

# A rare case of translocation (12;22) (p13;Q) in Ewing's sarcoma

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## ABSTRACT

Cytogenetic or immunohistochemical studies are often required to differentiate Ewing's sarcoma (ES) from other small round cell tumors. Herein we report a case of 13-year-old boy who presented with a large presacral lesion. Hemogram and biochemical parameters were normal except lactate dehydrogenase showing value of 96.40 IU/L, magnetic resonance imaging of the spine showed a large mass in presacral lesion (8 cm × 7 cm × 9 cm), with destruction of the sacrum (S2 S3 and S4) with interspinal extension. Bone scan showed multiple pelvic bone lesions, radiograph of chest, ultrasound of abdomen, pelvis and electrocardiogram were within normal limits. Bone marrow was not involved. Cells from the fine needle aspirate were cultured for short term using RPMI medium and karyotype obtained showed a t(12;22)(p12;q12) instead of the classic t(11;22). Diagnosis of ES was also confirmed by studies using immunohistochemistry for MIC2 which was positive, synaptophysin was inconclusive and leukocyte common antigen, desmin negative. This case provides evidence of the importance of chromosome 22, in the etiology of the disease.

**Key words:** Cytogenetics, Ewing's sarcoma, karyotype

## INTRODUCTION

The t(11;22)(q24;q12) is the most common translocation for ES and is present in more than 85% of cases.<sup>[1]</sup> This translocation leads to an EWS/FLI1 fusion gene in all cases. In a few instances, complex translocations, involving chromosomes 11 and 22 and a third chromosome or other variant translocations not involving chromosome 11 also have been reported.<sup>[2]</sup> They are molecularly characterized by expression of chimeric transcripts generated by specific chromosomal translocations, most commonly involving fusion of the EWS gene to a member of the Ewing tumors family of transcription factors, which include FLI1, ERG, ETV1, E1AF and FEV.<sup>[3]</sup> Sarcomas are a heterogeneous group of cancers derived from the connective tissue lineage. The

etiology of these tumors is unknown, and the vast majority of the cases occur without known hereditary factors. In the last two decades, the finding of specific acquired chromosomal alterations in sarcomas has helped in many cases to understand the underlying genetic basis of this tumor.<sup>[4]</sup>

Ewing's sarcoma (ES), peripheral primitive neuroectodermal tumors (pPNETs) and Askin tumors are referred to as Ewing tumors (ETs). ES is the second most common malignant bone tumor in children and young adults, occurring with a male/female ratio of 1.5:1.<sup>[1,5]</sup> The most frequent primary site is the femur, followed by the pelvis arising from the ileum, ischium, pubic bone, or sacrum. Tumors may also originate in the tibia, fibula, or the bones of the feet. ES is the result of a translocation between chromosome 11 and 22, which fuses the *EWS* gene on chromosome 22 to the *FLI1* gene on chromosome 11. *EWS/FLI1* functions as the master regulator.<sup>[6]</sup>

A review of literature shows the chromosome partners involved in translocation with chromosome 22 in sarcomas are 1, 2, 4, 6, 7, 9, 11, 12, 17, 19, 20 and 21 as shown in the Chart 1. Nearly 85% cases of ES contain the t(11;22)(q24;q12) Chromosomal translocation that encodes the EWS/FLI

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oncprotein. Besides the t(11;22), however, many cases have otherwise simple karyotypes with no other demonstrable abnormalities. Furthermore, it seems that an underlying genetic susceptibility to ES, if it exists, must be rare.<sup>[7]</sup> Although Pro-B acute lymphoblastic leukemia showed t(12;22)(p13;q12) with EWSR1 → ZNF384 genes and expression of myeloid antigens, having relatively good prognosis,<sup>[8]</sup> the present case showed same translocation t(12;22)(p13;q12) but in ES.

### CASE REPORT

The case we present here is a 13-year-old boy who presented with the complaints of left lower limb weakness for 5 months and low back pain for 6 months. Hemogram and biochemical parameters were normal except lactate dehydrogenase (LDH) showing value of 96.40/IU/L, magnetic resonance imaging of the spine showed a large mass in presacral lesion (8 cm × 7 cm × 9 cm), with destruction of the sacrum (S2 S3 and S4) with interspinal extension. Bone scan showed multiple pelvic bone lesions, radiograph of chest, ultrasound of abdomen, pelvis and electrocardiogram were within normal limits. Bone marrow was not involved. Fine-needle aspiration cytology of the mass proved it to be a small round cell tumor. Immunohistochemistry for MIC2 was positive. Synaptophysin was inconclusive and leukocyte common antigen, desmin were negative [Figure 1a and b].

