

# Supplementary Materials: SNP-SNP Interaction between TLR4 and MyD88 in Susceptibility to Coronary Artery Disease in the Chinese Han Population

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**Table S1.** Primer sequences and reaction conditions.

SNPs	Primer Sequence	Annealing Temperature	Restriction Endonuclease	Digestive Temperature	Product Size
rs10116253	F: GGGTGTAAAGC CAGGTAGAGGAGGT R: TGGAAAGTAGCAA GTGCAATGTAAGT	63 °C	BsmAI	37 °C	TT: 470 + 131 bp TC: 601 + 470 + 131 bp CC: 233 bp
rs10983755	F: TGCCAGAACATCAA GAACACAGAAAAG R: TGAAAGTAGCA AGTCAATGTAAGT	60 °C	TaiI	65 °C	GG: 342 + 121 bp GA: 463 + 342 + 121 bp AA: 463 bp
rs11536889	F: GCAGGAAGGA AGTGGGATGAC R: TGTTCTGAGG AGGCTGGATG	63 °C	TaaI	65 °C	GG: 391 bp GC: 391 + 292 + 99 bp CC: 292 + 99 bp
rs7744	F: AGGGAGCCTAA CCATGTCCCTGA R: AGGGTCTGT CCTGGGGCACA	61 °C	FspBI	37 °C	AA: 291 bp AG: 291 + 172 + 119 bp GG: 172 + 119 bp

**Table S2.** The genotype frequencies and HWE in this study.

Genotype	Cases	Controls	p Value of HWE	Chr (Chr Position)
TLR4 rs10116253 (T > C)			0.333	9 (117702042)
TT (%)	154 (36.3)	122 (28.8)		
TC (%)	199 (46.9)	220 (51.9)		
CC (%)	71 (16.7)	82 (19.3)		
T allele (%)	507 (59.8)	464 (54.7)		
C allele (%)	341 (40.2)	384 (45.3)		
TLR4 rs10983755 (G > A)			0.186	9 (117702392)
GG (%)	218 (51.4)	189 (44.6)		
GA (%)	170 (40.1)	197 (46.5)		
AA (%)	36 (8.5)	38 (9.0)		
G allele (%)	606 (71.5)	575 (67.8)		
A allele (%)	242 (28.5)	273 (32.2)		
TLR4 rs11536889 (G > C)			0.383	9 (117715853)
GG (%)	255 (60.1)	270 (63.7)		
GC (%)	144 (34.0)	133 (31.4)		
CC (%)	25 (5.9)	21 (5.0)		
G allele (%)	654 (77.1)	673 (79.4)		
C allele (%)	194 (22.9)	175 (20.6)		

<i>MyD88</i> rs7744 (A > G)			0.177	3 (38142530)
AA (%)	157 (37.0)	174 (41.0)		
AG (%)	213 (50.2)	185 (43.6)		
GG (%)	54 (12.7)	65 (15.3)		
A allele (%)	527 (62.1)	533 (62.9)		
G allele (%)	321 (37.9)	315 (37.1)		

HWE: Hardy-Weiberg equilibrium; *TLR4*: toll-like receptor 4; *MyD88*: myeloid differentiation factor 88;  
Chr: chromosome.



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