Supplementary Materials: miR-30-5p Regulates Muscle Differentiation and Alternative Splicing of Muscle-Related Genes by Targeting MBNL

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Table S1. Primers for the real-time PCR for miR-30-5p.

Name of Primers	Sequence of Primers (5′-3′)	Amplicon Size
Stem-loop RT-miR-30a-5p	gtcgtatccagtgcagggtccgaggtattcgcactggatacgacagcttc	
miR-30a-5p-F	GGCGTGTAAACATCCTCGACTG	
miR-30a-5p-R	GTGCAGGGTCCGAGGT	62 bp
Stem-loop RT-miR-30b-5p	gtcgtatccagtgcagggtccgaggtattcgcactggatacgacagctga	
miR-30a-5p-F	GGCGTGTAAACATCCTACACTC	
miR-30a-5p-R	GTGCAGGGTCCGAGGT	60 bp
Stem-loop RT-miR-30e-5p	gtcgtatccagtgcagggtccgaggtattcgcactggatacgacagcttc	
miR-30e-5p-F	GGCGTGTAAACATCCTTGACTG	
miR-30e-5p-R	GTGCAGGGTCCGAGGT	62 bp
U6-F	GCTTCGGCAGCACATATACTAAAAT	
U6-R	CGCTTCACGAATTTGCGTGTCAT	107 bp

Stem-loop RT-miR-30a-5p, Stem-loop RT-miR-30a-5p and Stem-loop RT-miR-30a-5p were used to reverse transscription of miR-30a-5p, miR-30b-5p and miR-30e-5p.

Table S2. Primers for the constructs.

Name of Primers	Sequence of Primers (5′-3′)	Amplicon Size
MBNL1-3′ UTR-F	GCTCTAGATCAGCCACAAGACATCCACA	448 bp
MBNL1-3′ UTR-R	GCTCTAGATCAGATCCCTCCCTCACCAC	
MBNL2-3′ UTR-F	GCTCTAGAGGGTTGTAACTGACTACAGCAT	220 bp
MBNL2-3′ UTR-R	GCTCTAGAAATTGTATCGCTATTACCTTGA	
MBNL3-3′ UTR-F	GCTCTAGATTCAACCCGCCTAGATAGAT	375 bp
MBNL3-3′ UTR-R	AAATACTGTGGAATAACCCT	
Pre-miR-30a-5p-F	CCCAAGCTTTTGGGAGAAGACTTAATGGTGT	319 bp
Pre-miR-30a-5p-R	GGGGTACCTAATGAAAATGTAGGGATGGGT	
Pre-miR-30b-5p-F	CCCAAGCTTTCATGTCAATCTTTGTACCTCCTG	291 bp
Pre-miR-30b-5p-R	GGGGTACCTTGCCATATCCTCTATCCGTGT	
Pre-miR-30e-5p-F	GGGGTACCAGGAGGAACTGAGCCGTGGACA	241 bp
Pre-miR-30e-5p-R	CCCAAGCTTGACCCTGCCTGGGGACCTTTGG	

The red font represents the sequence sites recognized by restriction enzyme.

Table S3. Primers for constructs of the mutant 3′ UTR.

Name of Primers	Sequence of Primers (5′-3′)	Amplicon Size
Mut-MBNL1-3′ UTR-F1	GCTCTAGATCAGCCACAAGACATCCACA	336 bp
Mut-MBNL1-3′ UTR-R1	GACCTTTGTTATTTGATTTGCTTGAAAGAAATA	
Mut-MBNL1-3′ UTR-F2	TATTTCTTTCAAACAAATCAAATAACAAAGGT	147 bp
Mut-MBNL1-3′ UTR-R2	GCTCTAGATCAGATCCCTCCCTCACCAC	
Mut-MBNL2-3′ UTR-F	GCTCTAGAGGGTTGTAACTGACTACAGCAT	187 bp
Mut-MBNL2-3′ UTR-R	GCTCTAGAAGTATTTAAAAAAGAAAACAACATT	
Mut-MBNL3-3′ UTR-F1	GCTCTAGATTCAACCCGCCTAGATAGAT	185 bp
Mut-MBNL3-3′ UTR-R1	GAGCAAAAAGTTTTGTACATGTGGATTCT	
Mut-MBNL3-3′ UTR-F2	AGAATCCACATGTACAAAACTTTTTGCTC	210 bp
Mut-MBNL3-3′ UTR-R2	AAATACTGTGGAATAACCCT	

