

Table II. Multiple regression of independent predictors for SR-BI protein levels for the entire population of subjects with HALP.

A. Initial full model ($p=0.04$, $r=0.65$)

Covariates: gender ($p=0.03$), BMI ($p=0.01$), HT use ($p=0.25$), ethnicity ($p=0.12$), aspirin use ($p=0.70$), CETP activity ($p=0.46$), LCAT activity ($p=0.68$), rs4238001 [G2S] ($p=0.07$), rs2278986 [intron 3] ($p=0.05$), and rs5888 [exon 8] ($p=0.65$).

B. Final model ($p=0.001$, $r=0.59$)

1.	BMI	$p=0.008$
2.	rs4238001	$p=0.02$
3.	rs2278986	$p=0.02$