

Clinico- Radiologic and Genetic Study of Fraser Syndrome: A Case Report

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Abstract: Cryptophthalmos syndrome or Fraser syndrome is a rare autosomal recessive disorder, characterized by major features such as cryptophthalmos, cutaneous syndactyly, malformation of the larynx, abnormal genitalia, craniofacial dysmorphism, orofacial clefting, mental retardation and musculoskeletal anomalies. It is inherited in an autosomal recessive pattern where both the parents must be possessing one defective allele for Fraser syndrome. Since both the parents are carriers for the Fraser syndrome, there is a chance for 25% of their children being affected, 50% of their children to be carriers and 25% of their children to be normal. A 22 year old pregnant woman with a family history of third degree consanguineous marriage was diagnosed to have a fetus with many of the above findings in her anomaly scan which was done to manage child birth and as it is crucial to get prenatal diagnosis early in the pregnancy. Parents decided to terminate the pregnancy. Soon after the abortus was expelled, cord blood was collected and subjected to Karyotype test. Karyotype analysis revealed the absence of a short sequence on chromosome number 4 in q arm at 4q21 region, which may be responsible for the above mentioned deformities. Due to the high morbidity and mortality associated with Fraser syndrome, early detection by amniotic fluid karyotyping may be helpful in early intervention.

Keywords: Cryptophthalmos Syndrome, Fraser Syndrome, Syndactyly

1. Introduction

Fraser syndrome is an autosomal recessive congenital disorder [1]. The initial tale of Fraser Syndrome is attributed to Zehender and coworkers in 1872, when it was known as ‘cryptophthalmos syndrome’, as reported by Khoury *et al* [2]. Fraser syndrome is named after the geneticist, George R. Fraser, who first described the syndrome in 1962. Considering this syndrome as a complex disorder with multiple anomalies, with or without cryptophthalmos, Fraser syndrome is the preferred terminology [3]. As per the diagnostic criteria in the opinion of Thomas *et al*. [6], Fraser syndrome is characterized by developmental defects, including cryptophthalmos (where the eyelids fail to separate

in each eye) and malformations in the genitalia (such as micropenis, cryptorchidism or clitoromegaly) [4]. Congenital malformations of the nose, ears, larynx and renal system, as well as mental retardation manifest occasionally. Syndactyly (fused fingers or toes) has also noted [5, 6]. The reported incidence of Fraser Syndrome is 0.043 per 10,000 neonates and 1.1 in 10,000 stillbirths, worldwide [7]. The reported incidence of Fraser Syndrome is 0.043 per 10,000 live born infants and 1.1 in 10,000 stillbirths, worldwide [7].

Cryptophthalmos (hidden eye) refers to a group of rare congenital eyelid deformity, in which the eyelids fail to separate. Patients with a syndromic combination of acrofacial and urogenital malformations with or without cryptophthalmos should be taken into consideration for Fraser syndrome [8].

Subjecting amniotic fluid sample for karyotyping test between 16-20 weeks of gestation can help diagnosis and early intervention in cases of congenital malformations.

The genetic background of this disease has been linked to FRAS1, a gene involved in skin epithelial morphogenesis during early growth [9]. The FRAS1 gene is pin pointed on the long arm of chromosome 4 (at 4q21 region). This disorder has also been coupled among FREM2 gene that originate in chromosome 13 [10] and with mutations in GRIP1 gene of chromosome 12 [11]. FRAS1 gene is responsible for proper development of skin, internal organs and tissues. Protein from the gene GRIP1 is responsible for the proteins from the genes FRAS1 and FREM2 to get to the actual location in the cell to work together [12]. The proper identification of Fraser syndrome can be accurately made after birth based on its distinctive association of malformations. Occasionally, it is possible to establish this disorder due to abnormal prenatal ultrasonographic features such as, polyhydramnios or oligohydramnios, echogenic lungs and renal abnormalities/agenesis [1].

2. Case Report

A 22 year old pregnant woman came with a history of previous spontaneous abortion at 12 weeks of gestation and

wanted to undergo anomaly scan for the present pregnancy. She had obstetric history of gravida 2, para 0, abortion 1, living 0. Family history revealed second degree consanguineous marriage. Her first trimester was uneventful-she had regular antenatal checkups and was supplemented with folic acid, iron and other general nutritional supplementations. Nuchal translucency scan was normal.

Second trimester was uneventful, further she underwent anomaly scan at 24 weeks of gestation. Anomaly scan report at 24 weeks of gestation revealed the following findings suggestive of abnormal thorax, abnormal abdomen, abnormal kidney and urinary bladder, and abnormal extremities in the fetus:

- 1) The lungs were massively enlarged and hyperechogenic, resulting in compression of the heart, and development of ascites.
- 2) The bronchial tree and trachea were dilated and the diaphragm inverted.
- 3) Abdomen was distended with massive ascites.
- 4) Bilateral club foot.
- 5) Chronic high airway obstruction syndrome.
- 6) The bladder was small and cystic dysplasia of the kidneys was noted (Figure 1).
- 7) Cystic hygroma was noted in the fetus.

