



X 가

1/28,000-38,000

X p22 3) X

2) X X 가 가

DNA (low-copy repetitive DNA segment) (nonhomologous pairing) (azoospermia)

3) 3 X 가

Xp X (critical region) Xq13-22 q22-26 (gonadal dysfunction) 3, 4) X 40

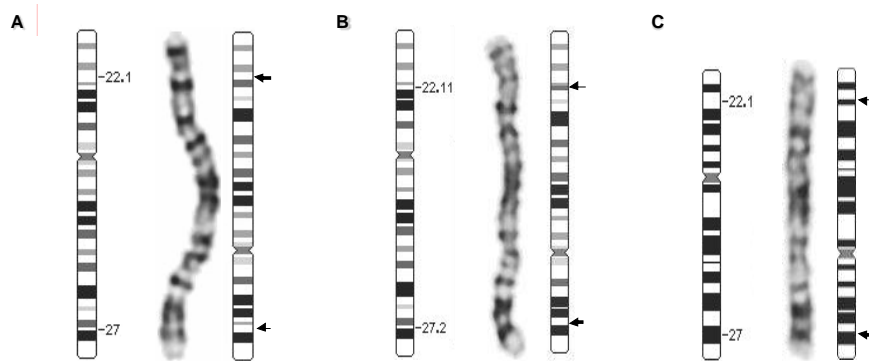
q27 SOX3(sex-determining region Y box3) 가 sY84, sY129, sY134, sY254, sY255, SRY Y 1

5) SHOX(short stature homeo- pseudo-autosomal region(PAR) (chondrocyte) (spermatocyte), 2 가

homeodomain 6) SHOX (point mutation) LH 5.1 ( 1.5-9.2 mIU/mL), FSH 20.0 ( 1-14 mIU/mL), prolactin 4.8 ( 0-15 ng/mL), estradiol 19.0 ( 15-80 pg/mL), testosterone 3.7 ( 1.31-8.13 ng/ml) FSH가

가 (disruption) 6) X GTG-

(random inactivation) (skewed X inactivation) 가 X 20 /



**Fig. 1.** Ideograms (normal X on the left, inv(X) on the right) and GTL-banded pericentric inversion X chromosome (middle) for each of the two cases. Arrows show breakpoints. (A) Case 1: A 38-year-old male carrier with azoospermia; 46,Y,inv(X)(p22.1q27). (B) Case 2: A 38-year-old pregnant woman with a family history; 46,X,inv(X)(p22.11q27.2). The patient had an uneventful pregnancy at term. (C) The female fetus of case 2; 46,X,inv(X)(p22.1q27).

. International System for Human Cytogenetic Nomenclature (ISCN) 550-

46,X,inv(X)(p22.11q27.2),  
46,Y,inv(X)(p22.11q27.2)  
. X

(Fig. 2 I-1)

X  
46,Y,inv(X)(p22.1q27)

가  
(Fig. 1A).

(0.001M) 6 가

3 BrdU

2

33258(0.5ug/mL) 30 , 2XSSC  
, 55 2XSSC 2

Hoechst

2

Giemsa

118

1

1

가 38

RBG-

118

17 X  
(Fig. 2 II-6)  
가

가 38  
(Fig. 2 III-5)

가 36 : 82 (30.5% : 69.5%)  
가 (Fig. 3).

X : inv(X)  
X

X : inv(X)가 118 : 5 (95.9% : 4.1%)

X

가

GTG-

20

ISCN

700-

118

46,X,inv(X)(p22.11q27.2) (Fig. 1B)

*in-situ*

X

550-

37

46,X,inv(X)

(p22.1q27) (Fig. 1C)

가

,

2005

<sup>3)</sup> Wenger

<sup>2)</sup>

39

(Fig. 2 III-3)

1/28,000-38,000

, Schorderet

<sup>3)</sup> X

46,Y,?inv(X)

