

**Suppl Table VI.** Influence of *VDR FokI* rs2228570 gene polymorphism on risk of rheumatoid arthritis

Models	Genotype	Cases, <i>n</i> (%)	Controls, <i>n</i> (%)	<i>P</i> -value <sup>a</sup>	EMP1	OR <sup>c</sup>	95% CI
Genotypic	AA	26 (12.15)	142 (18.98)	0.0613	0.0592	0.76	0.46-1.23
	AG	103 (48.13)	328 (43.85)			1.29	0.94-1.80
	GG	85 (39.72)	278 (37.17)			1	
Dominant	A	129 (60.28)	423 (57.16)	0.4623	0.4141	1.14	0.84-1.55
	GG	85 (39.72)	317 (42.84)				
Recessive	AA	26 (12.15)	127 (17.16)	0.0980	0.0791	0.66	0.41-1.03
	G	188 (87.85)	613 (82.84)				
Allelic	A	155 (36.2)	550 (37.2)	0.7207	0.7338	0.96	0.77-1.20
	G	273 (63.8)	930 (62.8)				
Additive	-	-	-	0.1160 <sup>b</sup>	0.7317	0.96	0.77-1.19

*CI* – confidence interval, *OR* – odds ratio. <sup>a</sup>*P*-value for  $\chi^2$ -test. <sup>b</sup>*P*-value for the Cochran-Armitage test for trend.

<sup>c</sup>*Unadjusted or crude ORs.*