







Homer Known Motif Enrichment Results (../results/KDM5C_unbound_germDEGs)

[Homer de novo Motif Results](#)
[Gene Ontology Enrichment Results](#)
[Known Motif Enrichment Results \(txt file\)](#)

Total Target Sequences = 43, Total Background Sequences = 29226

Rank	Motif	Name	P-value	log P-value	q-value (Benjamini)	# Target Sequences with Motif	% of Targets Sequences with Motif	# Background Sequences with Motif	% of Background Sequences with Motif	Motif File	SVG
1		RFX(HTH)/K562-RFX3-ChIP-Seq(SRA012198)/Homer	1e-4	-9.772e+00	0.0251	11.0	25.58%	1847.1	6.32%	motif file (matrix)	svg
2		Rfx2(HTH)/LoVo-RFX2-ChIP-Seq(GSE49402)/Homer	1e-4	-9.447e+00	0.0251	11.0	25.58%	1915.9	6.56%	motif file (matrix)	svg
3		X-box(HTH)/NPC-H3K4me1-ChIP-Seq(GSE16256)/Homer	1e-3	-8.606e+00	0.0268	10.0	23.26%	1746.1	5.98%	motif file (matrix)	svg
4		Rfx1(HTH)/NPC-H3K4me1-ChIP-Seq(GSE16256)/Homer	1e-3	-7.821e+00	0.0441	13.0	30.23%	3127.4	10.71%	motif file (matrix)	svg
5		Oct2(POU,Homeobox)/Bcell-Oct2-ChIP-Seq(GSE21512)/Homer	1e-2	-6.216e+00	0.1758	9.0	20.93%	1968.9	6.74%	motif file (matrix)	svg
6		Oct11(POU,Homeobox)/NCIH1048-POU2F3-ChIP-seq(GSE115123)/Homer	1e-2	-4.737e+00	0.6427	9.0	20.93%	2464.7	8.44%	motif file (matrix)	svg