number of studies estimated the frequency of primary amenorrhea based on the causes including gonadal dysgenesis was due to chromosomal abnormalities as the largest cause accounting in 45%, followed by Mayer Rokitanski Kuser Hauser * Corresponding author: Sultana MH Faradz E-mail: sultanafaradz@gmail.com

syndrome (MRKH) in 15%, and Hypothalamic Idiopathic HypogonadotropicHypogonadism (IHH) was estimated to be 15%. Complete androgen insensitivity syndrome (CAIS) assessed for 10% of all primary amenorrhea patients,⁷ imperforated hymen and transverse septum hymen reported to be 5%,8 while the remaining 5% was distributed among congenital adrenal hyperplasia (CAH) and ovarian insensitivity syndrome.³Mullerian Agenesis-Mayer Rokitansky-Kuster-Hauser Syndrome (MRKH) has a prevalence of 1:5000 female births.⁶ Affected individuals have a 46,XX karyotyping and a normal secondary sex characteristics. Type A (MRKA) patients show Symmetric uterine buds and fallopian tubes. Type B (MRKH) shows asymmetric uterine buds and fallopian tubes and being associated with other congenital anomalies (skeletal, renal, ovarian, ear and cardiac).^{6,9}

This is the first study in Indonesia to provide the profiles of primary amenorrhea patients including the karyotype results and their clinical profiles. The new of this study that we made a scoring system and the patients had been distributed to match the scores according to their clinical criteria and then confirmed it with their karyotype results. This scoring system can help to distinguish roughly possible karyotype results in situation of deficient genetic facilities before cytogenetic analysis hold out. Scoring system can be used to predict that PA patient with normal secondary sexual signs can be female or male karyotype results and cannot be a chromosomal abnormality based on clinical phenotypes. In contrast, PA with poor secondary sexual development with short stature and web neck can predict of chromosomal abnormalities. However, this scoring system still needs further studies to measure validity and reliability, and whether it could be used as a tool to predict the clinical phenotype and the genetic cause/karyotype.

METHODS

Population and Sample

The sampling method was purposive sampling; patients had been selected according to the purpose of the study. All patients who referred to CEBIOR with primary amenorrhea and had available data, with the criteria: thirteen years old with undeveloped secondary sexual signs or 16 years old with welldeveloped secondary sexual signs. All patients with inclusioncriteria Whom referred to Cytogenetic and Molecular unit Center for Biomedical Research (CEBIOR), Facultyof Medicine Diponegoro University from the period of January 2004 to January 2017 were included in this study.

In this study we made a scoring system consisted of clinical criteria and karyotype results the score system had 4 scores, score 1 for primary amenorrhea symptom only, score 2 for primary amenorrhea and poor secondary sexual signs, score 3 for primary amenorrhea, poor secondary sexual signs and short stature, score 4 for primary amenorrhea, poor secondary sexual signs, short stature, and webbed neck. The clinical data was obtained from the medical records and were grouped into the scoring system. Patients without complete cytogenetic and clinical data were excluded. All the patients had been distributed to match the scores according to their clinical profiles and then confirmed with the karyotype results.

Poor secondary sexual signs considered if Tanner stage less than 4 with spare or absent axillary hair. Short stature considered if the height is below 2 slandered deviation of the mid parental hight.

Chromosomal analysis had been performed by blood as a usual method in CEBIOR.

Table 1. Scoring system for primary amenorrhea patients

PA= primary amenorrhea

Poor secondary sexual signs = Tanner stage less than 4 with spare or abscent axillary hair

Score	Clinical Criteria
1	PA
2	PA, poor secondary sexual development
3	PA, poor secondary sexual development and short stature
4	PA, poor secondary development, short stature, webbed neck.

RESULTS

Seventy nine PA patients were included in this study. The distribution of patients according to the karyotype revealed that 55 (69.6%) of patients had 46, XX karyotype; 6 (7.6%) patients had 46, XY karyotype, and 18 (22.8%) patients with chromosomal abnormalities as showed in Figure 1.

