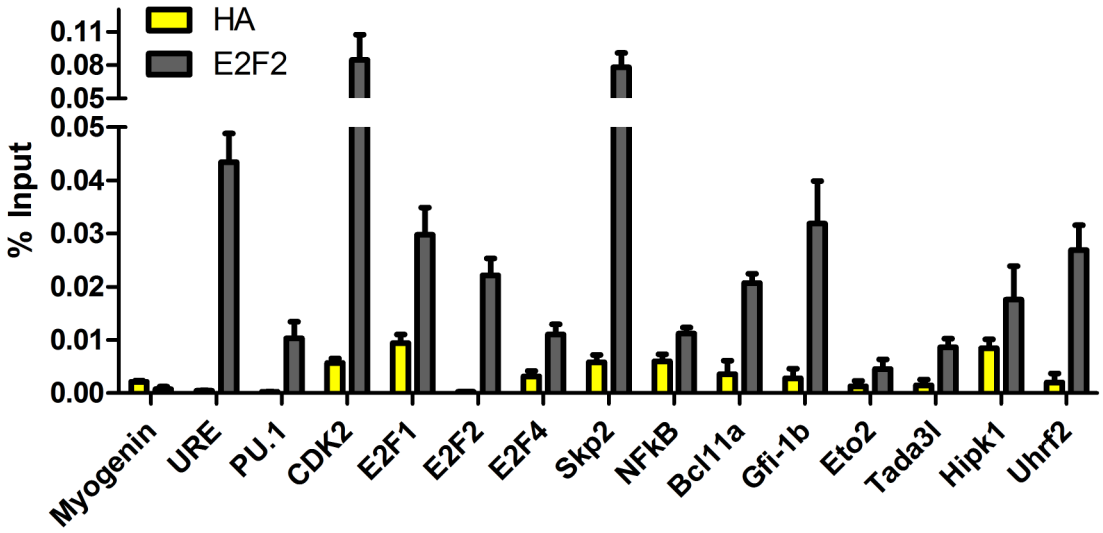
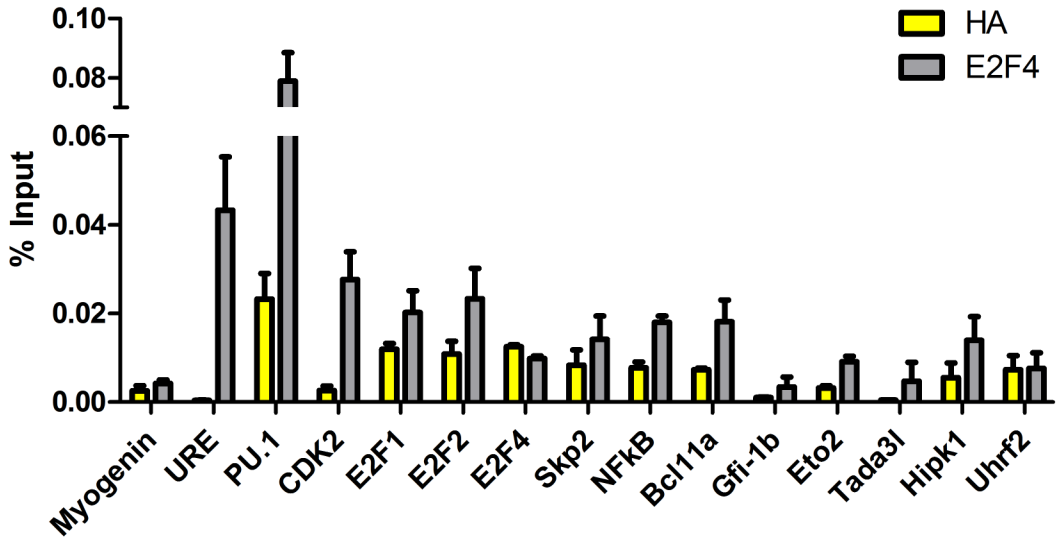
Figure S5 (Wontakal et. al.)

A.

Total Genes

Motif	Observed	Expected	Log_2Ratio
ZF5	50%	14%	1.82
E2F	38%	9.5%	2.01
AP2	30%	5.6%	2.42
ETF	22%	0.49%	5.51
CHCH	28%	5.0%	2.47
GC	22%	3.4%	2.68
Sp1	22%	3.6%	2.58
ABI4	19%	3.1%	2.59
Movo-B	16%	2.0%	3.03
Muscle	15%	5.3%	1.53

B.



C.