

## *itga5*<sup>b926</sup>

### Nature of the mutation

The *b926* allele contains a single T-to-A point mutation that substitutes tyrosine for asparagine at amino acid 218 (Crump et al., PloS Biol. 2(9): E244, 2004).

### Genotyping assay

Genotyping of the *b926* allele is based on the RFLP assay (Restriction Fragment Length Polymorphism; Botstein *et al.*, Am. J. Hum. Genet. 32: 314-331, 1980). This method is used to detect a mutation that either creates or abolishes a site recognized by a specific restriction enzyme. In the RFLP assay, a sequence of interest is first PCR-amplified and then the PCR product is subjected to restriction enzyme digestion. The presence or absence of the mutation is determined by the resulting restriction pattern. The *b926* mutation abolishes a site recognized by the BciVI restriction enzyme.

### **Primers:**

**ITG\_03:** 5' GTG ACC TTC AGC TCA ATG TAA ACG 3'

**ITG\_04:** 5' CTC ACC TTC ATC ATC ATC ACC G 3'

### **PCR program (55\_30\_30):**

1. 94°C for 3 min
2. 94°C for 30 sec
3. 55°C for 30 sec
4. 72°C for 30 sec
5. Go to step 2 (above) for 39 cycles
6. 72°C for 5 min
7. 8.0°C hold
8. END

### **Product size: 324 bp**

### **Digestion of the PCR product with the BciVI restriction enzyme:**

Product type	Product digestion	DNA fragments after digestion (bp)
PCR product derived from the WT template	cleaved	221 bp and 103 bp
PCR product containing the mutation	unaffected	324 bp

**IMPORTANT NOTE:** It is highly recommended to use WT positive controls to monitor whether enzyme digestion has been carried out to completion. Without this control, partially digested WT samples can be mistakenly regarded as heterozygous samples.

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