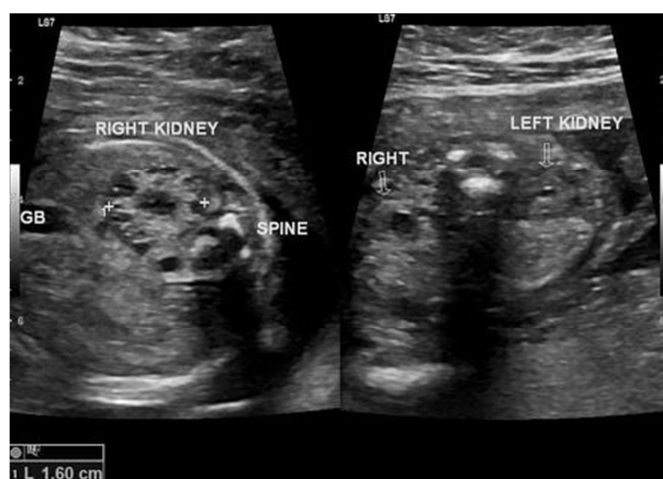


Figure 1. Ultrasonographic image of fetus, showing cystic dysplasia of kidneys.

Ultrasonography also revealed oligohydramnios. Due to the above mentioned anomalies, medical termination of pregnancy was done. On gross examination, the fetus showed grade 3 microtia of ears, saddle depression of dorsum of nose, syndactyly in fingers, cryptorchidism and micropenis and bilateral clubfoot (Figure 2).



Figure 2. Fetus showing club foot, syndactyly, malformed ears and cryptophthalmous.

Soon after termination of pregnancy, cord blood was collected and subjected to Karyotype analysis, using standard Karyotyping protocol. Karyotype of the sample revealed the absence of a short sequence on chromosome number 4 in q arm, at 4q21 region (Figure 3).

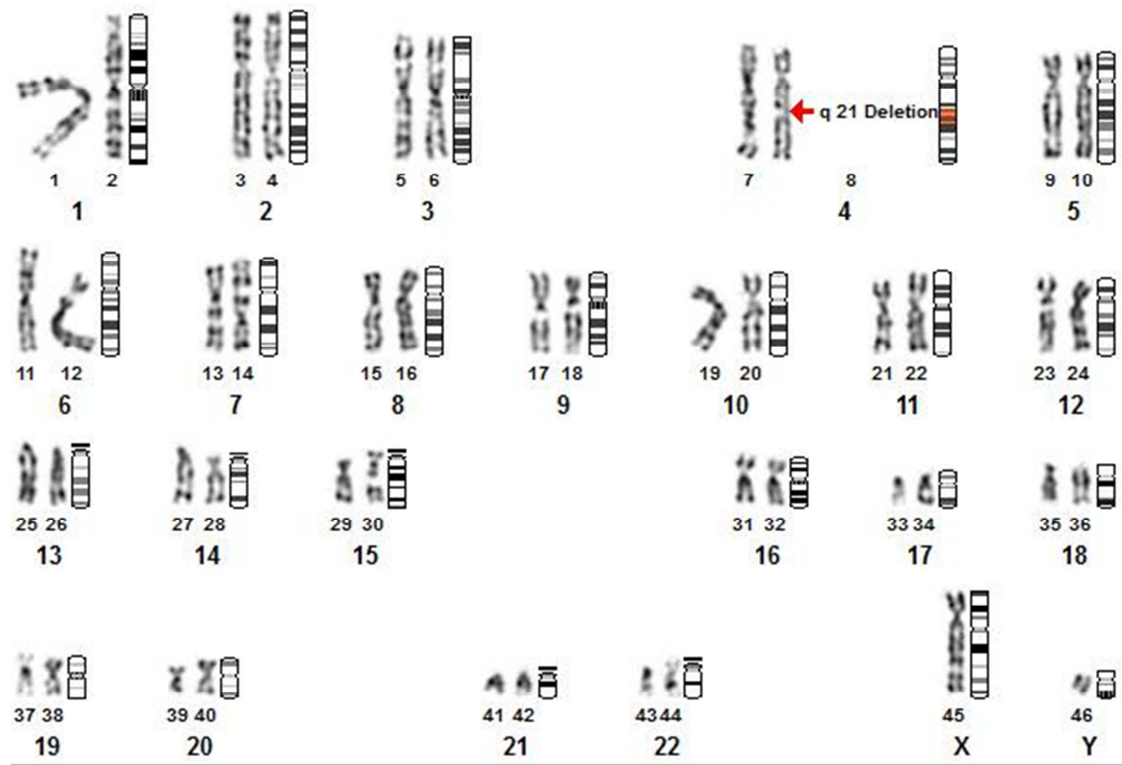


Figure 3. Karyotype analysis of cord blood of abortus with Fraser syndrome showing deletion on one of the chromosome number 4 at 4q21 region.

