

Supplementary Data

SUPPLEMENTARY TABLE S1. CHARACTERISTICS
OF ANALYZED POLYMORPHISMS OF THE VDR GENE

<i>rs no.</i>	<i>SNP function</i> ^a	<i>Location</i> ^b	<i>Alleles</i>	<i>MAF</i> ^c
rs1544410	Intronic	chr12:48239835	<u>A</u> /G (<u>B</u> /b)	0.31
rs2228570	Missense	chr12:48272895	<u>C</u> / <u>T</u> (<u>F</u> /f)	0.44

Underline denotes the minor allele in the control samples.

^aAccording to the dbSNP.

^bBased on UCSC Human Genome Browser, February 2009 human reference sequence (GRCh37).

^cMAF, minor allele frequency calculated from the control samples.
dbSNP, single-nucleotide polymorphism database; VDR, vitamin D receptor.