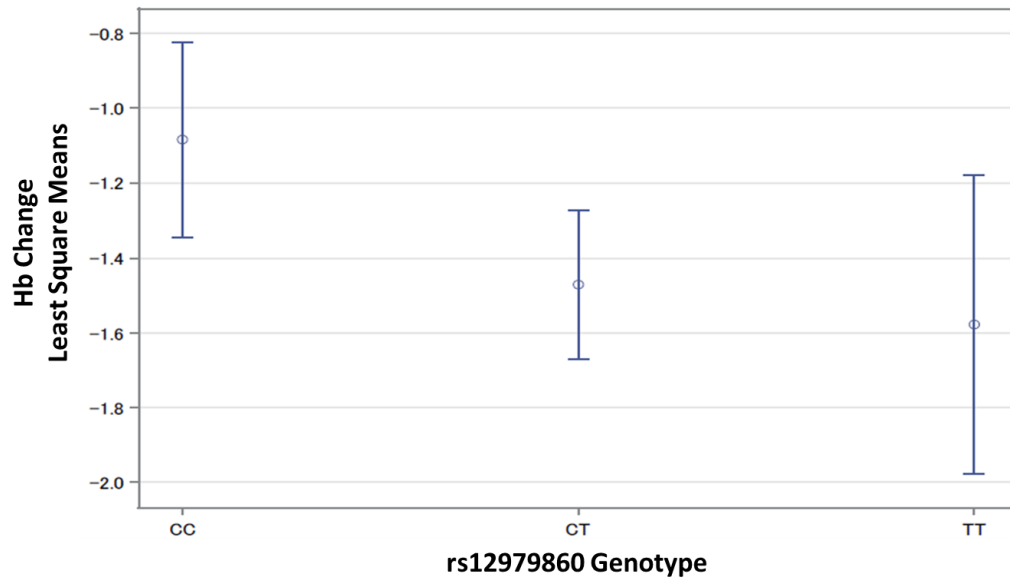


**S2 Fig. Hb Change Least Square means by rs12979860 Genotype**



**S2 Fig.** The least square means were calculated for Hb change using age, sex, arm, ITPase function for the *rs12979860* genotype. Circles indicate the LS means, error bars indicate the 95% confidence interval of the LS-means. We found a larger, significant reduction ( $P=0.0367$ ) in Hb when there was at least one 'T' allele present.