Supplementary information, Figure S2

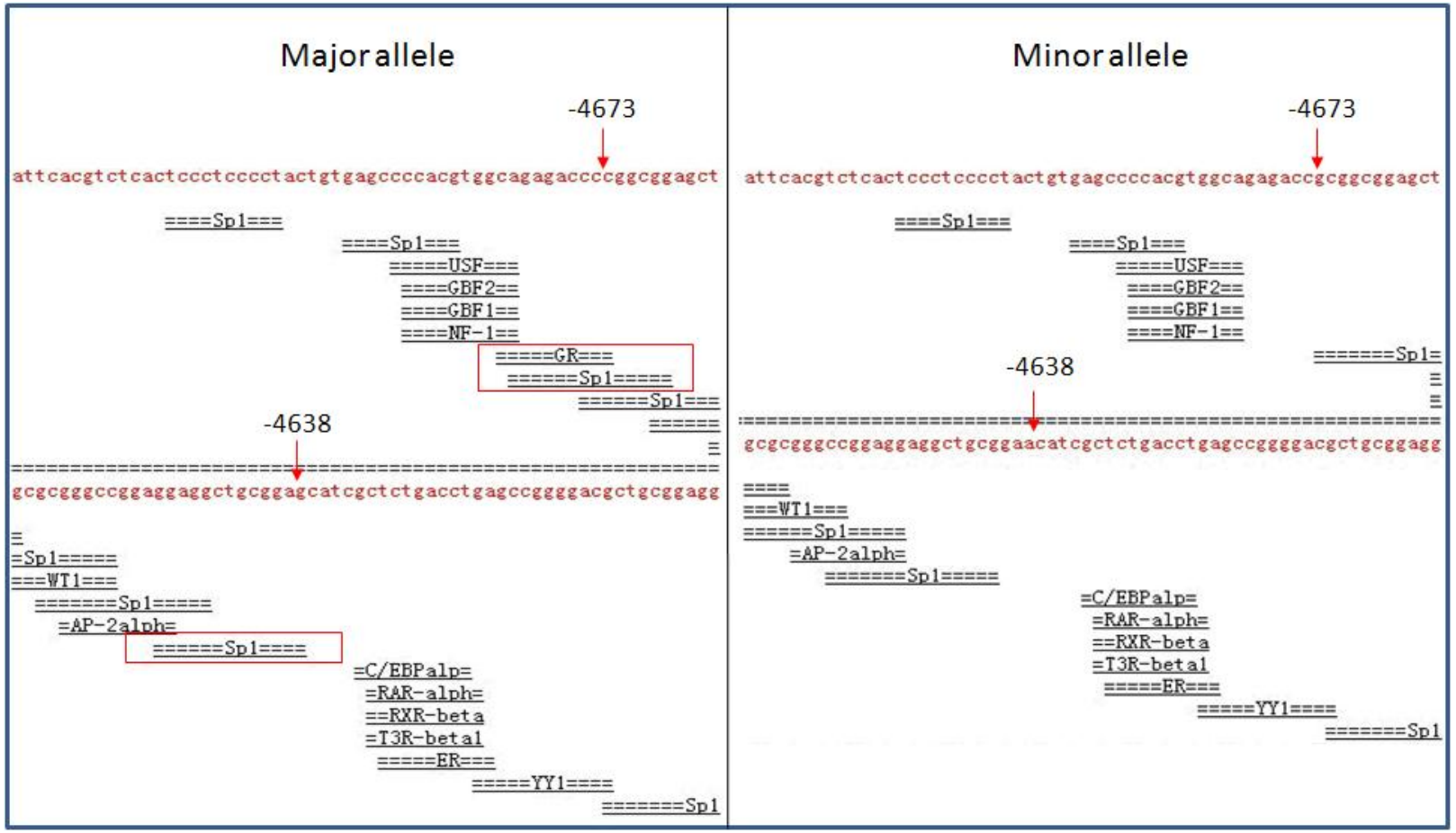

Figure S2 Computational analysis predicted transcription factors for *CBS* variant -4673C>G and variant -4638G>A.