The red font represents the sequence sites recognized by restriction enzyme *Xba*I; the underline font represents the mutant sites in the target sequences recognized by miR-30-5p.

Table S4. Primers for real-time PCR of genes.

Name of Primers	Sequence of Primers (5′-3′)	Amplicon Size
MBNL1-F	CAACAACATCTGCCACAA	109 bp
MBNL1-R	TACATCTGGGTAACATACTTG	
MBNL2-F	CGTAACCGTTTGTATGGATTACAT	76 bp
MBNL2-R	GTGTGCAGGAGGGTGAAA	
MBNL3-F	TGATAATACTGTGACCATCTGC	79 bp
MBNL3-R	AGGAGGATGAAAATACTTGC	
MyoG-F	GTCCCAACCCAGGAGATCATT	70 bp
MyoG-R	GACGTAAGGGAGTGCAGATTGTG	
MHC-F	CAATAAACTGCGGGCAAAGAC	75 bp
MHC-R	CTTGCTCACTCCTCGCTTTCA	
MyoD-F	GGAAGGGAAGAGCAGAAG	82 bp
MyoD-R	AAGGACTACAACAACAACAAC	
Trim55-F	AGTGAGTGGTAAGGAGTC	97 bp
Trim55-R	CCAGATGTAGTAGAGAATAAGAA	
INSR-F	TGGAGGAGTCTTCATTCA	100 bp
INSR-R	CTACTGTCCTCGGCACCAT	
GAPDH-F	AACTTTGGGATTGTGGAAGG	222 bp
GAPDH-R	ACACATTGGGGGTAGGAACA	

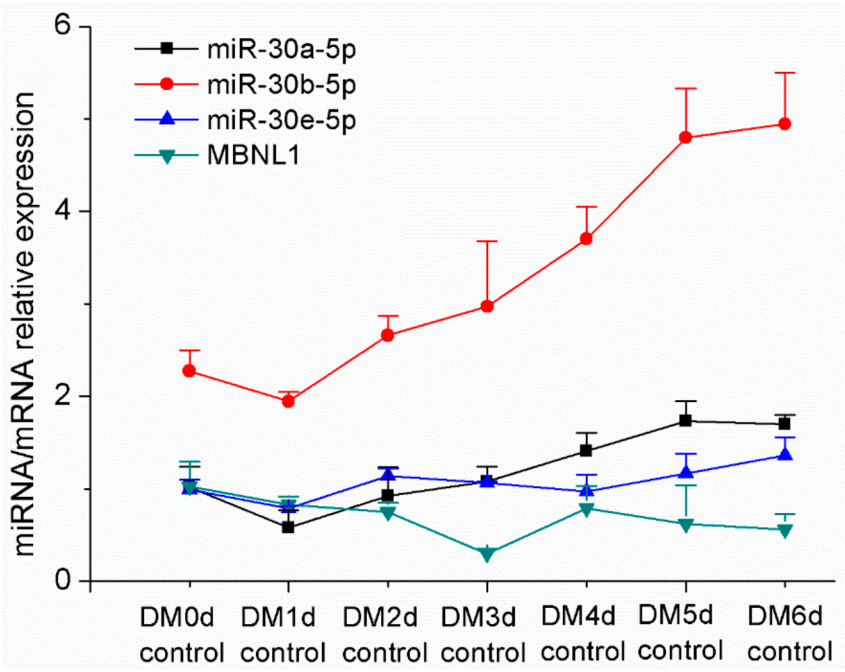

Figure S1. The expression of these miR-30-5p and MBNL1 in non-transfected C2C12 cells.