3. Exchange of Views

Fraser Syndrome is a rare genetic disorder with just more than 200 cases reported till date in the indexed literature [7]. It is a combination of acrofacial and urogenital malformation with cryptophthalmos. In 1986, Thomas *et al* laid down the diagnostic criteria for Fraser syndrome. Two major and one minor criterion or one major and at least four minor criteria are needed for the diagnosis [7].

Table 1. Diagnostic criteria.

Diagnostic criteria by Thomas <i>et al.</i> [7]	
Major	Minor
1. Cryptophthalmos	1. Congenital malformation of the nose
2. Syndactyly	2. Congenital malformation of the ears
3. Abnormal genitalia	3. Congenital malformation of the larynx
4. Sibling with cryptophthalmos syndrome	4. Skeletal defects
	5. Umbilical hernia
	6. Renal agenesis
	7. Mental retardation
	8. Cleft lip or palate

The findings in this case are compatible with the diagnosis of Fraser syndrome according to the major and minor criteria proposed by Thomas *et al.* [7]. In the present case, the major criteria were syndactyly and abnormal genitalia and the minor criteria were malformations of the nose, ears, skeletal defects and cystic dysplasia of the kidneys.

Syndactyly has been considered as a major feature of Fraser syndrome that occurs in 77% of the patients. Syndactyly is always cutaneous and in most cases involves

fingers and toes [15]. In the present case, there was syndactyly in the fingers.

In males, according to AM Slavotinek, CJ Tifft *et al* [1], abnormal genitalia such as cryptorchidism – (31.5%), micropenis, phimosis, chordee, hypospadias, and scrotal hypoplasia were noted [14]. In the present case, cryptorchidism and micropenis were noted.

Congenital malformation of the nose and ears has been considered as minor criteria. According to AM Slavotinek, CJ

Tiftt *et al* [1], nasal anomalies were common with having a broad nose or nasal bridge (20.5%), depressed or flat nasal bridge (11.1%), and bifid nose or a midline nasal groove (15.4%). Coloboma of the nares or notched nares were existing (11.1%). In the present case, there was saddle depression of dorsum of nose.

Malformed and/or low set ears (53.8%), microtia (16.2%), and atresia or stenosis of the external auditory meatus (17.9%) were noted [14]. In the present case, there was grade 3 microtia of ears.

According to Peer *et al* [13] laryngeal stenosis or atresia was reported in 30.8%. Choanal stenosis or atresia (76%) and subglottic stenosis (8.5%) were also described. In the present case lungs were massively enlarged and hyperechogenic resulting in compression of heart. The bronchial tree and trachea were dilated.

Skeletal defects has been considered as a minor feature of FS. According to AM Slavotinek, CJ Tiftt *et al* [1] absence or hypoplasia of the orbital or skull bones (10.2%) and defects in skull ossification (6.8%) were among the most frequent musculoskeletal anomalies. Talipes (8.5%) and abnormalities involving the pubic symphysis (7.7%) were relatively common. In the present case, bilateral clubfoot was noted.

According to AM Slavotinek, CJ Tiftt *et al* [1] out of 117 patients bilateral renal agenesis with or without agenesis of the ureters was present in 23.1% and unilateral renal agenesis with or without ureteral agenesis in 22.2%. The bladder was small or absent in 17.1% and cystic dysplasia of the kidneys was reported in 12%. In the present case, cystic dysplasia of the kidneys and small bladder was observed.

Fraser syndrome is an autosomal recessive inherited disorder. The parents of the affected children are sometimes but not always consanguineous. Consanguinity of marriage is reported in 15-24.8% of the cases [14, 15]. Parental consanguinity was also noted in this case.

The clinically variable anomalies corresponding with Fraser syndrome support the genetic heterogeneity of the syndrome. The gene of the Fraser syndrome was localized to 4q21 FRAS1 gene [1]. In the present study also, deletion in 4q21 region was noted.

Fraser syndrome is otherwise known as cryptophthalmos syndrome. However, cryptophthalmos is not an essential component of this syndrome. A review of 87 cases by Thomas *et al*, showed the absence of cryptophthalmos in 14 cases [7]. According to R. Koenig and J. Spranger *et al* [14] Fraser syndrome was noted without cryptophthalmos. In the present case also, there was absence of cryptophthalmos.

4. Conclusion

Fraser syndrome is an autosomal recessive disorder, and is associated with high mortality and morbidity. Hence, subjecting amniotic fluid for karyotyping between 16-20 weeks of gestation might confirm the presence or absence of the Fraser syndrome which may be helpful in early intervention.

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