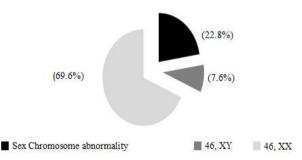


Figure 1. Distribution of karyotype results in primary amenorrhea patients

The most frequent chromosomal abnormalities was 45,X in 10.1% followed by 45,X/46,XX mosaicism in 3.7%. Isochromosomemosaicism of 45,X/46,X,i(Xq) accounted in 3.8%. Mosaicism with Y constitution 45,X/46,XY was found in 2 patients 2.5%; marker chromosome 45,X/46,X +mar (2%) in 1 patient and chromosome 1 and X translocation 46,XX,t(1;X)(p34;q25) detected in 1 patient. The

Karyotype results of all the patients showed in Table 2.

Chromosomal Categories	Karyotype	Number of patients	
Categories		n	%
Female Karyotype	46, XX	55	69.6
Male karyotype	46, XY	6	7.6
Numerical			
Abnormality			
Monosomy	X 45,X	8	10.1
Turners Mosaic	45,X/46,XX	3	3.8
Presence of XY	45,X/46,XY	2	2.5
constitution			
Structural			
Abnormality			
Marker	45,X/46,X +mar2%	1	1.3
chromosome			
Isochromosome	45,X/46,X,i(Xq)	3	3.8
Translocation X;1	46,XX,t(1;X)(p34;q25)	1	1.3

Table 2. The karyotype results of 79 patients with primary amenorrhea.

The distribution of diagnosis among 55 patients with female karyotype 46, XX revealed 14 patients with MRKH, 2 patients with CAH and 1 patient with pure gonadal dysgenesis, while the remaining 38 patients still with unknown diagnosis. The diagnosis among 6 PA patients with male karyotype, 2 were CAIS, 1 was PAIS, and 3 patients with pure gondal dysgenesis (PGD) of undetermined cause.

According to the cytogenetic results of 18 patients with chromosomal abnormalities, the diagnosis were classical Turner Syndrome in 8 patients, mosaic Turner Syndrome in 9 patients and 1 patient with autosomal X translocation. The distribution of diagnosis among all patients showed in Figure 3.

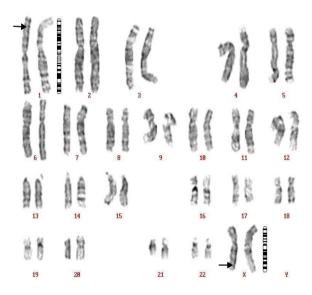


Figure 2. Karyotype result of patient with balanced translocation between 1p (arrow) and Xq (arrow)

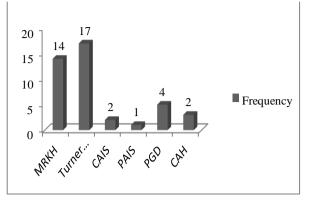


Figure 3. Frequency of diagnosis among 40 patients with primary amenorrhea

The result of scoring system and their karyotype revealed that scored 1 which was presented with primary amenorrhea only were 30 patients with female karyotype (46,XX); 1 patient with male karyotype 46,XY; and 1 patient with mosaic Turner Syndrome 45,X(1%)/46,XX(99%). There were 25 patients with female karyotype (46,XX) and 5 patients with male karyotype (46,XY) matched with score 2. Score 3 was demonstrated in 4 patients with classical Turner Syndrome (45,X); 4 patients with mosaic Turner (45X,46,XX) and 1 patient with karyotype

46,XX,t(1;X)(p34;q25). In This study 4 patients with classical Turner Syndrome (45,X), 3 patients with isochromosme 46,X,i(Xq)/45,X and 1 patient with marker chromosome 45,X(98%) /46,X,+mar (2%) were matched with score 4. (See Table 3)

Table 3. Scoring system and its related karyotype

Score	Frequency of Patients	Karyotype Results
1	30	46,XX
	1	45,X(1%)/46.XX(99%)
	1	46,XY
2	25	46,XX
	5	46,XY
3	4	45,X
	1	45,X(20%)/46,XX(80%)
	1	45,X(2%)/46,XX(98%)
	1	45,X(90%)/46,XY(10%)
	1	45,X(80%)/46,XY(20%)
	1	46,XX,t(1;X)(p34;q25)
4	4	45,X
	1	46,X,i(Xq)(10%)/45X(90%)
	1	46,X,i(Xq)(24%)/45X(76%)
	1	46,X,i(Xq)(20%)/45X(80%)
	1	45,X(98%)/46,X,+mar(2%)

