Homer Known Motif Enrichment Results

/ifs/venom/Crotalus_genomes/Ctigris_CLP27411/ATAC_Seq/HOMER/Macs2_Runs1/Macs2_Filtered_Hard50Overlap_ATAC_All_VG_Combined_peaks)

Home at seven Motif Results
Known Motif Enrichment Results
Known Motif Enrichment Results (vs Bkd)
Total Target Sequences = 7671, Total Background Sequences = 6171

Rank	Motif		P-value	Log P-value	q-value (Benjamini)	# Target Sequences with Motif	% of Target Sequences with Motif	# of Background Sequences with Motif	% of Background Sequences with Motif	Match	TF	STD
1		GRHL2(CP)/HMEC-GRHL2-ChIP-Seq(GSE86189)/Homer	1e-246	-567.196	0.000	5310.0	69.26%	3483.9	56.49%	seed edit logo		
2		NF1-half(var)(TGCNCAG-NF1-ChIP-Seq)/Lymphoblast(Blood)/Homer	1e-216	-497.526	0.000	5313.0	69.31%	3482.3	56.41%	seed edit logo		
3		FOXA1(Forkhead)/LNCAP-FOXA1-ChIP-Seq(GSE27825)/Homer	1e-148	-341.116	0.000	2360.0	29.98%	755.8	12.48%	seed edit logo		
4		FOXA1(Forkhead)/MCF7-FOXA1-ChIP-Seq(GSE26831)/Homer	1e-143	-329.878	0.000	2878.0	36.39%	1088.8	17.75%	seed edit logo		
5		NF1-FOXA1(CTGNNWGCAG)/LNCAP-FOXA1-ChIP-Seq(GSE27825)/Homer	1e-92	-212.776	0.000	458.0	5.82%	27.7	0.61%	seed edit logo		
6		FOXM1(Forkhead)/MCF7-FOXM1-ChIP-Seq(GSE76137)/Homer	1e-90	-208.024	0.000	2078.0	26.39%	850.8	13.75%	seed edit logo		
7		Foxo3(Forkhead)/Treg-Foxo3-ChIP-Seq(GSE66848)/Homer	1e-63	-146.426	0.000	1450.0	18.90%	586.9	9.10%	seed edit logo		
8		Foxo1(Forkhead)/Adipocyte-Foxo1-ChIP-Seq(GSE70261)/Homer	1e-51	-117.496	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
9		Foxo3(Forkhead)/Myocyte-Foxo3-ChIP-Seq(GSE70261)/Homer	1e-47	-109.326	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
10		EBF1(EBF)/Adipocyte-EBF-ChIP-Seq(GSE81475)/Homer	1e-46	-107.146	0.000	1225.0	15.55%	627.9	9.00%	seed edit logo		
11		Foxo1(Forkhead)/GlutamatergicNeuron-Foxo1-Isoform1-ChIP-Seq(GSE98678)/Homer	1e-44	-102.446	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
12		Foxo3(Forkhead)/CD8-Foxo3-ChIP-Seq(GSE92810)/Homer	1e-43	-100.016	0.000	1235.0	15.69%	684.7	7.90%	seed edit logo		
13		FOXK1(Forkhead)/MELJUNA-FOXK1-ChIP-Seq(GSE76137)/Homer	1e-42	-97.756	0.000	1809.0	20.46%	1052.1	11.31%	seed edit logo		
14		ELF5(ETS)/T470-ELF5-ChIP-Seq(GSE84187)/Homer	1e-41	-94.626	0.000	1092.0	13.87%	523.4	6.62%	seed edit logo		
15		Tbx19(T-box)/Pituitary-Tbx19-ChIP-Seq(GSE141262)/Homer	1e-32	-74.226	0.000	723.0	9.17%	353.9	5.16%	seed edit logo		
16		ESE3(ETS)/MDM2 E10-ChIP-Seq(GSE105600)/Homer	1e-32	-73.816	0.000	1213.0	15.42%	1028.9	13.86%	seed edit logo		
17		Tbx21(T-box)/Th1-Tbx21-ChIP-Seq(GSE39756)/Homer	1e-31	-71.606	0.000	1328.0	16.85%	1126.7	15.17%	seed edit logo		
18		ZEB1(Zinc finger)/SW620-ZEB1-ChIP-Seq(GSE90567)/Homer	1e-29	-67.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
19		Zic(Zinc finger)/Cerebellum-Zic-ChIP-Seq(GSE60731)/Homer	1e-28	-65.416	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
20		Foxo1(Forkhead)/T470-Foxo1-ChIP-Seq(GSE84187)/Homer	1e-27	-63.216	0.000	1092.0	13.87%	523.4	6.62%	seed edit logo		
21		Foxo3(Forkhead)/Myocyte-Foxo3-ChIP-Seq(GSE70261)/Homer	1e-26	-61.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
22		Foxo1(Forkhead)/Adipocyte-Foxo1-ChIP-Seq(GSE70261)/Homer	1e-25	-59.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
23		Foxo3(Forkhead)/Treg-Foxo3-ChIP-Seq(GSE66848)/Homer	1e-24	-57.016	0.000	1450.0	18.90%	586.9	9.10%	seed edit logo		
24		Foxo1(Forkhead)/GlutamatergicNeuron-Foxo1-Isoform2-ChIP-Seq(GSE98678)/Homer	1e-23	-55.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
25		Foxo3(Forkhead)/CD8-Foxo3-ChIP-Seq(GSE92810)/Homer	1e-22	-53.016	0.000	1235.0	15.69%	684.7	7.90%	seed edit logo		
26		FOXA2(Forkhead)/HEPG2-FOXA2-ChIP-Seq(GSE51987)/Homer	1e-21	-51.016	0.000	1809.0	20.46%	1052.1	11.31%	seed edit logo		
27		Tbx5(T-box)/Heart-Tbx5-ChIP-Seq(GSE35151)/Homer	1e-20	-49.016	0.000	1092.0	13.87%	523.4	6.62%	seed edit logo		
28		Tbx1(T-box)/Mesoderm-Tbx1-ChIP-Seq(GSE60474)/Homer	1e-19	-47.016	0.000	723.0	9.17%	353.9	5.16%	seed edit logo		
29		Tbx20(T-box)/Heart-Tbx20-ChIP-Seq(GSE53391)/Homer	1e-18	-45.016	0.000	1213.0	15.42%	1028.9	13.86%	seed edit logo		
30		EBF1(EBF)/Pro-EBF1-ChIP-Seq(GSE21512)/Homer	1e-17	-43.016	0.000	1328.0	16.85%	1126.7	15.17%	seed edit logo		
31		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-16	-41.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
32		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-15	-39.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
33		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-14	-37.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
34		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-13	-35.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
35		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-12	-33.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
36		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-11	-31.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
37		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-10	-29.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
38		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-9	-27.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
39		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-8	-25.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
40		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-7	-23.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
41		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-6	-21.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
42		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-5	-19.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
43		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-4	-17.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
44		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-3	-15.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
45		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-2	-13.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
46		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-1	-11.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
47		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-0	-9.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
48		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-1	-7.016	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
49		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-2	-5.016	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
50		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-3	-3.016	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
51		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-4	-1.016	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
52		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-5	0.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
53		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-6	2.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
54		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-7	4.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
55		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-8	6.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
56		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-9	8.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
57		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-10	10.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
58		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-11	12.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
59		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-12	14.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
60		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-13	16.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
61		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-14	18.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
62		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-15	20.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
63		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-16	22.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
64		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-17	24.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
65		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-18	26.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
66		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-19	28.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
67		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-20	30.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
68		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-21	32.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
69		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-22	34.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
70		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-23	36.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
71		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-24	38.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
72		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-25	40.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
