Recessively inherited lower incisor hypodontia

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Hypodontia, congenitally missing teeth, is common in modern man. The teeth most often missing in populations of European origin are the upper lateral incisors and second premolars. The condition is known to have a strong genetic component. At present two mutated genes in humans, MSXI\(^1\) and PAX9\(^2\), are known to cause missing permanent teeth. Mutations in MSXI\(^1\) can also cause orofacial clefting.\(^3\) Several experimental and clinical studies indicate that other genetic components are also involved.\(^4\) Hypodontia is also often seen in syndromes, particularly in those which present with other ectodermal anomalies,\(^5\) and in non-syndromic patients with cleft lip/ palate.\(^6\) Alveolus with or without cleft palate.

The population prevalence of the common congenital hypodontia (IPH, MIM 106600) is 8-10% in healthy European children. Some or all third molar teeth are missing in one-fifth of the population,\(^10\) and missing third molars are seen in varying combinations in IPH patients and/or family members.\(^11\) Family studies also indicate that peg shaped upper lateral incisors, impacted canines, rotated bicuspids, and short root anomaly (SRA) are caused by the same genetic components that cause missing incisors and premolars.\(^12\) The condition is inherited as an autosomal dominant trait\(^15\) with reduced penetrance and is mostly restricted to the permanent dentition. When a large number of teeth (>6) are congenitally missing, the term used is oligodontia (MIM 604625). The prevalence of oligodontia in European populations is estimated at 0.08%,\(^17\) but this figure also includes syndromic patients.

We describe 37 Finnish patients from 34 families with several lower incisors and upper lateral permanent incisors congenitally missing. In half of the patients, the corresponding deciduous teeth had either been missing or peg...
Method

Ten patients with missing lower and upper lateral incisors were first seen at the Hypodontia Unit of the Department of Pedodontics and Orthodontics, University of Helsinki over the past five years. Their parents did not display similar hypodontia, but sometimes had one congenitally missing permanent tooth. In two families, the condition was seen in sibs. In order to discover the prevalence of this condition in Finland, a questionnaire was sent to all active orthodontists and municipal health centres in the country (362 letters). We received radiographs of 220 patients and 65 patients selected from these were further studied after giving their consent.

Inclusion criteria were three or four congenitally missing lower incisors, at least two congenitally missing or peg shaped lower permanent incisors together with missing upper lateral incisors, preceded by agenesis of at least one lower deciduous incisor, and a pedigree consistent with an autosomal recessive mode of inheritance. Exclusion of obvious dominant oligodontia, anhidrotic ectodermal dysplasia (EDA, also carriers), and incontinentia pigmenti (IP) was attempted by careful clinical examination, anamnestic information, and pedigree analysis. The final sample consisted of 37 patients from 34 families. Three pairs of sibs were included (fig 1).

The patients, parents, some grandparents, and sibs were examined radiographically and interviewed, and facial and oral photographs were taken. Alginate impressions were made of all cooperative patients. Dental age was calculated from radiographs by the age medians for tooth formation and standard deviations (SD) for dental ages.

Parents and sibs showing dental anomalies such as hypodontia of one or two teeth, peg shaped teeth, retained cuspids or taurodontic teeth are shown in fig 1. The ancestors were traced back to 1850 from local church registries using the names, dates, and birth places of parents. Microfilm copies in The National Archives of Finland were used for earlier periods. To analyse the mode of inheritance, the ratio of affected to healthy sibs was corrected for the absence of healthy sibships born to two heterozygous parents.

For DNA analysis, samples of venous blood were taken from patients, grandparents, parents, and sibs. The study was approved by the Ethics Committee of the Institute of Dentistry, University of Helsinki.

Results

All 37 patients were under 22. Their clinical characteristics are presented in table 1 with typical panoramic radiographs in fig 2. Nine of the patients had no permanent lower incisors, two of them had also lacked the corresponding deciduous teeth. In five, the permanent lower molars were taurodontic (fig 2B). In one, the upper permanent lateral incisors were peg shaped with an anomalous structure (fig 2B). In four patients, three missing lower permanent incisors were evident. Of these 13 severely
affected patients, 10 were boys. The remaining 24 had a varying combination of missing or peg shaped lower and upper permanent and deciduous incisors. A deciduous tooth/teeth was missing in 19/36 (53%). In addition to incisors, other permanent teeth were also missing (fig 3). Mean dental age based on tooth formation was slightly delayed (SDS 0.8). Taurodontism was noted in the molar teeth of 16/26 patients (62%).

