

Jump to: Multiple Alignment Motif Tree Motif Matching

Input file: 3 motifs loaded

Settings: Metric=PCC, Alignment=SWU, Gap-open=1000, Gap-extend=1000, -nooverlapalign Multiple Alignment=IR, Tree=UPGMA, Matching against: JASPAR

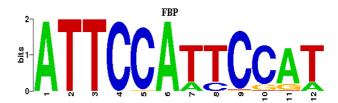
Note: All results files are removed nightly at midnight EST. Please save your results by saving "Webpage, complete".

<u>Download results as a PDF</u> <u>Click here to run STAMP again.</u>

#### **Multiple Alignment**

(Consensus sequence representations shown, but multiple alignment was carried out on the matrices)

Familial Profile: (click for matrix)

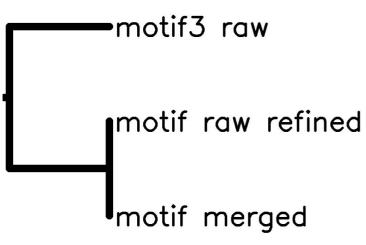


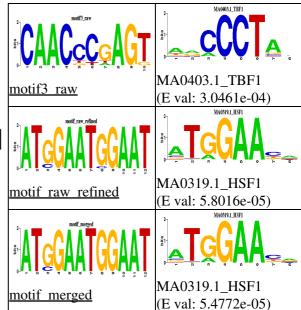
#### **Motif Tree**

Tree (drawn by <u>Phylip</u>)
<u>Click here for Newick-format tree</u> (viewable with <u>MEGA</u>)

**Input Motif** 

Best match in JASPAR





## **Motif Similarity Matches**



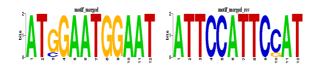
	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0403.1_TBF1	3.0461e-04	ACTCGGGTTG -NTAGGGYT-	MANALTERI COLOR OF THE PROPERTY OF THE PROPERT
MA0362.1_RDS2	8.0816e-04	CAACCCGAGT CCCCGAN-	TCGGGG
MA0133.1_BRCA1	1.1558e-03	-CAACCCGAGT NCAACNS	MANISS, LBRCAI
MA0074.1_RXRA_VDR	1.7325e-03	-CAACCCGAGT TGAACYCNNTGACCC	MANOTAL EXTRA VDR
MA0278.1_BAS1	3.1305e-03	ACTCGGGTTG NNNWNYTGACTCNGGCYNNNN	

# motif raw refined



	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0319.1_HSF1	5.8016e-05	-ATTCCATTCCAT	MARINAL HISTO
MA0406.1_TEC1	1.2589e-04	ATGGAATGGAAT -NGGAATGY	CATT CC
MA0045.1_HMG-I_Y	1.3035e-04	ATTCCATTCCAT- RTTTYKNYNTTTNTNN	MANNEL INVOLV
MA0370.1_RME1	7.4004e-04	ATTCCATTCCAT -TNCCTTTNNA-	AAGG A
MA0090.1_TEAD1	1.2863e-03	ATGGAATGGAAT CNSWGGAATGTR	ANALYTICATE E

### motif\_merged



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forward	1010130 00	ompliment

Name E value Alignment

MA0319.1\_HSF1 5.4772e-05 -ATTCCATTCCAT NNTTCCAT----



MA0045.1_HMG-I_Y	Y 1.1330e-04	ATTCCATTCCAT- RTTTYKNYNTTTNTNN	MAGNEL HINGLY
MA0406.1_TEC1	1.3134e-04	ATGGAATGGAAT -NGGAATGY	E CATT CC
MA0370.1_RME1	6.8200e-04	ATTCCATTCCAT -TNCCTTTNNA-	AAAGG A
MA0090.1_TEAD1	1.2949e-03	ATTCCATTCCATYACATTCCWSNG	2 AND THAN THAN THAN THAN THAN THAN THAN THAN

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