

SUPPLEMENTAL MATERIAL

Male et al., <http://www.jem.org/cgi/content/full/jem.20132398/DC1>**Table S1.** Transcription factors reduced in E4bp4^{-/-} lineage-depleted bone marrow, defined by microarray screen

Transcription factor	Gene symbol	Fold reduction	P
Interferon activated gene 205	<i>Ifi205</i>	1.86	0.00351
Inhibitor of DNA binding 2	<i>Id2</i>	1.64	0.03575
Nuclear factor, interleukin 3 regulated (E4bp4)	<i>Nfil3</i>	1.50	0.00121
GATA-binding protein 3	<i>Gata3</i>	1.46	0.00121
Forkhead box protein H1	<i>Foxh1</i>	1.34	0.000813
Ikaros family zinc finger 5 (Pegasus)	<i>Ikzf5</i>	1.25	0.001428
Leucine zipper transcription factor-like 1	<i>Lztf1</i>	1.24	0.000909

Table S2. Primers for the amplification of putative E4bp4-binding regions

Region	Forward primer	Reverse primer	Amplicon bp
Gene desert	5'-CATCCAGGAGCCACTGAAAT-3'	5'-GACATGGACGCTACCATGCTC-3'	184
Per2B	5'-TTGACCGCGGAAGCGGTGAGTG-3'	5'-GGGACGCGAGTGTGAACCTGG-3'	220
Eomes 1	5'-GCGTCCGTAATTGCTTATTAG-3'	5'-CGTTTCCGCAAACAGTCAAG-3'	125
Eomes 2	5'-CAAATTCAACTGCCTAGGTATC-3'	5'-AGAATATCTCCAAGATATGAATC-3'	77
Eomes 3	5'-GTATAGCTCTGTCTTCTGTAC-3'	5'-GAGAGAGAGACTGTTCACTCT-3'	125
Id2 1	5'-ACTTTGCCACCTACACTGAAG-3'	5'-CAAGAATGAGGCAGGGTTAC-3'	124
Id2 2	5'-GTCTGTAAATGTTTGGTTTGG-3'	5'-TTCATAGTAAGTTACTGGCATTTC-3'	188
Id2 3	5'-AGATTTCTCTGGCTAGCTGAA-3'	5'-CTCTAACAGCCCGCTCACA-3'	149