Photographs of the younger patients are shown in fig 4. Hair, nails, eyelashes, eyebrows, and perspiration were normal. Heights and weights were also normal for age. A large proportion of the patients reported allergies (62%), such as atopic skin (52%) and asthma (43%), diagnosed by a doctor.

The male/female ratio of the patients was 19/18 (1.05). The proportion of affected sibs was calculated from 30 sibships with 76 children. The apparent proportion of affected sibs was 0.43. After mathematical correction for the absence of healthy sibships born of two heterozygous parents, using the a priori correction of truncate complete ascertainment, the proportion was 0.22. Pedigrees of 31 families were traced back at least to the fifth generation, and, when possible, to the late 17th century (10 generations). In two families, the parents of the proband had a common ancestor six and seven generations back (fig 5), but no other family linkages between the families were found. A map of Finland, where the birth places of the great grandparents of the patients are marked, is shown in fig 6. In the case of 23/31 index cases, the maternal and paternal ancestors originated from the same rural area.

Minor dental anomalies in the form of a missing upper lateral incisor or a missing third molar/molars was seen in 41% of the parents and in 27% of the healthy sibs (fig 1). Both parents and sibs reported allergies (46% and 40% respectively) and skin problems (46% and 35% respectively). Ten of the 29 examined mothers (34%), eleven of the 22 fathers (50%), six of the 22 healthy sisters (27%), and six of the 20 healthy brothers (30%) are shown in the pedigrees on the basis of their minor dental anomaly (fig 1).

Discussion
Our attempt at a nationwide ascertainment resulted in a response to more than half of our questionnaires, but because the oldest patient was only 22, it is likely that we only found a proportion of the total number of RIH patients in Finland. By using strict criteria in an effort not to overdiagnose, some of the cases reported to us and some of the mildly affected sibs were perhaps misclassified.
There are many published reports of conditions with missing lower incisors. In Japan the prevalence of tooth agenesis is of the same order as in Europeans, but the lower lateral incisor is the most commonly missing tooth.\textsuperscript{21} Witkop syndrome is an autosomal dominant condition with missing lower incisors and dysmorphic nails.\textsuperscript{22} In a patient from Minnesota,\textsuperscript{23} dentition and a face very similar to that of our patients can be seen. First cousins of Egyptian origin, born of consanguineous marriages, with absent or conical lower deciduous incisors and thin hair and finger nails, with cleft palate in one and a branchial cyst in the other, were described by Fried,\textsuperscript{24} who suggested autosomal recessive inheritance. A similar patient from Turkey, a child of first cousins, has also been described.\textsuperscript{25} The Norwegian sibs reported\textsuperscript{26} could also well have the autosomal recessive condition described here. A brother and sister from Lebanon with only a few permanent teeth and thin nails were recently reported\textsuperscript{27} with the suggestion that they may have the condition described by Fried.\textsuperscript{24} The pattern of missing teeth, including deciduous teeth, can also be similar in incontinentia pigmenti\textsuperscript{28} and in Kabuki syndrome.\textsuperscript{29}

Congenitally missing deciduous teeth, as seen in 53% of the present patients, is not common in IPH. The prevalence figures are commonly close to 0.5%\textsuperscript{30–32} and, interestingly, are higher (0.9%) in Finland.\textsuperscript{33} All these figures may also include children with syndromic hypodontia or oligodontia. The genetics of hypodontia in the deciduous dentition has not been systematically studied, but has been assumed to be a symptom of IPH, as the corresponding permanent tooth is also usually missing. Detailed descriptions of the distribution of missing teeth in hypodontia patients are rare. However, in two Scandinavian studies, the pattern of missing lower and upper lateral incisors typical of RIH is evident.\textsuperscript{16,34} This also suggests that the condition also exists in other populations but has perhaps been overlooked because of overlapping symptoms with IPH. Retarded dental development and short rooted (taurodontic) molar tooth form are characteristic of hypodontia and oligodontia and were clearly seen in the condition described here.

