

Additional file 8

Table S1. Sequences of RT-qPCR/ChIP primers and EMSA probes

| RT-qPCR primer | | Sequence |
|----------------|----------|--|
| CD274 | | Forward: 5'- GGAGCCATCTTATTATGCCTT-3' Reverse: 5'- GTTTGTATCTTGGATGCCACA-3' |
| GAPDH | | Forward: 5'- TCTGACTTCAACAGCGACACC-3' Reverse: 5'- CTGTTGCTGTAGCCAAATTCGTT-3' |
| ChIP primer | Location | Sequence |
| E1. | -1289 | Forward: 5'-GCAGTACCTGTAAACTGTATTGCC-3' Reverse: 5'-GTTCTCTTTGGCCCAATAA-3' |
| E2. | -364 | Forward: 5'-ATATGGGTCTGCTGCTGACTTT-3' Reverse: 5'-ACAACAAGCCAACATCTGAACG-3' |
| E3. | -21 | Forward: 5'-ATTTACCGAAGGTCAGGAA-3' Reverse: 5'-AGCGAGCTAGCCAGAGATACTG-3' |
| EMSA probe | | Sequence |
| Wild type | | Forward: 5'-AGAGGTGGGCGGGACCCCGCCTC-3' Reverse: 5'-GAGGCGGGGTCCCGCCCACCTCT-3' |
| Mutant | | Forward: 5'-AGAGGTGTGCATCTGCATGCCTC-3' Reverse: 5'-GAGGCATGCAGATGCACACCTCT-3' |