

Suppl. Table S5: Oligonucleotides used for RT-PCR, cloning, and ChIP-PCR.

<u>Name</u>	<u>Sequence</u>
F_mTnni2	CCTGAAGAGTGTGATGCTCCA
R_mTnni2	CCCGTTCTTCAGTGTCTT
F_mMyh2	TCGTTGCCAGTAAGGGCT
R_mMyh2	TGCATCTTGCTCTGAATGG
F_mActa1	CACCAGGGTGTCAATGGTAGG
R_mActa1	TGGTACGGCCCGAAGCATAG
YY1-RT-F	ATGTGGTCCTCGGATGAAAA
YY1-RT-R	CTTTGTGCAGCCTTATGAG
F_mGAPDH	CACGGCAAATTCAACGGCACAGTCAAGG
R_mGAPDH	GTTCACACCCATACAACATGG
F_mMyog	ATGGTGCCAGTGAATGCAA
R_mMyog	ACCCAGCCTGACAGACAATC
primary miR1-2/133a-1-mib-F	AGACGTGGCTAAAGTGGAAGA
primary miR1-2/133a-1-mib-F	TATTACGAGCATTCAAGGTCAAGC
miRNA-1-2/133a-1-E1-F	GCATGGTACCGGAGGAGCAGGAGTTGAGTT
miRNA-1-2/133a-1-E1-R	GCATGCTAGCCCAGCTGAGGTTTCCTAGT
miRNA-1-2/133a-1-E2-F	AAAAGGTACCGCTACATTAGTTATTCCACAGG
miRNA-1-2/133a-1-E2-R	AAAAGCTAGCCAACACAATCTCAGACAGTATGC
miRNA-1-1/133a-2-E3-F	AAAAGGTACCACACATCAGTCCCTCAGAAC
miRNA-1-1/133a-2-E3-R	AAAACTCGAGGATCCGTTCCAAACATG
miR-206/133b-E4-F	AAAAGGTACCGGACTGATCCAAGGCTGTTCT
miR-206/133b-E4-R	AAAAGCTAGCTTCTATTGGGTTGCTTGG
muta-1-2/133a-1-E1-A-F	AAAATAGCAGGACAAGGCCAAAGCAGTGGG
muta-1-2/133a-1-E1-A-R	CCCAACACTGCTTGGGCTTGTCTGCTATTTT
muta-1-2/133a-1-E1-B-F	GCTATGGCTGACTAAATGCTCTCAGGGAGAGAG
muta-1-2/133a-1-E1-B-R	CTCTCTTCCCTGAGGACATTAGTCAGCCATAGC
F_mMyHC_YY1 ChIP	CACATACCTACCTTAAGGG
R_mMyHC_YY1 ChIP	CTCCATTCCCTATCAGATGC
F_mTnni2_YY1 ChIP	GTGAGGCCAGCCATCTC
R_mTnni2_YY1 ChIP	CATTTCTGCTTGTGTGTC
YY1 ChIP_miRNA1-2/133a-1-E1-A-F	GGGAGAATCTGGAAATGTA
YY1 ChIP_miRNA1-2/133a-1-E1-A-R	AAAGCTGAGGAGGATTCTAT
YY1 ChIP_miRNA1-2/133a-1-E1-B-F	AGCAAGATAGAATCCTCCTCA
YY1 ChIP_miRNA1-2/133a-1-E1-B-R	AGGCAGCTAACGATTGAAACA

YY1 ChIP_miRNA1-2/133a-1-E2-C-F
YY1 ChIP_miRNA1-2/133a-1-E2-C-R
YY1 ChIP_miRNA-206/133b-E4-E-F
YY1 ChIP_miRNA-206/133b-E4-E-R
ChIP-PCR-NC-F
ChIP-PCR-NC-R
mPax7-RT-F
mPax7-RT-R
Pax7-miR-1-siteA-top
Pax7-miR-1-siteA-bottom
Pax7-miR-1-siteB-top
Pax7-miR-1-siteB-bottom
F_mMyoD
R_mMyoD
YY1-3UTR-miR1-mut-top
YY1-3UTR-miR1-mut-bottom

GACACTATCAGATGCTGGACT
GCACTTGCCTGGTTCTATT
AGTCGGGACCCCTTCTTCT
CAATCACCTGCGTCACAATC
GAAATGCTGTTAACCTGGCAG
GAGGCTGACCTTGGAGTCAC
CTCAGTGAGTTCGATTAGCC
CCAGACGGTCCCCTTGT
CTAGTCATGTCCTGCCCTCAGCCAGCAAGACATTCTAGAGAGAGGAACGAGCT
CGTTCCCTCTCTAGGAATGTCTTGCTGGCTGAGGGCAGGAGACATGA
CTAGTCAGGGCCCAGTCCCAGCCAAGGGAGCAACATTCCAACGCTTGGACCAGAGCT
CTGGTCCAAGCGTTGGAATGTTGCTCCCTGGCTGGACTGGCCCTGA
CTGCTCTGATGGCATGATGG
GTTCCCTGTTCTGTGTCGCT
CTAGTGCTAAATGGGACTTCTTCCGGGCCTTATAAATATGAAGCGAGCT
CGCTTCATATTATAAGGCCGAAAAGAAGTCCCATTTAGCA