Bilateral split hand/foot malformation and inv(7)(p22q21.3)

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Abstract
A boy with typical tetramelic split hands and feet is described. In addition, there was a large arteriovenous malformation of the right arm. Chromosome studies showed a pericentric inversion of chromosome 7: 46,XY,inv(7)(p22q21.3). Inspection of the extremities and chromosome studies in the parents were normal. This case confirms the suggested localisation of a locus, important for early limb differentiation, on the long arm of chromosome 7, most probably in the chromosomal region 7q21.2-7q21.3. Previously reported cases are reviewed briefly.

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Figure 1 Case 1 at the age of 1 day. Central reduction defects of all four limbs are visible, as is the haemangiomatous malformation of the right lower arm and hand.

To date 12 cases of split hand/foot anomalies and a chromosomal rearrangement involving 7q21-7q22 have been reported.1-12 We report the 13th case, who also had haemangioma of a limb, not described before in this chromosomal syndrome.

Case report
The proband is a boy born after an uneventful pregnancy. He is the first child of healthy, unrelated parents. The family history did not include any congenital malformations. Immediately after birth congenital anomalies of the hands and feet were noted, and he was referred to our hospital. Physical examination at the age of 2 days showed typical central reduction defects (split hands and feet) of all four extremities (fig 1). On the left hand the third finger was missing and on the right hand two rays were present; the most ulnar ray was swollen and red-blue coloured. On the left foot only the first and fifth toes were present, with a deep cleft in between. The right foot also showed absence of the second, third, and fourth toes and a deformity of the first toe, which was in a retroflexed position. A soft precordial systolic murmur was noted. Further examination showed no other abnormalities. X-ray studies of the hands and feet confirmed the split hand and foot malformation (fig 2). Arteriographic studies of the right arm showed an arteriovenous malformation on the ulnar side (fig 3). The subclavian artery was large, the radial artery was slightly hypoplastic, and the ulnar and interosseal arteries were clearly enlarged and drained into an arteriovenous conglomerate with two large veins attached to it. At the age of 2 years the boy seems mentally normal and the arteriovenous malformation, visible on the outside of the right hand, seems to have involuted somewhat. The patient is able to use both hands and can grip several objects. Simple orthopaedic footwear facilitates normal gait in this boy, who will attend playschool in due course.

Physical examination of the hands and feet of the parents was normal.

CYTOGENETIC STUDIES
Chromosomes of the patient were examined in lymphocytes from peripheral blood using
present (the second, third, and fourth rays and a bifid distal thumb phalanx. Above right is the left hand showing absence of the third ray. Below left is the right foot with only two rays present (the lateral one tilted upwards). Below right is the left foot with apparent absence of the second, third, and fourth rays.

Figure 2  X rays of all four extremities taken at the age of one day, showing typical central reduction defects (split hands/feet). Above left is the right hand with absence of the second, third, and fourth rays and a bifid distal thumb phalanx. Above right is the left hand showing absence of the third ray. Below left is the right foot with only two rays present (the lateral one tilted upwards). Below right is the left foot with apparent absence of the second, third, and fourth rays.

routine procedures and Giemsa staining. An apparently balanced inversion of chromosome 7 was found. The karyotype was: 46,XY,inv(7) (p22q21.3) (fig 4). The chromosomes were painted with a chromosome 7 library. The rearranged chromosome contained only chromosome 7 material and no segment of that chromosome was found elsewhere in the karyotype (results not shown).

A fibroblast culture from a skin biopsy was established and stored and is available for interested researchers.

The chromosomes of the parents were normal.

Discussion
In this case, 13 cases of split hand/foot anomaly and chromosomal aberrations involving 7q21-22 have been described so far. In addition there is one report of a combination of split hand/foot and a chromosomal aberration involving 7p11. The cytogenetic data of these published cases and of our case are summarised in fig 5. Two reports concern split hands/feet as part of an EEC syndrome.

The haemangiomatic malformation described in our case has to our knowledge not been described before in association with split hands/feet. However, a haemangiomata at the site of limb deficiency has been described recently in a case of unilateral transversal limb defect.

Split hands/feet can be inherited as an autosomal dominant trait with reduced penetrance and variable expression. Although the locus of the gene for ectrodactyly has not yet been assigned, the number of reports of split hand/foot malformations with chromosomal aberrations involving 7q21-22 strongly suggest a localization of the gene(s) responsible in that region. Cases of 7q aberrations and split hand/foot malformation published so far are graphically represented in fig 5.

As can be seen in fig 5, the only discrepant breakpoint on chromosome 7 in these published cases was described by Hasegawa et al, who found a balanced t(7;9)(q11.2;p12) in three people with EEC syndrome belonging to the same family. This has led to the suggestion that the critical region for EEC syndrome is on 7q11.2-7q21.3. We think, however, that the case of Hasegawa et al might still be in keeping with a hypothetical localisation of the gene responsible to 7q21-22, admitting that no recombination has occurred between 7q11.2 and 7q21-22 in the two informative meioses of their family. It is conceivable that the structural abnormality of chromosome 7 entailed a reduced recombination in 7q, similar to that recently shown for chromosome 21, when involved in a Robertsonian translocation.
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Figure 4  Partial karyotype of case 1 showing chromosomes 7 from three metaphases, the normal chromosome 7 on the left and the abnormal chromosome 7 on the right. The idiogram of chromosome 7 shows the inversion breakpoints in bands 7p22 and 7q21.3 (arrows).

Figure 5  Schematic drawing of the cytogenetic breakpoints in reported cases with split hand/foot malformation and an aberration in the chromosomal region 7q. Represented are 12 previously published cases (DP, Pf, Ta, Mo, Ri, Ro, ME, Sh, Ah, Na, Ko, Ge) listed in the references, our case, and another published case (Ha) with a different breakpoint. For discussion see text.
the other hand, it is also possible that this is an example of locus heterogeneity, as it was reported recently that not all cases of split hand/foot anomalies are linked to chromosome 7q.18

It appears that chromosome band 7q21.3 or the proximal region of band 7q22 or both are the most likely candidate regions for such a gene or genes. In our case, there is either a submicroscopic deletion of euchromatin in 7q21.3, or the breakpoint falls in the supposed split hand/foot gene itself, leading to an abnormal gene product.

It is somewhat puzzling, however, that ectodactyly has been associated not only with balanced structural abnormalities involving breakpoints at 7q21-22, but also with deletions apparently involving these bands (fig 5). In fact, the former mechanism would still be compatible with partial disruption of the gene, whereas the latter implies loss of the locus where this disorder is supposedly located.

Dominantly inherited gene disorders are usually the result of point mutations or partial deletions leading to abnormal gene products, rather than to complete deletion/inactivation of one allele. Therefore, it is possible that other genetic mechanisms, such as parental imprinting, might also be involved in the pathogenesis of this malformation.19 Parental imprinting does not seem to be likely in view of a recently published family with split hand/foot anomaly and a balanced translocation involving 7q22.1,19 giving no indication of genomic imprinting. The determination of the parental origin of the rearranged chromosomes 7 associated with ectodactyly may shed light on possible (patho)genetic mechanisms involved in split hand/foot malformation.

Dominant split hand/foot is heterogeneous and has known variable expression and reduced penetrance. Elucidation of the genetic or epigenetic mechanisms responsible for these phenomena will be aided by the identification of the causative gene(s), and rare cases like ours may help in the gene localisation.