**X-linked Spondyloepiphyseal Dysplasia Tarda with Mutation in TRAPPC2 Gene: First Report from India**

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**Learning Point of the Article:**
Spondyloepiphyseal Dysplasia Tarda is a clinically, radiologically distinct syndrome which can be identified on genetic testing which enables definitive diagnosis, prognostication, counseling, management and prenatal diagnosis for such families.

**Abstract**

**Introduction:** X-linked spondyloepiphyseal dysplasia tarda (SEDT) is a type of short trunk skeletal dysplasia, occurring in males due to mutation in TRAPPC2 gene.

**Case Report:** We describe a large Indian family with multiple males affected with X-linked SEDT. The affected individuals presented with disproportionate short stature, short trunk, and barrel-shaped chest. Elder sibs aged 26 years and 31 years had back and hip pain. Premature osteoarthritis was seen requiring hip replacement surgery in one sib. The known pathogenic nonsense mutation c.209G>A (p.W70X) was identified in TRAPPC2 gene. This is the first mutation proven Indian kindred with X-linked SEDT.

**Conclusion:** Knowledge of molecular basis is essential to provide definitive diagnosis, accurate counseling, and prenatal diagnosis or early postnatal diagnosis for this rare condition.

**Keywords:** Genetic, India, spondyloepiphyseal dysplasia tarda, TRAPPC2 gene.

**Introduction**

X-linked spondyloepiphyseal dysplasia tarda (SEDT) (MIM) is a genetic disorder of the vertebral and epiphyseal growth, leading to short stature and degenerative osteoarthropathy. Based on the pattern of inheritance, there are three types, namely, X-linked recessive (MIM 313400), autosomal recessive (MIM 271600), and autosomal dominant (MIM 184100) [1]. The X-linked spondyloepiphyseal tarda is the most common form and mutations are attributed to TRAPPC2 gene found at locus Xp22 [1, 2]. We describe an Indian family with several males affected due to SEDT. Mutations were identified in TRAPPC2 gene in three male sibs and one clinically unaffected carrier sister.

**Clinical Summary**

Two male sibs born of non-consanguineous parentage, aged 15 years (patient III-6 in Fig.1) and 25 years (patient III-7 in Fig.1), respectively, were referred for short stature. They were noticed to be lagging behind their peers in height at around 5 years of age. Another maternal cousin aged 31 years (patient III-10, Fig.1) was also short and had a history of pain in joints of the knee, hip, and the back. He had left-sided partial hip replacement surgery at 30 years of age. Patient III-6 had the following anthropometry: Height 142.5 cm, weight 40 kg, head circumference 57 cm, lower segment (LS) 85 cm, upper segment (US/LS) ratio of 0.67, span 161 cm, and chest circumference 70 cm. Patient III-7 had the following anthropometry: Height 147 cm, weight 55 kg, head circumference 57 cm, lower segment (LS) 85 cm, upper segment (US/LS) ratio of 0.67, span 161 cm, and chest circumference 70 cm. Patient III-7 had the following anthropometry: Height 147 cm, weight 55 kg, head circumference 58 cm, LS 86 cm, US/LS ratio 0.70, span 166 cm, and chest circumference 98 cm. Patient III-10 had the following anthropometry: Height 148 cm, weight 58 kg, head circumference 56 cm, LS 87 cm, US/LS ratio 0.70, span 164 cm, and chest circumference 95 cm. All three had barrel-shaped...
Written informed consent was taken from the individuals for genetic testing. The research was conducted in accordance with the principles of Helsinki. Genomic DNA was extracted from peripheral blood. Polymerase chain reaction was performed covering all exons and exon-intron boundaries of the TRAPPC2 gene as per previously described methods [1]. Bidirectional Sanger sequencing was performed on automated capillary sequencer. Mutations are numbered from 1, beginning with the first nucleotide of the translation start [3]. The mutation nomenclatures were as per nucleotide ID NM_001128835.2 and protein ID NP_001122307.2. Patients III-6, III-7, and III-10 were confirmed to be hemizygous for the mutation c.209G>A (p.W70X) in exon 3 of TRAPPC2 gene and individuals II-6, II-9, and III-11 were confirmed as heterozygous carriers for the mutation.

Discussion

Spondyloepiphyseal tarda has been first described by Jacobsen, in 1939 [4]. Gedeon et al. identified the genetic defect in TRAPPC2 gene on locus Xp22 in three unrelated Australian families [5]. The TRAPPC2 gene has six exons and encodes a 140 amino acid protein with putative role in vesicular transport [5].

Several cases of spondyloepiphyseal tarda have been reported from India but none before with molecular confirmation. Pathare et al. reported the first case from Mumbai, Maharashtra, in 1991 [6]. Lakhkar and Raphael reported six unrelated individuals SEDT from Manipal, Karnataka [7]. SEDT does not exhibit any ethnic predisposition. Affected individuals have been described in European, American, Asian, and Australian populations (but not in African-Americans to date) [8]. One estimate suggests that the incidence is two persons per million [9].

