# GENE DOSAGE EFFECTS FOR COAGULATION FACTORS XII (F12) AND XIII SUBUNIT A (F13A) IN A CASE WITH PARTIAL MONOSOMY 6p RESULTING FROM A MATERNAL PERICENTRIC INVERSION OF CHROMOSOME 6

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Summary Genes encoding blood coagulation factors XII (F12) and XIII subunit A (F13A) have been shown to be localized in the distal 6p region. We studied gene dosage effects for F12 and F13A in a female case with a duplication-deficiency (partial monosomy for  $6p2307 \rightarrow pter$  and trisomy for  $6q25.1 \rightarrow qter$ ) due to a maternal inversion of chromosome 6. Plasma levels of factor XII were reduced in the patient (46%) and her mother (42%), while that in her father (56%) was within the normal range. On the other hand, plasma levels of factor XIII were normal in the three individuals and the phenotype of factor XIII subunit A was type 1 in all the three. The low factor XII values in the patient and the mother can be best explained by a duplex gene dosage for F12 but with the involvement of a variant gene which results in F12 with a low activity and exists in the two individuals. These results suggested that the loci encoding F12 and F13A can be excluded from the region  $6p2307 \rightarrow pter$ .

# INTRODUCTION

Blood clotting factor XII (FXII) is activated by contact with foreign surfaces, and initiates the intrinsic pathway of coagulation. The localization of the gene encoding this protein (F12) in the distal 6p region has first been indicated by de Grouchy *et al.* (1968) and been confirmed by Pearson *et al.* (1982). In human gene mapping 8 (HGM 8), the locus is assigned to 6p23. The activated form of factor XIII (FXIII) acts in the common pathway of coagulation where it stabilizes covalent

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bonds within fibrin strands. This protein consists of two pairs of subunits, *i.e.*, two alpha chains (FXIIIA) and two beta chains (FXIIIB). The detection of genetic polymorphism for FXIIIA (Board, 1979) has facilitated linkage studies with other various genetic markers, and the close linkage between F13A and HLA has been established (Eiberg *et al.*, 1984; Olaisen *et al.*, 1985). In HGM 8, its recombination fraction in males ( $\theta$ ) is reported to be 0.19. Although the loci, F12 and F13A, are assumed to form a coagulation gene cluster in the 6p23 region, their detailed localization remains to be determined at subband levels. We studied gene dosage effects for F12 and F13A in a case with a duplication-deficiency (trisomy for 6q25.1  $\rightarrow$  qter and monosomy for 6p2307  $\rightarrow$  pter) resulting from a maternal pericentric inversion of chromosome 6. The results excluded the two loci from the region 6p2307  $\rightarrow$  pter.

#### MATERIAL AND METHOD

*Case report.* The proband, a female infant, was born after an uneventful 40 weeks' gestation to unrelated parents (both 29 years of age). An elder sibling was healthy. There was no family history for mental retardation or multiple congenital anomalies. The birth weight was 2,120 g, and the length 45.2 cm. Immediately after birth, the infant was found to have brachymicrocephaly with high forehead, hypertelorism, antimongoloid eye slantings, exophthalmos, flat nasal bridge, small anteverted nose, long philtrum, full cheeks, low-set and malformed ears with folded helices and prominent antihelices, short neck with anterior as well as posterior webbings, heart murmurs (VSD and PH), pigeon breast, left-sided clubfoot and flexion contractures of all fingers. The subsequent clinical course was marked by episodes of cardiac failure, which were treated with digitalis and diuretics. The infant fed poorly and underwent permanent gavage feedings. Her psychomotor and somatic developments were grossly retarded.

Cytogenetic and genetic marker studies. Chromosome analysis was carried out on peripheral blood lymphocyte cultures from the patient, her parents and sibling. Prophasic or prometaphasic cells were collected according to the method of Ikeuchi and Sasaki (1979), and chromosome preparations were processed by GTG banding.

HLA-A, -B, -C, and -DR typing of peripheral blood lymphocytes was based on a standard dye-exclusion microcytotoxicity test. Glyoxalase of red blood cells (GLO1) was separated on a starch gel, using the method of Parr *et al.* (1977). FXII and FXIII were assayed by the method of Sirridge and Shanon (1983) and the caseindansyl cadaverine method (Iatro-FL FXIII Kit, Iatron Laboratories, Inc.), respectively. Phenotypes of FXIIIA were determined, using agarose gel electrophoresis as described by Nishigaki *et al.* (1981).

## RESULTS

### Cytogenetic studies

Banding analysis of the patient showed that the distal portion of the short arm of one chromosome 6 was stained differently from the equivalent portion of the normal homologue. Her mother was found to have a pericentric inversion of chromosome 6. Analysis of prophasic cells at the level of 850 bands per haploid set revealed the locations of breakpoints at 6p2307 and 6q25.1 (Fig. 1a). A meiotic crossing-over between the inverted and normal chromosomes 6 of the mother appeared to have yielded the recombinant chromosome 6 seen in the patient, where the segment 6q25.1  $\rightarrow$  qter was duplicated and the segment 6p2307  $\rightarrow$  pter was deleted (Fig. 1b). The patient's karyotype was therefore 46,XX,rec(6),dup q,del p,inv(6) (p2307q25.1)mat. Her father and sibling had a normal G-banded karyotype.

