

ChIP-PCR primers

Main Figures		Supplementary Figures	
Figure 1c		Figure S1c	
<u>Primer Name</u>	<u>Sequence</u>	<u>Primer Name</u>	<u>Sequence</u>
Bach2-F	GCCACTTACTTGGGAAAATG	Bcl2l1 F	CCGAGACGCAAAAAGGAGT
Bach2-R	GACAACCTGCCAACGTCTTGA	Bcl2l1 R	GGGACAGATGGATGAAGGAA
Bcl2l1-F	GTTGCCACACAAAAGTCAAG	Nfkb1a-F	CTGAGTGGCTGGAAAGTCCT
Bcl2l1-R	TTGCCATCATAGGGTTTTCC	Nfkb1a-R	CTGGCAAATCGCTAAGAGG
Ccl3-F	AGGTCAGGTGCTCTCTCAGC	Negative-control-F	GCCATCTTTCTACCTCCACA
Ccl3-R	GAGCCCTCGAAGTGTGACTC	Negative-control-R	CTCTGTCTGCCCTGTTAT
Cdc274-F	CTTTCCTCATTCCCCTTCA	Figure S4e	
Cdc274-R	CTGCGTGGGTCTGGAGTTA	<u>Primer Name</u>	<u>Sequence</u>
Dennd4a-F	CCTTTTCCCTGACCTCTGTCT	Arid3a-F	GGGGTTTTTAAGCCCATTTT
Dennd4a-R	TCCTCTTTCACAAGCATCCA	Arid3a-R	GCAAGATGGATAAGCCAAGC
Etv6-F	TGTTACTTCTGTCCCCACCA	Arid3a-neg-F	TAGACGGGGAATCTGGACAC
Etv6-R	TCCAATTCCAATTCTCTCTT	Arid3a_neg_R	GTCGGGTGAGCAACAAAGA
Gm14005-F	CAGGCACAATGACATCTGCT	Bcl11a-F	CATTTGGTGATTCTAAGGCTGA
Gm14005-R	CTCTGACTCAACGGGGAAAC	Bcl11a-R	GGTTTGAGGAGCAGGAAGAC
Gm20098-F	GTGACTTCCGCTCTGTAGCA	Bcl11a-neg-F	GTTGCTGGAGGTGGGTGTA
Gm20098-R	GGCAACCTATCTCTGTGTGG	Bcl11a-neg-R	GCAGTCTATGGTGAGAGTTGA
Gpr132-F	GAACCTCCCTAACCAACAG	Bcl6b-F	AGTGGGAGAAGGAGGAGTGA
Gpr132-R	CCTGAGAAGAGCCATCATCC	Bcl6b-R	GGACAATGGAAGGAAACGAA
Klf10-F	AGACGGGAGCACTGAGACA	Bcl6b_neg_F	GGACAGGAGGAAAGGGCTAT
Klf10-R	ACACCTTTGCCGCTGATT	Bcl6b_neg_R	CCCACACAGCAACTCTATT
Mcl1-F	GGAAAGGGTGGGATGTCA	Cbx3-F	GAGGAAGGAGACGGGAAAAT
Mcl1-R	TGCCTGAGAAGAAAAGCAAG	Cbx3-R	TAGGGGCTCGTTGTGTT
Mcm9-F1	TGGAATCGCCTCAAAGAAC	Cbx3-neg-F	ACAGTCTTTGTGGCAGCAGA
Mcm9-R1	TGCTTCAACCTCCCTTG	Cbx3_neg_R	TTCTACCCCTCCCTTT
Nfkb1a distal-F	AACTGGCTCGTCCCTCACT	Hhex-F	AAGCGCAGTGAAAAAGTTG
Nfkb1a distal-R	CACTCAGGGCTCATCAAAAAG	Hhex-R	AAGGTGCAGGAAAAGCAATC
Pdcd1lg2-F	ACGACTGCTTGGGGAAGTT	Hhex-neg-F	ACCTCAACAGGCGATGACTT
Pdcd1lg2-R	CACCTGGGGATGACTCTAA	Hhex_neg_R	CCAGCCACAAAGATGGAAT
Psm6-F	TGTGGGCTTTGCTAAACAGA	Ldha-F	AAGAGTCGAGTCCTTCTCT
Psm6-R	TGCCTCGATGGAAAGTTTG	Ldha-R	TCTACTTCACGCAAACTCAA
Slamf6-F	CAGACACACAACCTGGCACTACA	Ldha_neg_F	TTGCGTCAGGGAAATGTATG
Slamf6-R	AACCGAAGGTGGAACATCC	Ldha_neg_R	AAGGTCGTGGGAAAGATGG
Stim2-F1	TGGCTTCTTGGGTTGGTAAT	Mki67-F	AAAACGGCGGTTCAAATTC
Stim2-R1	CCTCTCTCCTTGGTCTTTG	Mki67-R	CCTCTGGACTTCCCAATC
Tnfaip3-F	TGATGGGAACTGGAAATGC	Mki67-neg-F	AGAGAGACACACTCAGGATGGA
Tnfaip3-R	CTGTGTGTCTGGGTGGTTTTT	Mki67-neg-R	TGTTGGGGAAGGATTAGGAG
Tpd52-F	ATGACCCAGAGGAAATGGAA	Nfkb1e-F	ATGATGGCTTGTGTGGAATG
Tpd52-R	CAGCCAGTGAGCCTAATCAA	Nfkb1e-R	AACAGGGAGGACAGGAGGA
Zfp619-F	GGTTTACGAGGATTCAACAAGG	Nfkb1e_neg_F	CTCAGCCTCAACTCTCTCC
Zfp619-R	AGACCTGGGAGGGTTTCAGT	Nfkb1e_neg_R	TCTCTTCTCCCAGCCTTC
		Ppp1ca-F	GAGAAGTCAGCGATTAACATGC
		Ppp1ca-R	GCCGCGTTCAGATGAGTTA
		Ppp1ca-neg-F	TCTGAGGAGGGCTTGGAGGT
		Ppp1ca_neg_R	CAAAAGGGTTCAGTTCCAC
		Xrcc-1-F	CACGTAGGACCCAATCACTG
		Xrcc-1-R	CTCAGGTTCCACCACTCTCC
		Xrcc1_neg_F	ATTTCAACCTCCCCTGATT
		Xrcc1_neg_R	GAGCGAGTCCAGCAAAAGTA