The short stature in this family was characterized as disproportionate with short trunk. London Medical Databases lists 74 conditions with short trunk dwarfism. The differential diagnosis is based on clinical presentation and radiological skeletal features. The list of common differential diagnoses includes brachyolmia, spondyloepiphyseal dysplasia congenita, Morquio syndrome, spondyloepiphyseal dysplasia, multiple epiphyseal dysplasia, and Stickler syndrome [2]. The SEDT form of spondyloepiphyseal dysplasia is milder than the congenita type and is X-linked. Presentation is between 5 and 10 years with back or hip pain like the patients in the present series. Hands, head, and feet appear to be normal size, and final adult height usually ranges from 4’9” to 5’3” (range 4’1”–5’8’’). The SED congenita patients frequently have retinal detachment and cleft palate not seen in SEDT patients. Tiller and Hannig stated that the US/LS ratio is around 0.8 [2]. The US/LS ratio was 0.54 and 0.58 in our patients. Savarirayan et al. stated that there is a great degree of clinical variability even in the same family. The arm span exceeds the height by 10–20 cm and this was observed in our patients too [2,10].

![Figure 1](https://www.jocr.co.in/charts/fig1.png)

**Figure 1**: Three-generation pedigree shows a typical X-linked inheritance of spondyloepiphyseal tarda in this family. The horizontal bar (—) indicates that these cases have been examined by the authors; the asterisk (*) indicates that molecular diagnosis was possible in these cases. Patients III-6, III-7, and III-10 were confirmed to be hemizygous for the mutation c.209G>A (p.W70X) in exon 3 of TRAPPC2 gene and individuals II-6, II-9, and III-11 were confirmed as heterozygous carriers for the mutation.

![Figure 2](https://www.jocr.co.in/charts/fig2.png)

**Figure 2**: Radiological features in affected patient: X-ray of skull (a) (lateral view), hands and feet, and elbow (anteroposterior view) (d-f) were normal, thoracodorsal spinal vertebrae (b) (lateral view) show humpback sign of vertebrae (black arrows), X-ray pelvis with hip (anteroposterior view) (g) shows deformed femoral heads with premature osteoarthritis (white arrows), short femoral neck with decreased angle.
Characteristically, the vertebrae show a posterior hump of bone on their superior and inferior aspects when viewed laterally [11]. These findings appear in late childhood (after 5 years) but usually before puberty [12]. Anterior beaking of vertebrae can be seen, especially in lower lumbar vertebrae in young patients (Fig. 2). Mild epiphyseal dysplasia is seen in the hips, knees, and shoulders, but rarely in other joints (Fig. 2). Femoral necks are short and bent (coxa vara). Atlantoaxial instability can be present secondary to odontoid hypoplasia or os odontoideum. Hence, neurological examination is necessary. Scoliosis and lordosis of spine can be progressive warranting treatment such as bracing [2, 8, 12].

The p.W70X mutation observed in our patients has been previously described in other Chinese and British families [13,14]. However, in the British family, the nucleotide substitution was c.210G>A rather than c.209G>A (seen in our patients and the Chinese family). Fifty-seven unique mutations have been described till date (Human Gene Mutation Database, Cardiff, UK) [15]. About 90% of the mutations involve exons 4, 5, and 6 [16]. Deletion mutations comprise a third of all mutations. This unusual high deletion frequency, particularly in a gene encoding only a small protein of 140 amino acids, may possibly be explained by the five truncated pseudogenes located on chromosome Yq11.23, which may cause homologous recombination and slipped mispairing [17].

The recurrence risk in pregnancies of carrier females is 50% of males who would be affected. Antenatal ultrasound will not able to diagnose affected fetuses. Due to availability of mutation, genetic counseling, extended family screening, and prenatal diagnosis or early postnatal diagnosis are possible in this family.

Treatment remains supportive. Hip replacement is required in several patients with premature degenerative hip arthropathy, during the third decade. Premature arthritis can be prevented by weight management and low impact exercise such as swimming and cycling. In view of possibility atlantoaxial instability, it is prudent to avoid high-risk physical activities such as football, rugby, and trampoline. The patient may require psychosocial counseling for the short stature. Role of growth hormone in increasing final height is not proven [2, 8, 10].

Conclusion

This is the first molecular proven kindred with X-linked SEDT from India. Knowledge of molecular basis is essential to provide definitive diagnosis, accurate counseling, and prenatal diagnosis or early postnatal diagnosis for this rare condition.

Clinical Message

X-linked SEDT is a clinically and radiologically distinguishable genetic syndrome. DNA diagnosis is essential for accurate diagnosis, genetic counseling, prognostication, management, prenatal diagnosis, and early postnatal diagnosis.

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