Neurological expression of an inherited translocation of chromosomal 1 and 7

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ABSTRACT

An unbalanced translocation of chromosome 1 and 7 (t(1;7)) associated with neurological phenotype and brain malformation has rarely been reported. This clinical report describes 3 siblings with brain malformations and a 13.5 Mb duplication of 1q42.3q44, and a 7.6 Mb duplication of 7q36.1q36.3 detected by array comparative genomic hybridization. This unbalanced t(1;7) was found to be inherited from a balanced translocation from the mother. Our objective in presenting this particular case is to compare our patients’ clinical and neuroimaging features with those of other cases reported in the literature. Our data can then contribute facilitating genotype-phenotype correlations.

Case Report. We evaluated 3 patients from a consanguineous Saudi family (Figure 1). The main clinical, MRI, and associated malformations are summarized in Table 1.

Array-comparative genomic hybridization performed in patients III-6, III-7, and III-8 detected a deletion at 1q42.3q44 spanning 13.5Mb, and a duplication at 7q36.1q36.3 spanning 7.6Mb. The fluorescence in situ hybridization studies confirmed both the deletion and the duplication (1q42.3q44, 7q36.1q36.3). Studies of the parents showed normal karyotype in the father (46, XY), and an unbalanced reciprocal translocation in the mother 46,XX,t(1;7)(q44;q36). Whole exome sequencing to rule out other possible genetic causes was negative. Clinically, our patients had severe intrauterine growth restriction and oligohydramnios. They showed dysmorphism in the form of wide forehead, hypotelorism, upturned nose, and narrow anterior fontanelle. The neurological evaluation showed microcephaly (head circumference...
between -3 and -4 standard deviation) in all 3 patients. All patients had axial hypotonia and normal deep tendon reflexes. They showed global developmental delay with slow improvement. Patients III-6 and III-8 developed generalized tonic-clonic seizures at the age of 8 months, and the electroencephalogram showed multifocal discharges. Other non-neurological abnormalities included the following malformations: undescended testis, hypospadias, and congenital heart disease. The family had a history of 4 spontaneous abortions.

The neuro imaging revealed multiple brain abnormalities. An MRI in patients III-6 and III-8 demonstrated complete agenesis of the corpus callosum, hypoplasia of the inferior vermis, and thin optic nerves, associated with mild delayed myelination (Figure 2), while in patient III-7 (Figure 2), it showed thick corpus callosum and inferior vermis hypoplasia.

**Discussion.** In humans, the incidence of balanced chromosome translocations is approximately one in 500. Balanced reciprocal translocations are associated with a 50% risk of spontaneous abortions and a 20% risk of genetic abnormalities. Unbalanced translocations involving chromosome 1 and 7 are well-recognized cytogenetic abnormalities in hematological malignancies. So far, only 7 patients with the t(1;7)-related neurological phenotype are reported in the literature. In this paper, we report 3 more patients. Our patients’ phenotype is dominant by hypotonia, developmental delay, microcephaly, seizures, and particular neuroimaging findings.

Two neurological phenotypes associated with t(1;7) can be described. The first phenotype is associated with behavioral disorders in the form of early onset schizophrenia, and autistic behavior. Neuroimaging in these cases is reported as normal. This phenotype may be explained by possible susceptibility loci for specific language disorder, or autism, reported from several linkage studies, both of which are positioned around 1q31-q34. The second phenotype, as in our patients, is associated with brain malformation. Two types of malformations have been reported: holoprosencephaly, as described by Chuang et al and Schinzel, and abnormal corpus callosum and cerebellum as in our patients. The occurrence of brain malformation in t(1;7) may be explained by the involvement of several genes. Deletion of 7q36 clearly involved the sonic hedgehog gene, as described by Chuang et al and Schinzel, and abnormal corpus callosum and cerebellum as in our patients. The occurrence of brain malformation in t(1;7) may be explained by the involvement of several genes. Deletion of 7q36 clearly involved the sonic hedgehog gene, and haploinsufficiency of the sonic hedgehog gene has been shown to be the responsible molecular event in some cases of holoprosencephaly. The sonic hedgehog gene exerts a strong influence on the differentiation of ventral and medial structures of the prosencephalon, and the defective expression resulting from a mutation of this gene is thought to be the molecular basis of the human malformation holoprosencephaly with a broad spectrum of clinical severity. The defect in cleavage also influence the development of other cerebral structures that occur later in ontogenesis resulting in an
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abnormality in the formation of the corpus callosum, and various migrational disorders, such as dysplasia.

In conclusion, we expand the number of patients with t(1;7)-related neurological disease to 10, and confirm the core features of the neurological phenotype of the disease as early psychiatric features or developmental delay, microcephaly, and seizures associated with brain malformation. We suggest that neuroimaging in these patients reveals a characteristic pattern that could help with early diagnosis. Further research is needed to understand the signaling network, and to shed light upon the role of chromosomes 1 and 7 in the developing brain.

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References