

TF	Pattern name	Source	Sequence name	Start	Stop	Strand	Score	P value	Q value	Matched motif
ZBTB10	m-dataset-3775-1	hTFtarget	hg38_ncbiRefSeq_NM_000188.3	1252	1267	+	10.0921	8.52e-05	0.168	GAGGGAGAGCCACCGG
ZBTB10	m-dataset-3775-1	hTFtarget	hg38_ncbiRefSeq_NM_000188.3	916	931	-	9.96053	9.14e-05	0.168	TAGACAGGTAAGGAAA