

Table S2. Primer sequences used to measure CncC and dKeap1 occupancy by ChIP

gene	position	5'-primer	3'-primer
<i>sad</i>	-5000	GCCAACGTCTGGTCAAAAAT	CCAAGTTCTTTTTGTGCTCTGTT
	-1200	CCCCTAATAGACCATAAATATGACAAG	CTTGACGAAGGTTGTGAAAATG
	60	GAGCAGGTGCAGGTGGTTAC	TCTTCTCGGTCATCGGTGTA
	4100	ATCGCTCATGTTTGCCTTTT	GCTCAAAGTGTGATGGATCG
<i>dib</i>	-100	TTCACCACAGGTAGGCACAA	CATTCCGGGAAAAAGGCATTA
<i>nvd</i>	-100	CGCGTATGCCGGAAAAATA	GTTCACCGTTTTCGTTAGGG
<i>spok</i>	300	GTCCGCCAAAGTATTTTCCA	TTACGGTTTTTGGCACATTGA
<i>phm</i>	0	CTGATGTTACCATTGTCTGAACG	TCAACGTGTGTCTACGTCAGC
<i>shd</i>	-200	GGATCGATGGAAGCACTTGT	ACGATGGGAGGCGATTTA
<i>dKeap1</i>	700	AAGGTTTTGCCTCTTCAGCA	CGGCTTTAAATTTCCGCATA
<i>gstD1</i>	-1000	CATGTGCCTTTGACTCATCATT	GACCATAAAAATATAACCGTTTTTCG
<i>Rp49</i>	0	TCCTTCCAGCTTCAAGATGAC	GTGCGCTTCTTCACGATCT
<i>Actn3</i>	1200	CGAAGAAACAGCCAAAATCG	GGACGCCACTTTGTATTGAAC
<i>Gapdh1</i>	300	GAAAAGCGGCAGTCGTAAT	AATTCGATCTTCGACATGG