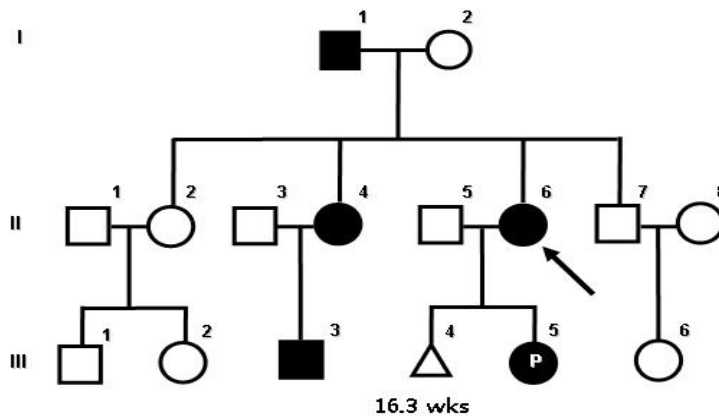
, 25 가

12

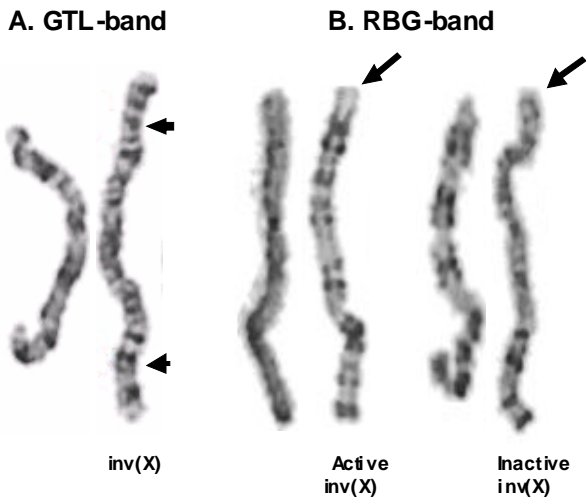
Xp

p22

(Fig. 2. II-4)



**Fig. 2.** Pedigree of the patient in case 2. All carriers had a normal phenotype and exhibited normal fertility. The family exhibited balanced pericentric inversion of the X chromosome. The karyotype of each family member was as follows: I-1: 46,Y,inv(X)(p22.11q27.2); II-4: 46,X,inv(X)(p22.11q27.2); II-6(patient): 46,X, inv(X)(p22.11q27.2); III-3: 46,Y,?inv(X); III-5: 46,X,inv(X)(p22.1q27). The arrow shows the pregnant woman in case 2.



**Fig. 3.** Partial karyotype of case 2; 46,X,inv(X)(p22.1q27.2). (A) The normal X chromosome and pericentric inversion of X chromosome of a 850-band level ideogram. (B) Arrows show breakpoints on a GTL-banded inv(X). (C) Inactivation pattern of normal X and the inv(X). The proband had a 69.5% incidence of inv(X) late replication. Arrows indicate each active/inactive inv(X).

Xp22.3 (low-copy number repeat sequences) contiguous gene syndrome  
 7). Xp  
 (chondrodysplasia punctata), steroid sulphate deficiency, Kallmann 8). X  
 Xq26.3-qter (disomy)  
 가  
 (profound developmental delay), (hypotonia), (microcephaly), (corpus callosum agenesis), (dysmorphic facial features), (left multicystic, dysplastic kidney) 8).  
 X  
 (positional effect)  
 가 가  
 9). 25  
 X

8)  
 46,Y,inv(X)(p11.4q11.2) 9)  
 inv(X)  
 가 X 가  
 (spermatogenesis) Xp-Yp  
 PAR1 (pseudoautosomal region 1)  
 X-Y  
 Xq27.1 SOX3 가  
 가 1  
 1, 2  
 FSH 가 SOX3  
 FSH 가 SOX3  
 , X  
 (hypopituitarism) 5).  
 Bleyl 6) Xq27.1  
 X  
 (corneal opacity), 가 (adherent iris strands), (keratolenticular cataract) (Peter anomaly)  
 Xq27 SOX3 가  
 Xq13 (X-inactivation center, XIC) XIST(X-inactivation specific transcript) X  
 X inv(X)가  
 , X 가 X  
 X



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