Transcription factor	Binding motif (PWM)	Enrichment score	p-value
TBX-40	T_CCAA	0.937	0.403
TBX-33	AT-T-CA	1.04	0.895
TBX-43	$_{AG}GTG_{T}$	A 0.5	0.997
TBX-39	CçAAgT	0.141	0.997
TBX-38	g T G.	0.806	0.998