The number of cases of atopic diseases, commonly seen in our patients and also reported by the family members, exceeds the reported population prevalences\textsuperscript{35} and is an interesting finding in the present context. Of the similar published conditions, only the Norwegian report mentions asthma as a symptom of the patient. Recently, a distinct anhidrotic ectodermal syndrome with missing teeth and immunological abnormalities, EDA-ID, has been described. The condition seems to be caused by impaired NF-κB signalling.\textsuperscript{36}

In recessively inherited conditions heterozygous manifestations may occur. Here, minor dental anomalies were noted in less than half of
recessive mode of inheritance, characterised by missing deciduous and permanent incisors, and an increased inclination to eczema and asthma. We have named the condition Recessive Incisor Hypodontia (RIH). The patients resemble reported patients from consanguinous marriages from various parts of the world. Attempts to clarify the molecular pathology of this condition are at present being carried out and the results will be of interest in developmental studies and in the study of molecular mechanics of atopic diseases.

We warmly thank our patients and the numerous families for making this study possible. We thank dentists throughout Finland, in particular Professor Satu Alaluusua for informing us about patients with missing lower incisors. We thank Professors Reijo Norio and Leena Palotie-Peltonen for valuable advice and support. The study was supported by the Foundation of Paediatric Research, the Ulla Hjelt Fund, and the Academy of Finland.

EDITOR—Congenital diaphragmatic hernia (CDH) is seen in 1/2000 to 1/5000 fetuses and liveborn infants. 1 2 Around 60% of fetuses diagnosed by antenatal ultrasound scanning at 20 weeks' gestation die in utero and the mortality rate in those surviving to term remains 30–50%. Coexistent major structural malformations are seen in a large proportion of cases, the commonest in liveborn infants being cardiac anomalies and neural tube defects. 3 The genetic contribution to the aetiology of CDH is poorly understood. Although no large scale, population based, offspring recurrence study exists, familial clustering of CDH has been attributed to polygenic inheritance, which predicts an offspring recurrence risk of 1–2%. Familial congenital diaphragmatic hernia is, however, well described with autosomal dominant inheritance in most reported families, although no linkage studies have been performed. Candidate genes may therefore be localised by studying the large proportion of patients with CDH and an underlying chromosome abnormality. Autosomal trisomies, typically of chromosomes 13, 18, and 21, are reported in some series, although many of these cases have additional organ malformations. 4–6 However, a number of de novo structural anomalies associated with CDH have been documented, defining candidate loci for future study; these are summarised in table 1.

### Table 1 Candidate loci for congenital diaphragmatic hernia

<table>
<thead>
<tr>
<th>Locus</th>
<th>Reported anomaly</th>
<th>Other phenotypic features</th>
<th>Reference</th>
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<tbody>
<tr>
<td>1q</td>
<td>Del(1)(q32q42)de novo</td>
<td>Low set ears</td>
<td>17</td>
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<tr>
<td></td>
<td>Del(3)(q11.1q13.2)de novo</td>
<td>No</td>
<td>37</td>
</tr>
<tr>
<td></td>
<td>Del(3)(q21q23)de novo</td>
<td>Bilateral equinovarus</td>
<td>16</td>
</tr>
<tr>
<td></td>
<td>Del(4)(p16)de novo</td>
<td>BPES</td>
<td>6</td>
</tr>
<tr>
<td></td>
<td>Del(8)(p23.1)de novo</td>
<td>Facial dysmorphism</td>
<td>18</td>
</tr>
<tr>
<td></td>
<td>Del(8)(p23.1)de novo</td>
<td>Facial dysmorphism</td>
<td>18</td>
</tr>
<tr>
<td></td>
<td>Del(15)(q24qter)</td>
<td>Tricuspid stenosis</td>
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<td></td>
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<td>7</td>
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<tr>
<td></td>
<td>12p</td>
<td>Mosaic tetrasomy 12p</td>
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</tr>
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</tr>
<tr>
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</tbody>
</table>

Case report

We report a male infant with CDH associated with a proximal deletion of the long arm of chromosome 3 in mosaic form. Clinical genetic evaluation was sought during the third pregnancy of a 29 year old female and a 32 year old male. She had previously had two healthy children and there was no history of pregnancy loss. Fetal ultrasound examination at 22 weeks' gestation showed a large, left sided diaphragmatic hernia with mediastinal shift but no hydrops. No other structural abnormality was seen. A placental biopsy was taken for cytogenetic analysis. Both direct and long term preparations showed an abnormal mosaic male karyotype with an additional, unidentified small chromosome in approximately 50% of cells examined. Analysis of cultured amnioncytes confirmed the marker chromosome in 50% of cells. Fluorescence in situ hybridisation (FISH) studies indicated that the marker was derived from the centromeric region of chromosome 3 (fig 1A) and contained euchromatic material. The karyotype was assigned as...