### CYTOGENETICS

#### Materials and methods

The fine-needle aspiration material obtained from the primary tumor mass was suspended in RPMI-1640 medium supplemented with 20% qualified fetal calf serum (GIBCO). Overnight cultures (16-18 h) were set up. The following day, Karyomax Colcemid solution (final concentration 0.05 µg/ml) was added for 30 min. Cells were subjected to hypotonic treatment (KCl-0.075 M) for 30 min at 37°C and fixed in methanol:acetic acid (3:1). A chromosome analysis was performed on GTG-banded metaphases and karyotype analysis [Figure 2a and b] was interpreted according to ISCN.<sup>[9]</sup>

### DISCUSSION

Typical ET can be diagnosed on morphology and immunohistochemistry, but the variants need cytogenetic or molecular confirmation of the diagnosis. ES and peripheral PNET strongly express the MIC2 (CD99) antigen in very high amounts which represents a highly selective and almost unique feature of these cells, therefore MIC2 analysis is a useful part of the panel of tests used in the differential

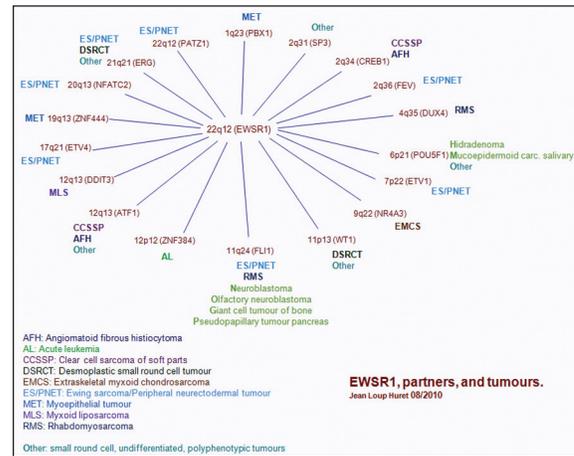


Chart 1: EWSR1 partners and tumors

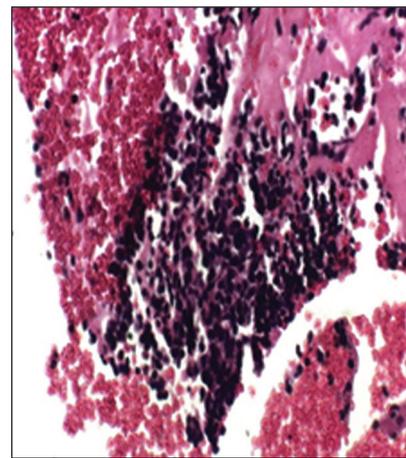


Figure 1a: Cell block of Ewing's sarcoma (H and E)