DISCUSSION

Attainment of menarche is important for female confidence and feminism. Primary amenorrhea can be a cause of psychological trauma in any female in reproductive age group.² World Health Organization (WHO) ranked the primary amenorrhea as the sixth most common cause of infertility, hence the amenorrhea account 20% of all patients of infertility.⁶ The cause of primary amenorrhea are different and the role of genetic factors are significant. Several studies

for cytogenetic analysis of primary amenorrhea patients have been done and aiming to understand the frequency and chromosomal constitution in those patients.¹⁰

In this study we demonstrated that 22.8% (18/79) PA patients with chromosomal abnormalities either numerical or structural. However, our studv coordinate to the world wide estimated range for abnormalities primary chromosomal among earlier studies amenorhea patients. In the chromosomal abnormalities among primary amenorrhea patients reported to be counted in 20.63% Egyptian patients,¹¹ 27.8% in Indian in population,^{1,4,12} 20% in Iranian population¹³ and 41% in Mexican population.¹⁴ However, our study results was coordinated to the world wide estimated range for abnormality chromosomal among primarv amenorrhea patients which was between 15.9% and 63.3%.5,7,15

All patients with sex chromosomal abnormalities in our study were Turner Syndrome either classical Turner Syndrome or Mosaic. This result was in line with previous studies which reported that Turner Syndrome is the most common observed chromosomal abnormality in primary amenorrhea and also strengthened the role of sex chromosome in the reproduction of female.^{4,9,16}

The previous result revealed that classical Turner Syndrome detected in 30% (16/52) of primary amenorrhea patients in Turkish population,¹⁶ and 26.9% (7/ 26) of Indian patients.^{1,4,12,17}In current study Turner syndrome was detected in 17 out of 18.The high percentage of Turner Syndrome in our study could be due to patients selection and lack of genetic diagnostic in many referral hospitals in Java Island. So, most of patients referred to our center to do cytogenetic analysis.

The clinical profiles of Turner Syndrome patients showed that all patients with classical Turner Syndrome had short stature. Despite short stature, which seems to be the general clinical characteristic of Turner Syndrome, all other clinical stigmata were inconsistent, even in individuals with classical 45,X. The possible explanation for this issue that the physical manifestations of Turner Syndrome patients largely depends on the karyotype.¹⁶ The cause of short stature in Turner Syndrome suggested to be a homeobox gene, short stature homeobox (SHOX) gene in the pseudoautosomal region 1 (PAR1) is the major player and that haploinsuffiency of this gene leads to the growth failure.^{16,20} The other possible cause of short stature in Turner syndrome is inadequate production of Estrogen, many of those with Turner Syndrome developosteoporosis. This can decrease height further, as well as exacerbate the curvature of the spine.14,20

Mosaic Turner Syndrome with XY constitution seen in 11.7% (2/17) of the patients, detection of Y chromosome and its component is very important due to the risk of gonadoblastoma since the risk is quite high ranging from 10 to 20%. While patients with mosaicism for 46,XY cell line or structural rearrangement of the Y chromosome mostly have masculinized external genitalia and are at increased risk for having gonadoblastoma and other gonadal tumor.^{12,20,21} Thus, early intervention should be done for orchidectomy.²² In this study we observed 1 patient with marker chromosome 45,X(98%)/46,X mar(2%) with clinical profile of this patient showed short stature and no other dysmorphic features, compared to the previous study which reported severe phenotype manifestations in those type of Turner variants,¹⁶ this could be due to the low percentage of marker X. Although, the conventional cytogenetic cannot detect the origin of marker chromosome, the molecular cytogenetic techniques, FISH, can accurately detect it.^{16,17} In this study we observed a patient with translocation between chromosome 1 and X. Besides primary amenorrhea, patient also had short stature. Translocation between X chromosome and autosome are known very often leads to fertility problems including PA and clinical features of Turner Syndrome and variable phenotype including developmental delay, thus, may due to nonrandom Xinactivation.¹⁸ Especially when involving q arm of X chromosome, therefore, the finding may very well presented a balanced translocation resulting in the described phenotype (see figure 2). Some candidate genes located in the q arm of X chromosome may interrupted due to X-autosome translocation such as DIAPH2, XPNEP2, DACH2, POF1B, CHM and NXF5.19 However, further cytogenetic/molecular analysis should be done to see the genotype and phenotype association in this patient

