

TBR1 HT-SELEX motif:

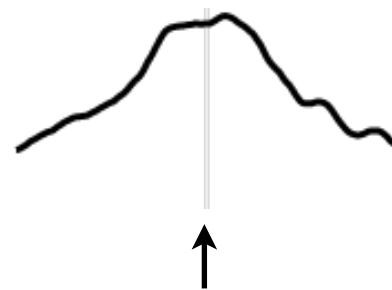


Enriched in our peaks:



E-value:
 $8.3e-101$

Distribution of
motif matches:



ChIP-seq peak summit