



ampli set FV H1299R ^{CE IVD} **45 tests**
detection of H1299R polymorphism of the factor V gene.

cat 1315

The heterogeneity of clinic phenotype and the variability of thrombotic events showed by patients with familiarity for thrombotic disease have led to the hypothesis that the predisposition to these type of disorders may be due to many genetic factors. Recently, a complex haplotype of Factor V (HR2), which includes 13 different polymorphisms, has been reported. Among them, 7 cause an amino acid substitution and a functional modification of the protein, leading to an excess of plasmatic isoform FV1 concentration, more thrombogenic. It isn't clear if haplotype HR2 alone could be a factor of thrombotic risk. It is sure that the risk of clinical thrombotic events in subjects carriers of the F V Leiden mutation is increased.

The detection of the H1299R polymorphism is carried out starting with an amplification using specific primers for the exon 13 of a fragment of 703 bp, following by a restriction section due to *Rsa I*. The mutation is confirmed by the gain of a cleavage site for the enzyme *Rsa I*.

Principle of method: A) extraction of genomic DNA; B) amplification; C) enzymatic digestion; D) detection on agarose gel

Applicability: on extracted and purified genomic DNA from whole blood samples.

ANALYSIS OF RESULTS

The yield of amplification is a fragment of 703 bp. The next restriction section made by the *Rsa I* enzyme can be done the following results:

1	2	3
Absence of mutation	Presence of mutation	Presence of mutation
Homozygote	Eterozygote	Homozygous mutant patient
Normal Patient	Mutant patient	
1 fragment	3 fragments	2 fragments
703 bp	703 bp	493 bp
	493 bp	210 bp
	210 bp	

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