#### Genetic marker studies

Table 1 showed the results of genetic marker studies. None of HLA-A, -B, -C or -DR loci were deleted in the patient. Phenotypes of GLO1 from the patient and her parents were all type 2. The levels of FXII were markedly reduced in both the patient (46%) and the mother (42%), while that was within the normal range in



Fig. 1. Partial G-banded karyotypes of the patient (b) and the mother (a). Arrows indicate breakpoints of the inversion.

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	Patient	Father	Mother
HLA	A2,A26,BW35,B(-), CW3,CW7,DR1,DR4	A26,AW24,BW62,B(), CW3,CW7,DR1,DR1	A2,AW24,BW35,BW51, CW3,C(-),DR4,DRW8
GLO1	Type 2	Type 2	Type 2
FXII (normal, 50–150%)	46% <sup>a</sup>	56% <sup>a</sup>	42% ª
FXIII (normal, 50–150%)	80%	94%	81%
FXIIIA	Type 1	Type 1	Type 1

Table 1. Results of gene marker studies.

<sup>a</sup> Means of two separate measurements.



Fig. 2. Agarose gel electrophoresis of factor XIIIA from the patient (C), father (P) and mother (M).

the father  $(56^{\circ}_{0})$ . In all the three, on the other hand, levels of FXIII were normal. Phenotypes of FXIIIA were all type 1 (Fig. 2).

## DISCUSSION

Figure 3 summarizes physical and genetic mapping data for chromosome 6 loci presented in HGM 8. The shortest region of overlapping (SRO) for F12, HLA, GLO1 and PGM3 has been assigned to 6p23, 6p21.3, 6p21.31  $\rightarrow$  p21.2 and 6q12, respectively. On the basis of male recombination fractions, the gene order pter-ME2 (mitochondrial malic enzyme)-F13A-HLA-GLO1-PGM3 has been proposed. The genetic distance between the loci for ME2 and F13A is found to be 10 centimorgan (cM), that for F13A and HLA 19 cM, that for HLA and GLO1 4 cM, and that for GLO1 and PGM3 15 cM. As regards ME2 and F13A, however, there has been no evidence for physical assignment nor deletion mapping to the distal 6p region.

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Fig. 3. Physical and genetic mapping data for chromosome 6 loci (Lam and Olaisen, 1985). As regards F12, a solid bar represents a segment to which the locus is assigned, and an open bar a segment from which the locus is excluded: a, de Grouchy et al. (1968), "?" indicates undescribed breakpoints; b, Pearson et al. (1982); and c, Niebuhr et al. (1985).

The present study has shown markedly reduced levels of FXII not only in the patient with a duplication-deficiency (monosomy for  $6p2307 \rightarrow pter$  and trisomy for  $6q25.1 \rightarrow qter$ ) but also in the mother with a pericentric inversion of chromosome 6. This confounding result can be explained in three ways. First, the locus, encoding F12 lies within the region  $6p2307 \rightarrow p2400$ , and the patient, a duplicationdeficiency recombinant, is hemizygous for this locus. The involvement of a variant gene which results in F12 with a low activity and exists in the mother would be a prerequisite for this supposition. Second, the locus is located just on the breakpoint of the inversion, 6p2307. As a result of a gene disruption or a submicroscopic deletion of the locus for F12 during the process of the inversion, both the patient and the mother would become hemizygous for F12. Third, the locus exists proximal to the breakpoint. In this case, the patient should be heterozygous for a common F12 allele and a variant allele which results in F12 with a low activity and is inherited from the mother. In the former two explanations the FXII level of the patient depends only on that of the father, while in the third explanation the level is influenced by that of the mother as well as the father. The similarity of FXII level of the patient to the mean parental value is most in accordance with the third explanation, suggesting the exclusion of the F12 locus from the region  $6p2307 \rightarrow pter$ .

Niebuhr *et al.* (1985) have described a normal F13A level in a case with partial trisomy  $2p(p24 \rightarrow pter)$  and partial monosomy  $6p(p24 \rightarrow pter)$  derived from the paternal balanced translocation, t(2;6) (p24;p24). Our study has also showed the

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normal F13A level in the patient with monosomy for  $6p2307 \rightarrow pter$ . Although the two studies failed to substantiate the localization of F13A in the distal 6p region by deletion mapping, in view of the genetic distance between the loci for F13A and HLA (its SRO=6p21.3) in males whose chiasma map distance is reported to be 55 cM of length (Morton *et al.*, 1982), we would like to assume that the loci encoding F12 and F13A may form a coagulation factor gene cluster in the region  $6p2300 \rightarrow p2307$  as do the loci encoding F7 and F10 in the 13q34 region (Pfeiffer *et al.*, 1982). The confirmation of this hypothesis will have to await gene dosage studies in further cases with partial monosomy 6p or *in situ* hybridization studies using a complementary DNA for F12 or F13A.

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