brd4 -	0	0	0	0	0	0	0	0	0	0	0	0.14	
cdk7 -	0	0	0	0	0	0	0.036	0	0	0	0	0.067	0.10
ctcf -	0	0	0	0	0	0	0	0	0.032	0	0	0	- 0.12
erg -	0	0	0	0	0	0.047	0	0	0	0	0	0	
fancl -	0	0	0	0	0.008	0	0	0	0	0	0	0	- 0.10
gabpa -	0	0	0	0	0	0.085	0	0	0	0	0	0	
gata3 -	0	0	0.0067	0	0	0	0	0	0.00850	0.057	0	0	0.00
med1 -	0	0	0	0	0	0	0	0.031	0	0	0	0	- 0.08
myb -	0	0	0	0	0	0	0	0.11	0	0	0	0	
myc -	0	0	0	0	0	0	0	0	0	0	0	0.044	- 0.06
runx 0	.001	5 0	0	0	0	0	0	0	0	0	0	0	
runx1 -	0	0	0.008 D .	0044	1 0	0	0	0	0.0380	0.049	0	0	0.04
tal1 -	0	0	0.0032	0	0	0	0	0	0	0	0	0	- 0.04
tal1_scl -	0	0.003	0	0	0	0	0	0	0	0	0	0	
tcf12 -	0	0	0.0025	0	0	0	0	0	0	0	0	0	- 0.02
tcf3 -	0	0 9	9.3e-05	0	0	0	0	0	0	0	0	0	
znf335 -	0	0	0	0	0	0	0	0	0	0	0	0	
	GSE17954 -	GSE25000 -	GSE29180 -	GSE42575 -	GSE45864 -	GSE49091 -	GSE50622 -	GSE59657 -	GSE68976 -	GSE76181 -	GSE83116-	GSE83777 -	- 0.00