Supplementary Table S6. Predicted E2F1 binding site in the *TBX21* promoter in hTFtarget.

Motif	Score	P value
GGGGTGGAACGG	9.94737	4.15E-05
GAGGCGGAGCAG	9.53947	4.84E-05
CTGCGCGCACCC	9.23684	7.96E-05
GGGGCGAGGCGG	8.15789	8.01E-05
GGGGTGGGGCGA	15.3289	3.35E-06
CGAGAGGGCGGGGTGG	13.8939	8.82E-06
AGGGCGGGGTGG	13.4211	8.69E-06
CGTTTCCC	11.4648	1.21E-05