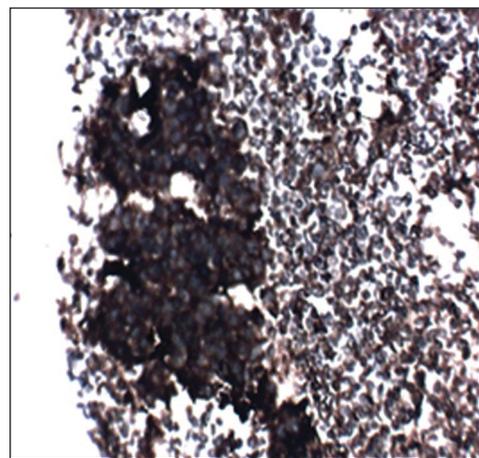
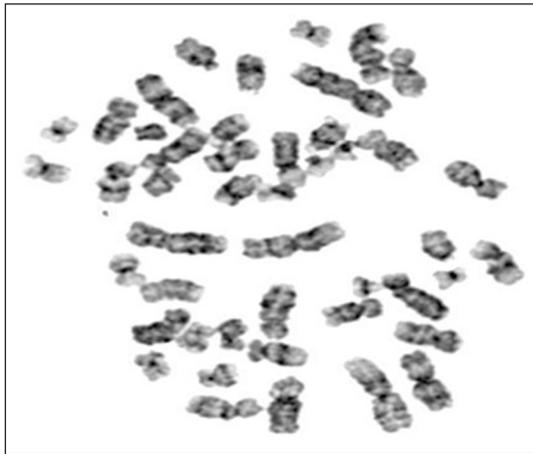
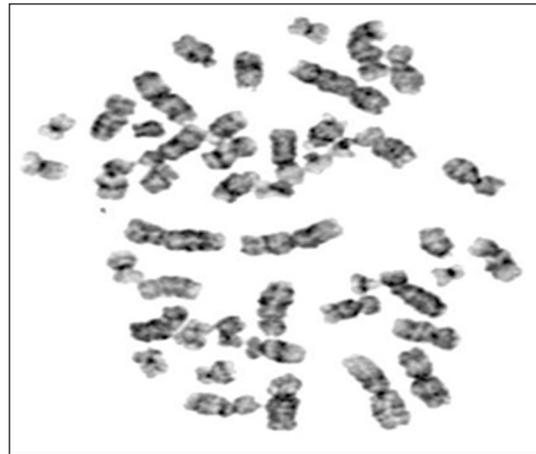


Figure 1b: Cell block of Ewing's sarcoma positive for MIC2

diagnosis of ES/pPNET.<sup>[10]</sup> But the histological grade is of no prognostic significance, however, as all ES are of high grade. Fever, anemia and elevation of the number and values of white blood cell, erythrocyte sedimentation rate and LDH have been reported to indicate more extensive disease and a poorer prognosis.<sup>[11]</sup>



**Figure 2a:** GTG banded metaphase



**Figure 2b:** GTG-banded karyotype showing 46, XY, t(12;22)(p13;q12). Arrows indicate break points

**Table: 1 The frequency of t(11;22) and its variants in ES**

Translocation	Gene fusion product	Frequency in Ewing tumors %
t(11;22)(q24;q12)	EWS-Flt1	>90
t(21;22)(q22;q12)	EWS-ERG	5
t(17;22)(q12;q12)	EWS-E1AF	<1
t(2;22)(q33;q12)	EWS-FEV	<1
t(7;22)(p22;q12)	EWS-ETV1	1
t(4;22)(q31;q12)	EWSR1-SMARCA5	1

ES – Ewing's sarcoma; EWSR1 – Ewing sarcoma region 1

The incidence of classical t(11;22) in ES is around 90% approximately. Involvement of other chromosomes participating in translocation with 22 such as 21 is 5%, 2%, 4%, 7% and 17% are 1%, respectively and are rare [Table 1].

The present case adds on to this rare list of abnormalities in ES. The *EWS-FLI1* translocation has been reported in two polyphenotypic tumors and two rhabdomyosarcomas.<sup>[11]</sup> Cytogenetic abnormalities are highly specific to ETs; chromosome 22 is most frequently involved in structural changes detected in cells of ES. Although t(11;22) is most frequent, chromosome 22 is also involved in other translocations in neoplasia, suggesting that the break point on chromosome 22 seen in ES cells may be a more important factor in the origin of this tumor than the loci to which the deleted segment is translocated.<sup>[1,5,6,12-14]</sup> In a case of ES which Whang-Peng *et al.*<sup>[15]</sup> examined it had a translocation involving chromosome 6 and 12, with a break point (12p13).<sup>[15]</sup> This finding with the absence of t(11;22) raises the possibilities that soft-tissue ES is a tumor significantly different in origin and pathogenesis from ES of the bone because our case presented with the tumor of the sacral bone. More cases need to be studied to determine the significance and incidence of this abnormality in ES. It is also important to determine whether additional structural chromosomal aberrations are present in

ES tumors because it appears that a more complex karyotype with multiple chromosomal aberrations is associated with a poor outcome in ES.<sup>[16,17]</sup>

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**How to cite this article:** Kousar Jahan SK, Mayanna MG, Kavitha BL, Patil A, Kumari P. A rare case of translocation (12;22) (p13;Q) in Ewing's sarcoma. *Indian J Med Paediatr Oncol* 2014;35:89-92.  
**Source of Support:** Nil, **Conflict of Interest:** None declared.