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 ± 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 ± SD from 3 independent experiments.