Association of 13q deletion and Hirschsprung’s disease

Sir,

It was interesting to read the paper by Lamont et al1 about the association of 13q deletion and Hirschsprung’s disease. We have observed a similar case.

An eight month old boy was investigated in our genetic service. The family history was negative. The mother’s first pregnancy ended with a spontaneous abortion. The proband was born of the second uneventful pregnancy with a birthweight of 3000 g. At two weeks of age Hirschsprung’s disease was diagnosed and sigmoidostomy was performed.

At eight months of age the following features were observed: mild craniofacial dysmorphism (fig 1) including hypertelorism, strabismus, small nose, and low set ears, a simian crease on the left palm, a small penis, and bilateral cryptorchidism. Cytogenetic analysis was carried out using GAG, G, and C banding techniques. The karyotype was 46,XY,del(13)(q22→qter) (fig 2). The karyotypes of the parents were normal. At three years of age the craniofacial dysmorphism was more pronounced (fig 3) and there was marked psychomotor retardation.

It is well known that Hirschsprung’s disease may be associated with various dysmorphic syndromes and occasionally with structural autosomal aberrations,2 but it usually occurs as a single malformation. Further observations are needed to ascertain any association between 13q deletion and Hirschsprung’s disease. However, if the q22 region is involved in the deletion, as in the two cases observed by Lamont et al1 and in our case, this connection should be given serious consideration.

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Dominantly inherited cleft lip and palate

Sir,

Temple et al (J Med Genet 1989;26:386–9) reported two families with dominantly inherited cleft lip and palate. We have recently seen a similar situation in a family (figure) presenting with non-syndromic, unilateral cleft lip and palate in at least three and probably four successive generations.

All affected family members were personally examined and in none of them could associated clinical findings suggestive of Van Der Woude syndrome or another autosomal dominantly inherited syndrome associated with cleft lip and palate be found.

III.5 and his wife asked for genetic advice after his healthy sister (III.10) gave birth to her first child (IV.5) with a left sided cleft of the lip, hard and soft palates, and alveolus. No lip pits or fistulae were present and the child was otherwise normal. On thorough clinical and x ray examination, no minimal signs of cleft lip/palate were found in III.10. The child’s father (III.9) had no clinical abnormalities and his family history was negative.

One brother (III.2), one sister (III.4), and a niece (IV.3) of the consultand were born with a right sided cleft of the lip and palate. His mother (II.2) also had a cleft lip and palate at birth. She underwent several surgical repairs, including closure of the lip in early childhood, closure of the palate at 50 years of age, and a Lefort I type osteotomy because of retrognathia of the maxilla and hypoplasia of the upper cheek. No preoperative photographic data were available but clinical examination showed that the cleft had been right sided. According to all family members, the dead maternal grandmother (I.2) had also had a unilateral cleft lip and palate.

Except for II.2, whose parents were first cousins, consanguinity was excluded between the parents of all the other affected family members.

The linear pattern of inheritance in this family is suggestive of autosomal dominant rather than multifactorial inheritance and adds further evidence to the hypothesis, which has already been suggested in other family studies,1 that a single major dominant gene is responsible for cleft lip and palate in some families.

If we accept an autosomal dominant transmission pattern in this family, subject III.10 is a non-penetrant heterozygote. Under the assumption of autosomal dominant inheritance with reduced penetrance, III.5 and his wife were given a maximum risk for an affected child of 8.6%. A detailed report of this family study is in preparation.

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Reference

Are abortions more or less frequent once prenatal diagnosis is available?

Sir,

Harris et al1 have recently stated that the application of DNA techniques to inherited diseases reduces the risks of fetuses being aborted, since the majority will be found to be healthy after testing; they report 18 abortions of fetuses at risk for Duchenne muscular dystrophy, and state that this number is a reduction from previous years. However, our early experience from the West Midlands was different from that in Manchester, and showed that if prenatal diagnosis is not available, women at risk tend either to refrain from pregnancy or to take the risk and complete the pregnancy.2 Similar observations have been made in Ontario.3 We thought it important to be certain of women’s attitudes and behaviour before analysing any effect

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Footnotes: