

SUPPLEMENTARY TABLE S2. RESTRICTION FRAGMENT LENGTH POLYMORPHISM CONDITIONS FOR THE IDENTIFICATION OF POLYMORPHISMS GENOTYPED IN THE VDR GENE

<i>rs no.</i>	<i>Location</i>	<i>Alleles</i>	<i>Primers for PCR amplification (5'-3')</i>	<i>Annealing temp. (°C)</i>	<i>PCR product length (bp)</i>	<i>Restriction enzyme</i>	<i>Restriction fragment length (bp)</i>
rs1544410	chr12:48239835	A/G (B/b)	F: GGAGACACAGATAAGGAAATAC R: CCGCAAGAAACCTCAAATAACA	60	248	FspI	A (B) = 248 G (b) = 175+73
rs2228570	chr12:48272895	C/T (F/f)	F: GCACTGACTCTGGCTCTGAC R: ACCCTCCTGCTCCTGGCT	72.5	341	FokI	C (F) = 341 T (f) = 282 + 59

PCR, polymerase chain reaction.