Prenatal detection of short arm deletion and isochromosome 18 formation investigated by molecular techniques

Mazin B Qumsiyeh, Ana Tomasi, Mark Taslimi

Abstract
A patient was referred for amniocentesis because of advanced maternal age and polyhydramnios. The fetal karyotype was a mosaic 46,XX,del(18)(p11.1)/46,XX,−18, +i(18q)de novo. The deletion appeared to encompass the whole short arm as evidenced by G banding and in situ hybridisation. However, telomere sequences were found on both ends of the deleted chromosome as well as the isochromosome. The normal 18 and the isochromosome showed more alphoid sequences than the del(18). Subsequent passages of the cell lines showed an increase in the frequency of the isochromosome from 20% to about 30%. Possible mechanisms are discussed.

Case report
The patient was referred for cytogenetic testing because of advanced maternal age (42 years) and polyhydramnios. This was the fifth pregnancy in this patient. The first pregnancy ended in a stillborn male with breech presentation and placenta praevia (no other information was available). Three other pregnancies were apparently without complications and no phenotypic abnormalities were reported in these children (ages 6, 15, and 22 years). Polyhydramnios was observed at a gestational age of 25 weeks and an amniocentesis was performed (the patient had refused earlier prenatal diagnosis). Amniotic fluid alphafetoprotein was 6·2 and a screen for acetylcholine esterase was negative. Cytogenetic evaluation from amniotic fluid cells showed 16 cells with a 46,XY,del(18)(pcen or p11.1) and four cells with 46,XY,−18,+i(18q). Normal cells were not seen in any of the three coverslip cultures and two flask cultures examined. The patient was counselled that the child at birth could have a phenotype consistent with both non-mosaic deletion 18p and mosaic trisomy 18q. Parental karyotypes were normal. The patient elected to have a therapeutic abortion which was performed at another centre. The fetus was not available for our examination. However, photographs (fig 1) obtained after delivery were sent to us and they showed the following phenotypic abnormalities: brachycephaly, low hair line, prominent occiput, hypertelorism, low nasal bridge, broad nose, full lips with down turning corners, microgнатia, slightly low set ears, small pelvis, short fingers, broad hand, and mild club feet.

Methods
Fluorescence in situ hybridisation was performed using published methods with agents and probes from Oncor, Inc (Gaithersburg, MD). The probes used were an oligonucleotide repeat representing human telomeres and a centromeric probe for chromosome 18 (D18Z1). Biotin labelled probes were hybridised and detected according to the protocol recommended by the supplier (Oncor, Inc). Chromosome preparations were observed under a Zeiss Fluorescence Axioskop microscope with HBO 50W DC mercury lamp and filter combinations of BP 450-490, FT 50, and LP 52. Photographs were taken on ASA 400 Kodak gold film.
Results and discussion
Initial amniotic fluid cultures showed the presence of a mosaic 46,XX,del(18)(p11.1)/46,XX,−18, +i(18q). In addition to the 20 cells initially examined from three coverslips and two flasks, 50 other cells were evaluated from subsequent harvest of three flasks each. Each showed the presence of the mosaic condition with isochromosomes found in 12, 15, and 20% of the cells examined from these three flasks. The culture with the 20% mosaicism was subcultured and then harvested two weeks after the first harvest. Results from this harvest showed 14 of 47 cells examined with the isochromosome, a frequency of about 30%. The increase in the frequency of the isochromosome could indicate that this form of the aberration is more stable than the deletion or is selected for in culture. Thus, we speculated initially that the deletion was terminal with a loss of the telomere sequences. We performed in situ hybridisation with both telomere and centromere probes in these cells and the results did not agree with this simple explanation.

In situ hybridisation with the D18Z1 probe showed weaker hybridisation on the centromeric region of the deleted 18 than on the normal homologous chromosome 18 (fig 2). The isochromosome showed more centromeric hybridisation (fig 2) and C banding (data not shown) than the deleted chromosome 18. The formation of the isochromosome was not accompanied by a retention of the telomere interstitially; none of the 15 cells examined showed a fluorescent signal at the centre of the isochromosome (fig 2). Hybridisation with the telomeric probe showed the deleted chromosome 18 to possess telomeric sequences at both ends of this apparent telocentric chromosome (fig 2). Studies of the hybridisation of alphoid sequences and of C band polymorphisms failed to show the parental origin of the rearranged 18. The parents refused to supply blood for additional molecular studies. Vorsanova et al. reported a case in which amniotic fluid cells showed a short arm deletion of chromosome 18. In situ hybridisation with a radioactive alphoid probe also illustrated less hybridisation (as measured by the number of silver grains) on the deleted chromosome 18 than on the normal 18. These authors did not report the presence of any isochromosomes in these cultures. The phenotype appears milder than the case reported by van Essen et al. with holoprosencephaly and DiGeorge anomalies. Gudsen et al., on the other hand, reported a case very similar to ours and that of Fryns et al. of a deletion 13p11 and formation of isochromosome 13. This phenomenon of deletion/isochromosome formation may thus be more prevalent than initially anticipated.

The following hypothesis can explain these observations. The initial event could have been a destabilised alphoid sequence on 18 with deletion of sequences between the telomere and the centromere. This event resulted in a deleted 18p with reduced centromeric alphoid sequences but a functional telomere remained. This reduction in centromeric alphoid sequences is not a new phenomenon and earlier observations have been reported. The formation of the isochromosome (fig 2) could have been a subsequent event because it appears to have more centromeric material than the deleted chromosome. This is similar to what was reported for chromosome 17 short arm deletion and isochromosome formation. This hypothesis also fits the observation of homologous sequences at telomeres and centromeres of Drosophila and the model of a repeat unit for kinetochore DNA. An alternative hypothesis is that the isochromosome originated first followed by breakage and loss of
of the arms and a regeneration of a telomere. While all these steps are possible, the hypothesis is less parsimonious (that is, requires more steps). The postulated fragility of the alphoid sequences on 18 might also explain other cases reported, such as the patients reported with complementary isochromosomes of both the long and short arms with the karyotype of 47,XY or XX, -18, +i(18p), +i(18q); or the increased incidence of chromosome 18 whole arm translocations and isochromosome formation. More studies of the centromeres of specific chromosomes, and at the subtelomeres will help to further our knowledge of the mechanisms of these chromosome aberrations.

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