

# Supplementary Information

**Table S1.** The Sequence of Negative Control for miRNA mimic and inhibitor.

Name	Sequence
NC(5' to 3') (Control for mimics)	UUC UUC GAA CGU GUC ACG UTT ACG UGA CAC GUU CGG AGA ATT
NC(5' to 3') (Control for inhibitor)	CAG UAC UUU UGU GUA GUA CAA

**Table S2.** Primer sequences and parameters of Myoblast differentiation marker gene for real-time PCR.

Gene	Primer sequence (5' to 3')	Accession No.	T <sub>m</sub> /°C
<i>Myf5</i>	F: GGACCAGTTGAGCCAAGAG R: CGAAAGCTTGGTTGACCTTC	NM_008656	55
<i>MyoD</i>	F: AGTGAATGAGGCCTTCGAGA R: GCATCTGAGTCGCCACTGT	NM_010866	60
<i>MyoG</i>	F: CAATGCACTGGAGTTCTGGT R: CTGGGAAGGCAACAGACAT	NM_031189	60
<i>MyHC</i>	F: CGCAAGAACATGTTCTCAGGCT R: GCCAGGTTGACATTGGATTG	NM_030679	60
<i>MURF1</i>	R: GCTGGTGGAAAACATCATTGACATC F: GCTGGTGGAAAACATCATTGACATC	NM_001039048	59
<i>Atrogin</i>	R: CTGAATAGCATCCAGATCAGCAGG F: TTGATAAAGTCTTGAGGGAAAGTG	NM_026346	57

**Table S3.** Primer sequences and parameters of miR-199a-3p target genes for real-time PCR.

Gene	Primer sequence (5' to 3')	Accession No.	T <sub>m</sub> /°C
<i>IGF-1</i>	F: AGGCATTGTGGATGAGTGGT R: TCCTTTGCAGCTTCGTTT	NM_010512	58
<i>PIK3r1</i>	F: CTGAGATTGCTTCGTGAC R: CTTATCCCATGGCTATGA	NM_001077	61
<i>mTOR</i>	F: TCCGAGAGATGAGTCAAGAGG R: CACCTCCACTCCTATGAGGC	NM_020009	61
<i>RKS6KA6</i>	F: AACGGTTATGCGGGAGC R: AACTGTGCGGGATCTGC	NM_025949	60
<i>GAPDH</i>	F: GGCAAGTTCAACGGCACAG R: CGCCAGTAGACTCCACGACAT	NC_005103	60

**Table S4.** Primer sequences used for cloning 3' UTR of miR-199a-3p target genes.

Gene	Primer sequence (5' to 3')	Length	Position (NCBI)
<i>IGF-1</i>	F: CCG <u>CTCGAG</u> CTGTGGGTGGGTGAGTAC R: ATTT <u>GCGGCCG</u> CATGCTTGAAATGGGAATG	691	4851-5542
<i>PIK3r1</i>	F: CCG <u>CTCGAG</u> GGTTGACTCTGGGCTTTGC R: ATTT <u>GCGGCCG</u> CCATGGCTATGATCACTCTGGT	681	4046-4727
<i>mTOR</i>	F: CCG <u>CTCGAG</u> AAACCACGTCGTCTCCTC R: ATTT <u>GCGGCCG</u> CAGCATATCCCTCCCTCAC	625	7783-8408
<i>RKS6KA6</i>	F: CCG <u>CTCGAG</u> CTACCTTGAATCCTATGGCG R: ATTT <u>GCGGCCG</u> GACAAAATACTACACCAAGACTAAAT	660	3682-4342

Underlined symbols are the Restriction Enzyme cutting sites.

© 2013 by the authors; licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution license (<http://creativecommons.org/licenses/by/3.0/>).