

## StyI polymorphism at the D15S11 locus

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**Source and Description of Probe:** pIR4-3R, a 0.4 kb fragment subcloned into pUC19 (1, 2), originally isolated from a flow-sorted library enriched for proximal 15q sequences.

**Polymorphism:** StyI reveals a four allele polymorphism with band sizes 3.4 kb, 2.4 kb, 1.9 kb, 0.9 kb, and a constant fragment at 2.2 kb.

**Allele Frequency:** Estimated from 28 unrelated Japanese.  
 3.4 kb allele: 0.33  
 2.4 kb allele: 0.42  
 1.9 kb allele: 0.06  
 0.9 kb allele: 0.19

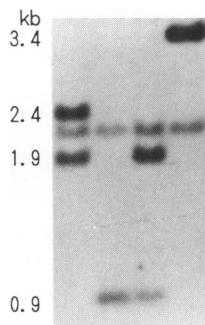
**Not Polymorphic For:** BanII, HaeIII, HincII, NciI, NsiI.

**Chromosomal Localization:** pIR4-3R has been assigned to 15q11.2-15q12, a segment missing in one #15 chromosome of a Prader-Willi syndrome patient with a prominent cytological deletion (1, 2).

**Mendelian Inheritance:** Codominant segregation observed in 8 families with Prader-Willi syndrome patients.

**Probe Availability:** Available from ATCC.

**References:** Donlon, T.A. *et al.* (1986) *Proc. Natl. Acad. Sci. USA* **83**, 4408-4412. 2) Tantravahi, U. *et al.* (1989) *Am. J. Med. Genet.* **33**, 78-87.



## PstI RFLP of the CGB gene

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**Source and Description of Clone:** pCGβ474 is a 580 bp cDNA clone of the beta sub-unit of the human chorionic gonadotrophin gene (1).

**Polymorphism:** PstI identifies two alleles:

A1-absence of 3.8 kb fragment

A2-presence of 3.8 kb fragment

Constant bands also at 3.0, 2.3, 1.7, 1.4, 1.2, 0.66 and 0.42 kb.

**Frequencies:** 14/28 unrelated Caucasians were of the 1,1 genotype.

Assuming Hardy-Weinberg equilibrium: A1 (-3.8 kb): 0.71  
 A2 (+3.8 kb): 0.29

**Not Polymorphic For:** EcoRI, HindIII, MspI, TaqI in 18 unrelated Caucasians.

**Chromosomal Localization:** CGβ and the beta chain of luteinizing hormone are part of a gene cluster at 19q13 (2).

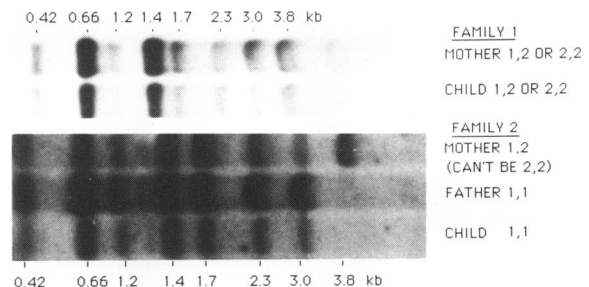
**Mendelian Inheritance:** Mendelian inheritance was demonstrated in two two-generation families (Figure).

**Probe Availability:** Contact Dr Boime, Department of Pharmacology, Washington University Medical School, St. Louis, MO 63115.

**Other Comments:** 2,2 homozygotes cannot be distinguished from 1,2 heterozygotes even if the pCGβ474 plasmid is cut down to the 80 bp XmnI-HincII fragment which only detects the variable 3.8 kb fragment and the constant bands at 0.66 and 0.42 kb. This suggests that the RFLP may be due to an insertion/deletion.

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**References:** 1) Policastro *et al.* (1983) *J. Biol. Chem.* **258**, 11492-11499. 2) Brook *et al.* (1985) *Cytogenet. Cell Genet.* **40**, 590-591.



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