

ERRATUM

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**A *Tsp509I* variant in exon 13
of the neurofibromatosis type 1 (NF1) gene
allows the identification of both alleles at the mRNA level**

Hum Genet (1995) 96:131–132

An unfortunate error occurred in the presentation of the alternative bases in the *Tsp509I* polymorphism. The residues at position 2034, reported as C and A, were actually G and A. The correct version reads:

The residues found at position 2034 are G (frequency: 0.71) and A (frequency: 0.29). Both codons (CCG and CCA) still encode proline.