

Table 1. Supplemental Material. Association between *IL-20* gene polymorphisms and subclinical atherosclerosis.

SNP	Model	Genotypes and Alleles	Women		P	OR	95% CI	Men		P	OR	95% CI			
			SA	Control				SA	Control						
			n	n				n	n						
rs1400986		CC	53	202	0.0009	0.43		124	134	0.087	0.67	0.45-1.00			
		CT	21	187				66	106						
		TT	1	26				9	17						
	codominant1	C	127	591	0.0002	0.44	0.28-0.71	314	374	0.036	0.71	0.52-0.97			
		T	23	239				84	140						
	codominant2	CC	53	202	0.002	0.42	0.24-0.73	124	134	0.048	0.67	0.45-0.99			
		CT	21	187				66	106						
	dominant	CC	53	202	0.036	0.14	0.01-1.10	124	134	0.219	0.57	0.24-1.33			
		TT	1	26				9	17						
	recessive	CC	53	202	0.0004	0.39	0.23-0.67	124	134	0.029	0.66	0.45-0.96			
		CT+TT	22	213				75	123						
	additive	CC+CT	74	389	0.046	0.2	0.03-1.51	190	240	0.33	0.67	0.29-1.53			
TT		1	26	7				17							
rs1518108		-	-	-	0.0002	0.42	0.26-0.68	-	-	0.03	0.71	0.52-0.97			
		CC	25	112				54	69				0.089	1.17	0.75-1.82
		CT	33	220				106	116						
	TT	17	83	39	72										
	codominant1	C	83	444	0.596	0.89	0.63-1.26	214	254	0.204	0.84	0.64-1.09			
		T	67	386				184	260						
	codominant2	CC	25	112	0.181	0.67	0.38-1.18	54	69	0.501	1.16	0.74-1.81			
		CT	33	220				106	116						
	dominant	CC	25	112	0.864	0.91	0.46-1.80	54	69	0.183	0.69	0.40-1.17			
		TT	17	83				39	72						
	recessive	CC	25	25	0.27	0.74	0.44-1.25	54	69	0.95	0.99	0.65-1.50			
		CT+TT	50	303				145	188						
additive	CC+CT	58	332	0.6	1.17	0.65-2.12	160	185	0.036	0.63	0.40-0.98				
	TT	17	83				39	72							
		-	-	-	0.67	0.93	0.65-1.32	-	-	0.2	0.84	0.65-1.09			

Table shows the models with significant association. SA: Subclinical atherosclerosis.