

Homer *de novo* Motif Results (./rslt/)

[Known Motif Enrichment Results](#)

[Gene Ontology Enrichment Results](#)






If Homer is having trouble matching a motif to a known motif, try copy/pasting the matrix file into [STAMP](#)

More information on motif finding results: [HOMER](#) | [Description of Results](#) | [Tips](#)

Total target sequences = 33701

Total background sequences = 26523

* - possible false positive

Rank	Motif	P-value	log P-value	% of Targets	% of Background	STD(Bg STD)	Best Match/Details	Motif File
1		1e-15302	-3.523e+04	97.42%	1.55%	8.2bp (28.3bp)	GATA20(C2C2gata)/colamp-GATA20-DAP-Seq(GSE60143)/Homer(0.966) More Information Similar Motifs Found	motif file (matrix)
2		1e-55	-1.284e+02	40.66%	34.38%	17.8bp (20.1bp)	Adof1(C2C2dof)/col-Adof1-DAP-Seq(GSE60143)/Homer(0.881) More Information Similar Motifs Found	motif file (matrix)
3		1e-20	-4.728e+01	18.66%	15.74%	17.9bp (22.9bp)	SPL5(SBP)/colamp-SPL5-DAP-Seq(GSE60143)/Homer(0.809) More Information Similar Motifs Found	motif file (matrix)
4		1e-17	-4.045e+01	4.74%	3.35%	17.6bp (18.1bp)	Tb_0216(RRM)/Trypanosoma_brucei-RNCMPT00216-PBM/HughesRNA(0.826) More Information Similar Motifs Found	motif file (matrix)
5		1e-16	-3.902e+01	21.29%	18.50%	18.1bp (19.4bp)	CST6(MacIsaac)/Yeast(0.894) More Information Similar Motifs Found	motif file (matrix)