73		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-26	42.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
74		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-27	44.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
75		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-28	46.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
76		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-29	48.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
77		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-30	50.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
78		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-31	52.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
79		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-32	54.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
80		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-33	56.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
81		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-34	58.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
82		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-35	60.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
83		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-36	62.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
84		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-37	64.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
85		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-38	66.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
86		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-39	68.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
87		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-40	70.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
88		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-41	72.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
89		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-42	74.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
90		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-43	76.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
91		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-44	78.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
92		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-45	80.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
93		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-46	82.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
94		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-47	84.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
95		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-48	86.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
96		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-49	88.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
97		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-50	90.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
98		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-51	92.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
99		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-52	94.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
100		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-53	96.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
101		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-54	98.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
102		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-55	100.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
103		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-56	102.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
104		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-57	104.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
105		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-58	106.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
106		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-59	108.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
107		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-60	110.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
108		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-61	112.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
109		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-62	114.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
110		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-63	116.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
111		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-64	118.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
112		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-65	120.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
113		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-66	122.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
114		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-67	124.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
115		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-68	126.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
116		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-69	128.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
117		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-70	130.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
118		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-71	132.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
119		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-72	134.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
120		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-73	136.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
121		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-74	138.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
122		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-75	140.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
123		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-76	142.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
124		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-77	144.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
125		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-78	146.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
126		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-79	148.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
127		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-80	150.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
128		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-81	152.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
129		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-82	154.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
130		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-83	156.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
131		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-84	158.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
132		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-85	160.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
133		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-86	162.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
134		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-87	164.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
135		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-88	166.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
136		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-89	168.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
137		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-90	170.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
138		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-91	172.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
139		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-92	174.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
140		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-93	176.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
141		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-94	178.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
142		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-95	180.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
143		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-96	182.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
144		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-97	184.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
145		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-98	186.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
146		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-99	188.984	0.000	1325.0	16.81%	681.9	9.77%	seed edit logo		
147		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-100	190.984	0.000	1180.0	14.97%	1000.9	13.48%	seed edit logo		
148		Foxf2(Forkhead)/Mesenchyme-Foxf2-ChIP-Seq(GSE65486)/Homer	1e-101	192.984	0.000	1016.0	20.53%	409.8	10.93%	seed edit logo		
149		Foxa2(Forkhead)/Liver-Foxa2-ChIP-Seq(GSE25253)/Homer	1e-102	194.984	0.000	1470.0	18.67%	752.9	11.98%	seed edit logo		
150		Foxf2(Forkhead)/Mesenchyme-