In our study the most common cause of primary amenorrhea in female karyotype patients was MRKH (14/55). The clinical profile of the patients showed normal Tanner stage in all patients, normal external genital and no dysmorphic features. With regards to the high number of MRKH patients, it needs molecular study to detect the associated gene mutations,²³ although, the cause of mullerian gene is unknown, but it may be due to a mutation in the gene of the anti mullerian hormone or the Anti Mullerian (AMH) receptor. The underlying Hormone mechanism would be exposure to AMH acivity. No activating mutation is reported, in contrast with inactivating mutations which cause persistence of mullerian structures.24

Male karyotype presented in a significant percentage (7.6%) of patients with primary amenorrhoea although they appeared to be physically normal with some just appearing the height is taller for their age. In our study Y chromosome had been detected in 10% of PA patients, compared to previous study which demonstrated that Y chromosome was observed in 20% patients that referred as complain of primary amenorrhea.¹⁴ Although, this study demonstrated lower percentage, it could be due to selection of patients, preservation of some patients for karyotype analysis. Furthermore, conventional cytogenetic method can missed the Y component up to 9.3%.²¹ Detection of Y chromosome complement and their composition is important in genetic counseling, because of the association with the risk of gonadoplastoma.9,12,20

In this study the most common cause of primary amenorrhea was gonadal dysgenesis Turner Syndrome followed by MRKH, pure gonadal dysgenesis, CAH, CAIS and PAIS. The remaining 39 (49.3%) patients still not diagnosed, it could be due to mutations and need molecular analysis for establishing the diagnosis. This study was agree with study in United States which demonstarated that gonadal dysgenesis as the commonest cause of primary amenorrhea and MRKH as the second most common cause.²³ The same result had been reported from Korean study, which reported that the common causes of primary amenorrhea were gonadal dysgenesis (28.0%, 37/132), followed by MRKH syndrome (20.0%, 27/132).²⁵ However, MRKH was the most prevalent cause of primary amenorrhea in Thailand, they reported that the 3 most common causes of primary amenorrhea were Müllerian agenesis/MRKH?

(39.7%), gonadal dysgenesis (35.3%), and hypogonadotropichypogonadism (9.2%).^{17,20,26}This verified that racial and environmental factors played an essential part in the causes of primary amenorrhea.

In the present study we made a scoring system from the clinical criteria and we matched the patients to appropriate scores. This scoring system can help to distinguish roughly possible karyotype results in situation of deficient genetic facilities before cytogenetic analysis hold out. Scoring system can be used to predict that PA patient with normal secondary sexual signs can be female or male karyotype results and cannot be a chromosomal abnormality based on clinical phenotypes. In contrast, PA with poor secondary sexual development with short stature and web neck can predict of chromosomal abnormalities. However, this scoring system needs further study with large sample population and measurement for validity and reliability.

A significant number of patients had sex chromosomal abnormalities, thus early cytogenetic investigation is prudent to guide further management. Patients with primary amenorrhea should be initially screened by primary physicians and gynecologists for presumptive diagnosis here in after, patients should receive prompt referral for genetic study. The reason for referral should be explained to the patient. If cytogenetic abnormalities are detected, a detail explanation should be given to the patient by a geneticist or gynecologist with experience in genetics. Counseling session should include the risk of premature ovarian failurefor patients withTurner's syndrome and the use of hormonal replacement therapy, the possibility of infertility in the future patients with mosaic Turner and the risk of gonadal malignancy for patients with XY gonadal dysgenesis. Counseling should be performed tactfully, bearing in mind that sensitive issues related to femininity are involved. An experienced counselor and clinical psychologist if needed would be helpful when patient facing with the gender identity.

CONCLUSION

Turner syndrome is the most common cause of primary amenorrhea in patients with chromosomal

abnormalities. Cytogenetic analysis to distinguish the causes of primary amenorrhea is an important step especially in developing countries, although molecular analysis is now mandatory to detect the gene mutation. The new scoring system needs further study for measuring the reliability and validity.

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