

**Relationship of polymorphisms in the tissue inhibitor of metalloproteinase (*TIMP*)-1
and -2 genes with chronic heart failure**

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Supplementary Table S1. Comparison of genotype and allele frequencies of the 372T>C polymorphism between heart failure patients with and without left bundle branch block (LBBB).

	Without LBBB (n = 209)	With LBBB (n = 84)	P*
Females			
Genotype	n = 63	n = 36	
TT, n (%)	25 (39.7)	14 (38.9)	0.979
TC, n (%)	25 (39.7)	15 (41.7)	
CC, n (%)	13 (20.6)	7 (19.4)	
Allele	n = 126	n = 72	
T, n (%)	75 (59.5)	43 (59.7)	>0.999
C, n (%)	51 (40.5)	29 (40.3)	
Males			
T, n (%)	75 (51.4)	33 (68.7)	0.053
C, n (%)	71 (48.6)	15 (31.3)	

*P-values were calculated using the Pearson chi-square test with Yates correction where appropriate. Left bundle branch block data were available for 293 patients (instead of 300).

Supplementary Table S2. Comparison of genotype and allele frequencies of the 372T>C polymorphism between white and non-white subjects stratified by clinical status.

	Blood donors (n = 304)			Heart failure patients (n = 300)		
	White (n = 238)	Non-white (n = 66)	P*	White (n = 214)	Non-white (n = 86)	P*
Females						
Genotype	n = 76	n = 17		n = 74	n = 25	
TT, n (%)	20 (26.3)	6 (35.3)	0.672	29 (39.2)	10 (40.0)	0.818
TC, n (%)	40 (52.6)	7 (41.2)		29 (39.2)	11 (44.0)	
CC, n (%)	16 (21.1)	4 (23.5)		16 (21.6)	4 (16.0)	
Allele	n = 152	n = 34		n = 148	n = 50	
T, n (%)	80 (52.6)	19 (55.9)	0.878	87 (58.8)	31 (62.0)	0.815
C, n (%)	72 (47.4)	15 (44.1)		61 (41.2)	19 (38.0)	
Males	n = 162	n = 49		n = 140	n = 61	
T, n (%)	87 (53.7)	23 (46.9)	0.504	83 (59.3)	30 (49.2)	0.241
C, n (%)	75 (46.3)	26 (53.1)		57 (40.7)	31 (50.8)	

*P-values were calculated using chi-square test with Yates correction where appropriate.