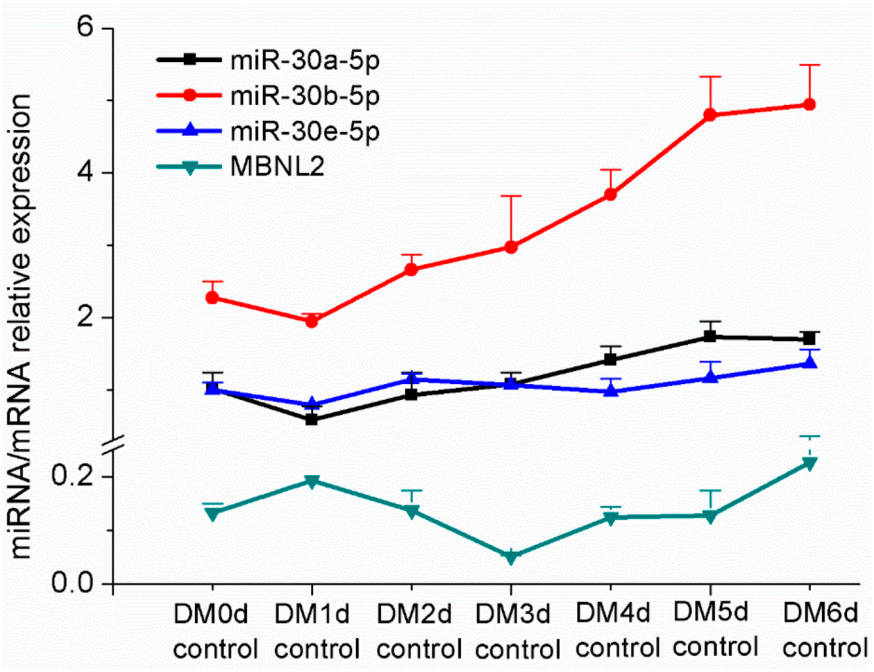

Figure S2. The expression of these miR-30-5p and MBNL2 in non-transfected C2C12 cells.

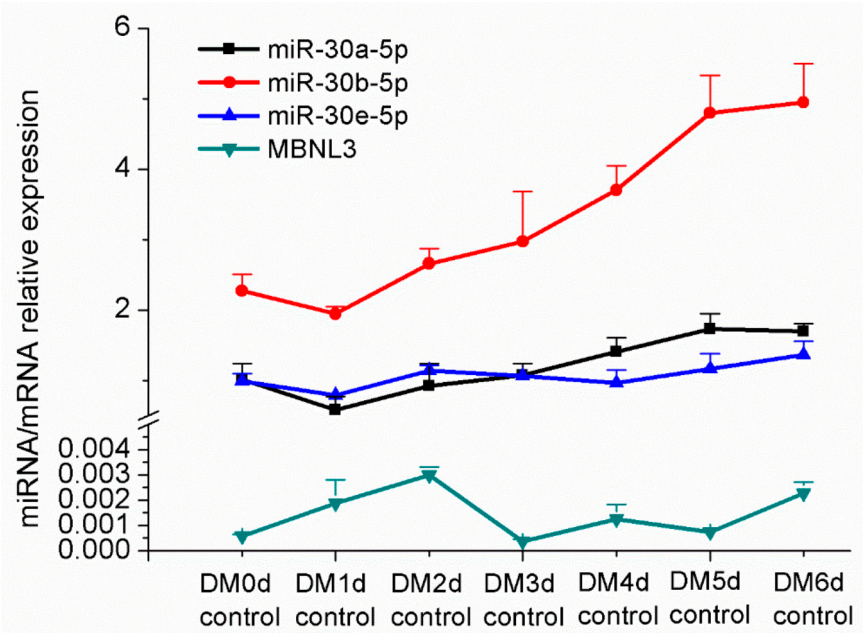

Figure S3. The expression of these miR-30-5p and MBNL3 in non-transfected C2C12 cells.