

Supplementary information, Figure S1

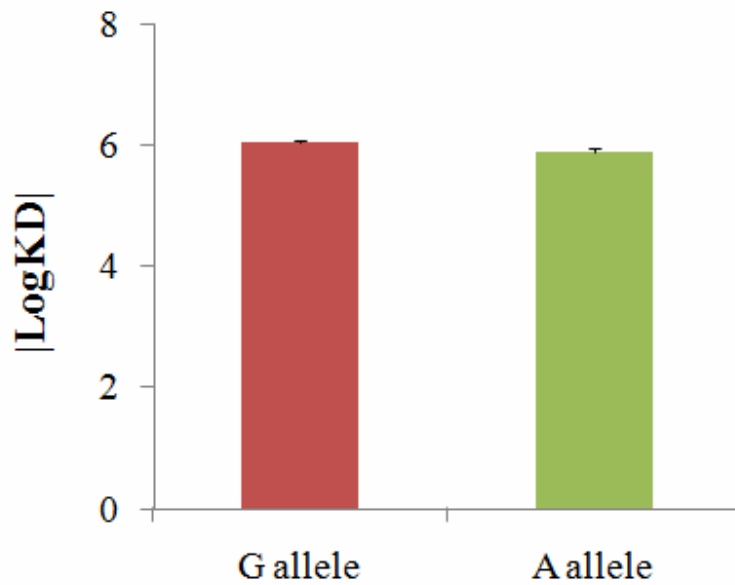


Figure S1 SPR analysis shows that *CBS* -4638G>A does not change the binding affinity with nuclear extracts. Data shown are mean \pm SD. The actual values for each group are as follows: G allele, 6.02 ± 0.06 ; A allele, 5.86 ± 0.14 . All values represent the mean \pm SD from 3 independent experiments.