

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)

cmyc_24h_merged: CCACGTGS
cmyc_24h_ref_1_raw: -CACGTGS
cmyc 24h 1 raw: -CACGTGN

Familial Profile: (click for matrix)

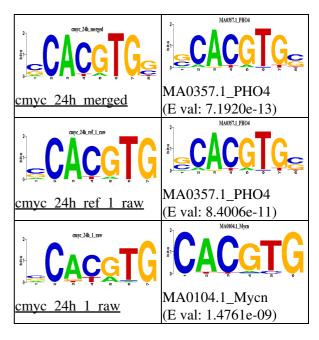


Motif Tree

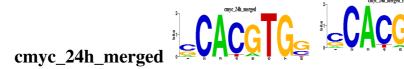
Tree (drawn by Phylip)
Click here for Newick-format tree (viewable with MEGA)

Input Motif

Best match in JASPAR



Motif Similarity Matches



	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0357.1_PHO4	7.1920e-13	CCACGTGS SCACGTGS	MARSTLEHOU EACH CGT C
MA0059.1_MYC_MA	X 1.3702e-11	-CCACGTGS ACCACGTGSTN	MANOSPI, MTC MAX CACCET CET S S S S S S S S S S S S S
MA0147.1_Myc	2.2250e-10	-CCACGTGS- NCCACGTGSN	EACGTGG
MA0128.1_EmBP-1	1.7804e-09	CCACGTGS CCACGTST	A ACCT GG

MA0104.1_Mycn

4.0657e-09

SCACGTGG -CACGTG-





	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0357.1_PHO4	8.4006e-11	SCACGTG- SCACGTGS	E CACGTGE
MA0128.1_EmBP-1	2.2991e-10	SCACGTG- CCACGTST	ACCT GG
MA0104.1_Mycn	9.2487e-10	SCACGTG -CACGTG	E CACGTG
MA0147.1_Myc	1.2123e-09	-SCACGTG NCCACGTGSN	CACGTGG
MA0004.1_Arnt	1.2659e-09	SCACGTG -CACGTG	EACGTG



forward

reverse compliment

Name E value Alignment Motif

	•		
MA0104.1_Mycn	1.4761e-09	NCACGTG -CACGTG	CACGTG
MA0004.1_Arnt	2.3317e-09	CACGTGN CACGTG-	EACGTG
MA0093.1_USF1	3.8273e-08	NCACGTG NCACGTG	E CACGTG
MA0409.1_TYE7	5.5190e-08	NCACGTG TCACGTG	CACGTGA
MA0257 1 DUO4	2.060%, 07	NCACGTG-	MARRET, LPHO4

SCACGTGS

Stamp Results

Sequence logo generation powered by <u>weblogo</u> <u>STAMP is written by Shaun Mahony</u>

MA0357.1_PHO4 2.0698e-07

02/17/20