Case Reports

S252W Mutation in Indian Patients of Apert Syndrome

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Two common mutations in the exon IIIa of fibroblast growth factor receptor 2 account for majority of the cases of Apert syndrome. They can be analyzed by amplifying the segment followed by testing for the abolition of restriction sites. We evaluated two children with typical features of Apert syndrome. A segment of FGFR2 exon IIIa was amplified by polymerase chain reaction. Restriction fragment length polymorphism was analyzed using enzymes MboI and BglI respectively for S252W and P253R mutations. The DNA segment was sequenced using ABI 310 automated DNA fragment analyzer. Both the patients showed S252W mutations. DNA sequencing confirmed the results of the restriction fragment length polymorphism. Our study is the first report from Indian subcontinent to show the prevalence of S252W mutation among Apert syndrome patients from Indian origin.

Key words: Acrocephalosyndactyly, Apert syndrome, Mutations.

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Craniosynostosis refers to premature fusion of sutures of skull. The overall birth prevalence is 343 per million newborns(1). Apert syndrome, with a birth prevalence of 15-16 per one million newborns accounts for about 4-5% of all cases of craniosynostosis. Apert syndrome is characterized by craniosynostosis, midface hypoplasia and symmetric syndactyly of hands and feet. Two common mutations in fibroblast growth factor receptor 2 (FGFR2) account for about 97% of the cases(2). The mutations S252W and P253R abolish a restriction site for MboI and BglI respectively. This provided the basis to study these mutations in two Indian patients with Apert syndrome.

Subjects and Methods

Case 1

Eleven months old female child was brought for evaluation of dysmorphism. She was born of a non-consanguineous marriage at full term by cesarian section in view of cephalopelvic disproportion. The child had apparently normal development. She was able to sit on her own and able to stand with support. She was babbling and understood spoken language. She did not have seizures or vomiting. Her weight was 6.5 kg (<5th percentile), length was 75 cm (50th percentile) and head circumference was 43.5 cm (within 2 SD). She had craniosynostosis, open anterior fontanel, and lambdoid suture, cleft palate, brachycephaly, bilateral proptosis, and syndactyly of both hands and feet (mitten hands). Anteroposterior diameter of skull was12.5 cm and transverse diameter was 14 cm. Deep tendon reflexes were brisk and plantar responses were extensor. Hearing and vision were apparently normal.

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Case 2

A one-month-old female, born to nonconsanguineous couple was referred for evaluation of dysmorphism. She had age appropriate physical growth. She had bilateral proptosis, midface hypoplasia, and brachycephaly, syndactyly of hands and feet, and mild clefting of uvula. She had high arched palate and broad thumbs and great toes in addition (*Fig.1*).

Informed consent was taken from the parents of both the subjects before performing the study. Polymerase chain reaction (PCR) followed by analysis of restriction fragment length polymorphism was done on DNA according to the published protocol(2). Both the subjects were heterozygous for S252W mutation. In both the individuals the two restriction sites were intact in the PCR product for BgII enzyme suggesting the absence of P253R mutations (*Fig. 2*). The mutation was confirmed by bidirectional sequencing of the PCR product using ABI 310 automated DNA fragment analyzer (*Fig.3*).

Discussion

We analyzed mutations in two patients with Apert syndrome. As described in the literature, we also found one of the two common mutations which accounts for 71% of the cases of Apert syndrome. To the best of our knowledge, no mutations have been reported earlier in Indian literature in patients with this condition.

The role of mutation detection in establishing the diagnosis is limited as the clinical phenotype is usually very characteristic. However, it is useful in situations where overlap between different craniosynostosis syndromes is noted. The same molecular methods can be used for prenatal diagnosis of the condition by chorionic villus sampling. The condition is usually due to de novo mutation



Fig. 1. Photograph of case 2 shows characteristic midface hypoplasia and syndactyly.



Fig. 2. Digested PCR products on 10% PAGE: Lanes 1, 2 and 3: normal, patient 1 and patient 2 after digestion with MboI; lane 4: undigested product; lanes 5, 6 and 7: normal, patient 1 and patient 2 after digestion with BgII.



Fig. 3. Sequencing of PCR products shows the nucleotide substitution.

and there is no significant increase in the risk of recurrence. But for many autosomal dominant and X-linked disorders germline

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mosaicisms are reported. The risk of recurrence due to germline mosaicism cannot be quantified, however indicates the need for prenatal diagnosis. Hence, it will be quite useful for relieving the anxiety of the couple with a previous affected child. The technique has been used earlier for prenatal diagnosis of the condition when sonographic features suggested the diagnosis of this particular condition(3).

Contributors: KMG was involved in clinical evaluation of patients, carried out molecular work and drafted the manuscript. SRP conceptualized the work and was involved in clinical evaluation of patients and correction of manuscript. FK was involved in DNA sequencing and contributed to drafting of manuscript. SA guided the molecular work and DNA sequencing. She also contributed to the correction of the manuscript. She will act as guarantor.

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REFERENCES

- Cohen MM Jr, MacLean RE, es. Craniosynostosis: Diagnosis, evaluation, and management. 2nd edn. New York: Oxford University Press, 2000.
- Park WJ, Theda C, Maestri NE, Meyers GA, Fryburg JS, Dufresne C, *et al.* Analysis of phenotypic features and FGFR2 mutations in Apert syndrome. Am J Hum Genet 1995; 57: 321-328.
- Hansen WF, Rijhsinghani A, Grant S, Yankowitz J. Prenatal diagnosis of Apert syndrome. Fetal Diagn Ther 2004; 19: 127-130.

Neonatal Bartter Syndrome

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Bartter syndrome is an inherited renal tubular disorder with hypokolemia, hypochloremic metabolic alkalosis, normal blood pressure with hyperreninemia and increased urinary loss of sodium, potassium and chloride. We report an infant with neonatal Bartter syndrome, who improved with potassium supplements.

Key words: Bartter syndrome, Neonate.

Bartter syndrome is a renal tubular disorder characterized by hypokalemia,

hypochloremic metabolic alkalosis, normal blood pressure with hyperreninemia and increased urinary loss of sodium, potassium and chloride. Usual clinical manifestations are failure to thrive, polyuria and episodes of dehydration. A 70-day-old boy, born to non-consanguineous parents presented with failure to thrive, persistent hypokalemia,

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