

Gene and polymorphism	Region	rs number	Primer pairs	Method	Genotype		
					Wild Ref	Heterozygote	Variant
ER1 (+397 T>C)	intron 1	rs223469 3	(F)5'CTGCCACCCTATCT GTATCTTT-3' (R)5'ACCCTGGCGTCGA TTATCTGA-3'	Pvu II based PCR- RFLP	TT	TC CC	/16/
CALCR (+1377 T>C)	Exon 13	rs180119 7	(F)5'AGGTCCAAACCAC CGTGAAG-3' (R)5'GCAGTGGGAGACT CCATTCC-3'	Alu I based PCR- RFLP	TT	TC CC	/16/