

Table S8. Primers used for ChIP-seq validation (qPCR)

Genomic Region	Primers	Sequence (5'-3')
Chr 15: 61,814,801 - 61,815,300	<i>Myc</i>	Fw ACAAATCCGAGAGCCACAAC Rev TCGAGCTCATTGCACAATTC
Chr 15: 61,687,900 - 61,688,400	<i>Negative (Myc)</i>	Fw TGGGGTACTCTGGGTTCATC Rev TAGGGACTGGAAATGGCATC
Chr 11: 67,817,666 - 67,818,084	<i>11qB3</i>	Fw TAGCTGAATTTCCCCCACTG Rev CTGAATCAGCACCGACAGAA
Chr 14: 84,838,181 - 84,838,727	<i>Pcdh17</i>	Fw TTCCCGGTGTCAATCAATTT Rev TACATACTGG TCCCCCTTG
Chr 5: 58,105,123 - 58,105,623	<i>Pcdh7</i>	Fw TTTGTCCCTTGGTGGAGTTC Rev AGGGTCGGGAGTCTTTCATT
Chr 18: 37,179,478 - 37,179,961	<i>Pcdha12</i>	Fw CCTCATGATGTGCTCTTCA Rev AGCATGGTACACACGCAGAA
Chr X: 117,401,147 - 117,401,603	<i>Pcdh11X</i>	Fw AGTCTGCTTTTGCCAGGTGT Rev ACCCTGAAAATCCTCCAAGG
Chr 18: 37,644,356 - 37,644,924	<i>Pcdhb17</i>	Fw GGCCAGTTTCTCACCACTA Rev GCAGTCTCCATGGCTCTTTC
Chr 18: 37,300,013 - 37,300,550	<i>Negative (Pcdh)</i>	Fw GGGCGTTTTCTGAGATTGAG Rev CAGGGCATTGTGCTTGTTTA
Chr 18: 37,901,429 - 37,901,923	<i>Pcdhgb6</i>	Fw TCATAACCCCATGTTTCAGAGC Rev GAGTTCCCCTTGGTGTGTGT
Chr 18: 37,672,967 - 37,673,448	<i>Pcdhb21</i>	Fw TGCAGGACAGGATGAGAGAA Rev GGCAGCTCAGAGAGTGGTTC
Chr 11: 100,030,172 - 100,030,721	<i>11qD3</i>	Fw TAGGCCAAACTGCACTTCCT Rev TCCTACTCTGGCTGGGAAGA
Chr 11: 99,482,356 - 99,482,857	<i>11qD3 (Negative)</i>	FW CTTGCGGCTGAAGTTTTAGG Rev CAGTTAGCAACCCTGCATGA
Chr 9: 77,391,786 - 77,392,244	<i>9qE1</i>	Fw CAAAATGAGGAGCTCCAAGC Rev GGAGAAACGTTCTGGTCCAC
Chr 15: 101,732,703 - 101,733,262	<i>15qF3</i>	Fw CCTTGCAAACACAGCATT Rev GGGCAAGCTAATCACCTGAA
Chr 10: 117,879,278 - 117,879,482	<i>IFNg</i>	Fw ATTTGTGGCCTTGCAGTTTC Rev TATTTACCAATTGCCCTGA