

Supplemental Figure 1

Target Sequences	Nam e	P- value	lg P- pvalue	q-value (Benjam in i)
	CTCF (Zf)/CD4+ -CTCF-ChIP-Seq (Baskiet al)/Homer	1e-134	-3.108e+02	0.0000
	BORIS (Zf)/K562-CTCF-ChIP-Seq (SE32465)/Homer	1e-109	-2.517e+02	0.0000
	Jun-AP1 (Zf)/K562-Jun-ChIP-Seq (SE31477)/Homer	1e-50	-1.162e+02	0.0000
	Fra1 (Zf)/BT549-Fra1-ChIP-Seq (SE46166)/Homer	1e-49	-1.133e+02	0.0000
	Fos2 (Zf)/3T3L1-Fos2-ChIP-Seq (SE56872)/Homer	1e-48	-1.113e+02	0.0000
	ATF3 (Zf)/GBM -ATF3-ChIP-Seq (SE33912)/Homer	1e-46	-1.067e+02	0.0000
	Fra2 (Zf)/Striatum -Fra2-ChIP-Seq (SE43429)/Homer	1e-46	-1.060e+02	0.0000
	BATF (Zf)/Th17-BATF-ChIP-Seq (SE39756)/Homer	1e-43	-9.918e+01	0.0000
	JunB (Zf)/Dendritic Cells-Junb-ChIP-Seq (SE36099)/Homer	1e-39	-9.106e+01	0.0000
	AP-1 (Zf)/ThioMacPU.1-ChIP-Seq (SE21512)/Homer	1e-34	-7.983e+01	0.0000
	Bach2 (Zf)/OCLy7-Bach2-ChIP-Seq (SE44420)/Homer	1e-31	-7.326e+01	0.0000
	HNF4a (NR), DR1/HepG2-HNF4a-ChIP-Seq (SE25021)/Homer	1e-28	-6.560e+01	0.0000
	FOXA1 (Forkhead)/MCF7-FOXA1-ChIP-Seq (SE26831)/Homer	1e-27	-6.296e+01	0.0000
	Foxa2 (Forkhead)/Liver-Foxa2-ChIP-Seq (SE25694)/Homer	1e-26	-6.197e+01	0.0000
	FOXA1 (Forkhead)/LNCAP-FOXA1-ChIP-Seq (SE27824)/Homer	1e-25	-5.922e+01	0.0000
	Foxa3 (Forkhead)/Liver-Foxa3-ChIP-Seq (SE77670)/Homer	1e-24	-5.714e+01	0.0000
	FoxEbox (Forkhead,bHLH)/Panc1-Foxa2-ChIP-Seq (SE47459)/Homer	1e-23	-5.386e+01	0.0000
	FOXM1 (Forkhead)/MCF7-FOXM1-ChIP-Seq (SE72977)/Homer	1e-23	-5.365e+01	0.0000
	Foxo3 (Forkhead)/U2OS-Foxo3-ChIP-Seq (EMTAB-2701)/Homer	1e-16	-3.714e+01	0.0000
	FoxL2 (Forkhead)/Ovary-FoxL2-ChIP-Seq (SE60858)/Homer	1e-14	-3.266e+01	0.0000
	FOXP1 (Forkhead)/H9-FOXP1-ChIP-Seq (SE31006)/Homer	1e-14	-3.251e+01	0.0000
	TEAD4 (TEA)/Tropoblast-Tea4-ChIP-Seq (SE37350)/Homer	1e-12	-2.951e+01	0.0000

Known Interactions

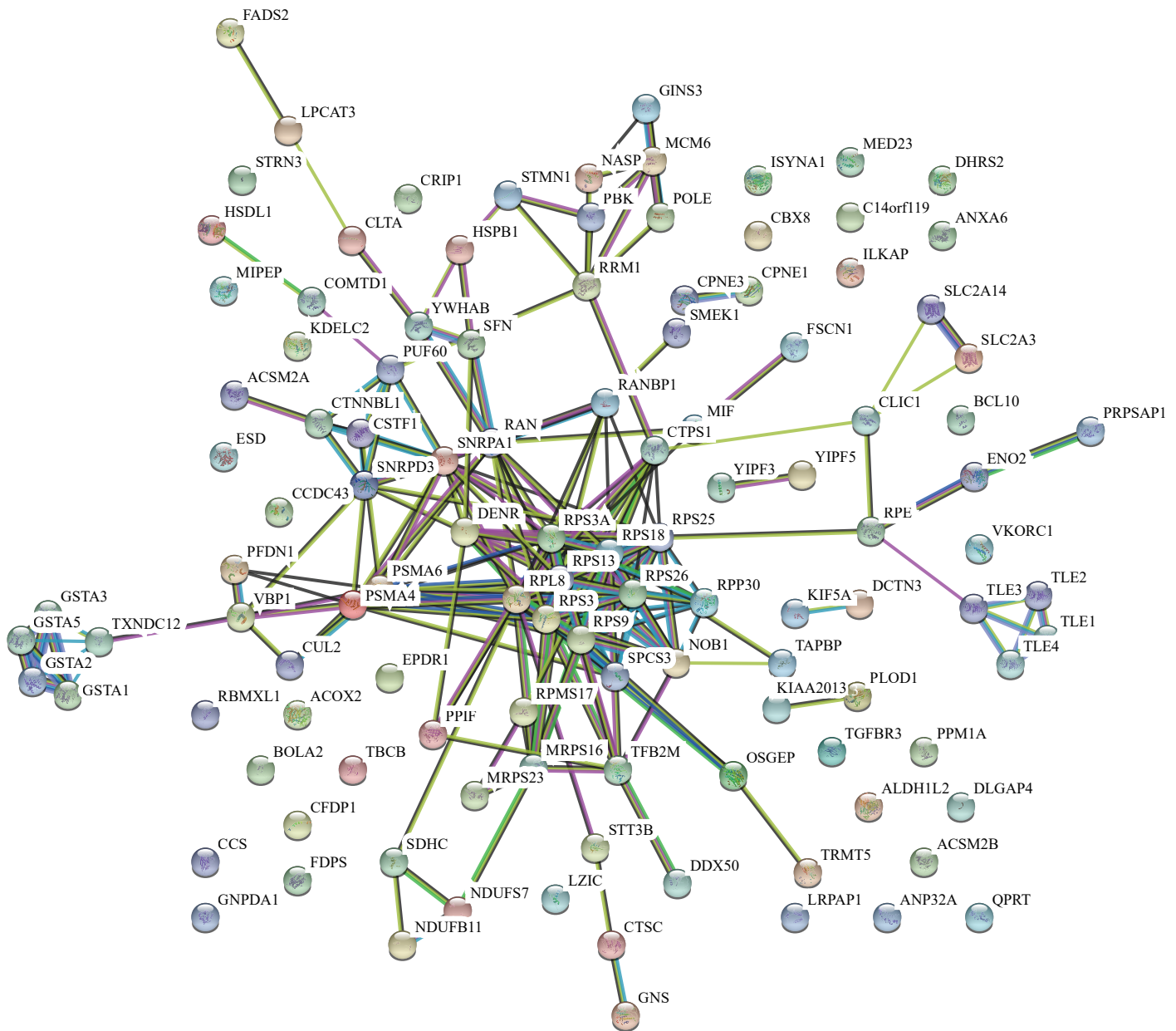
- from curated databases
- experimentally determined

Predicted Interactions

- gene neighborhood
- gene fusions
- gene co-occurrence

Others

- textmining
- co-expression
- protein homology



Supplemental Figure 2