

Supplementary material

The requirement of Mettl3-promoted *MyoD* mRNA maintenance in proliferative myoblasts for skeletal muscle differentiation

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Figure S1

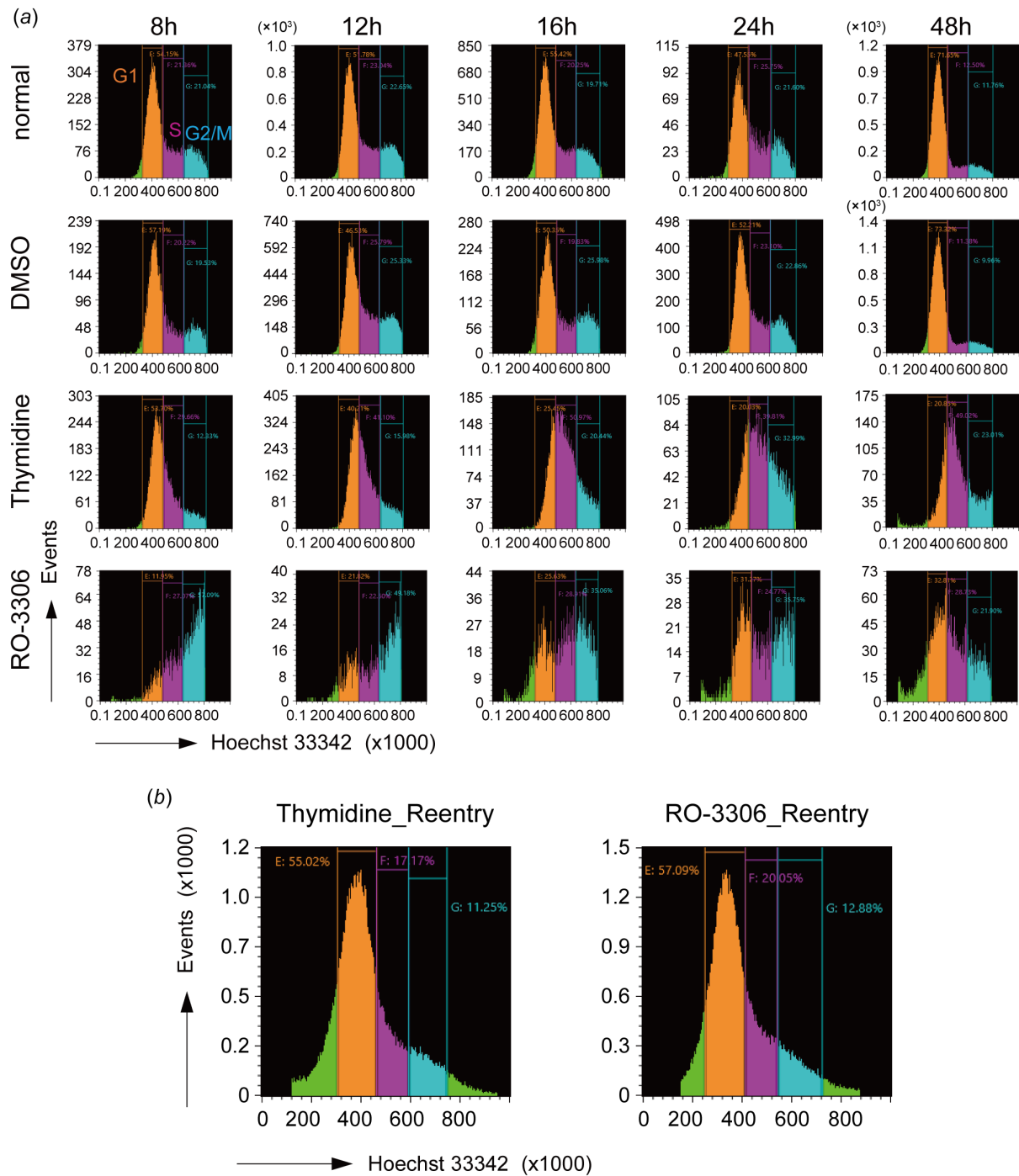


Figure S1. Cell treatments with thymidine and RO-3306. (a) Population changes of C2C12 cells in each cell cycle phase after incubation with thymidine, RO-3306, or DMSO. (b) Population of C2C12 cells treated with thymidine or RO-3306 in each cell cycle 24 h after re-entry into the cell cycle.

Figure S2

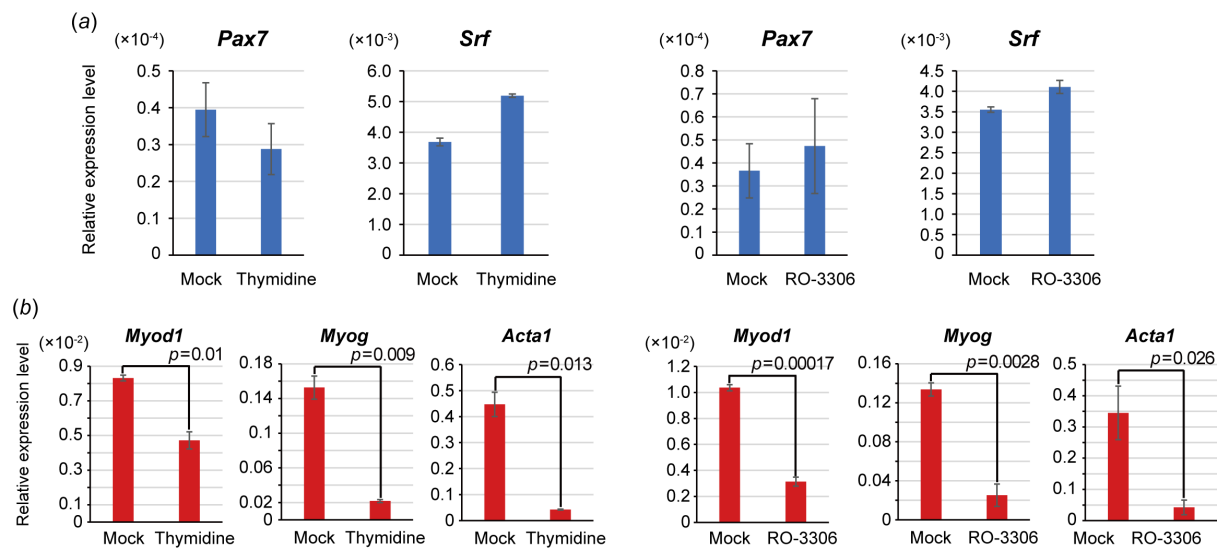


Figure S2. mRNA levels of skeletal muscle-specific factors upon cell cycle arrest. (a) qRT-PCR analysis of *Pax7* and *Srf* in undifferentiated C2C12 cells treated with thymidine (left panels) and RO-3306 (right panels) for 48 h. Data represent the mean \pm SD (n=3). (b) qRT-PCR analysis of *Myod1*, *Myog*, and *Acta1* in C2C12 cells treated with thymidine (left panels) or RO-3306 (right panels) at 48 h after the induction of differentiation.

Figure S3

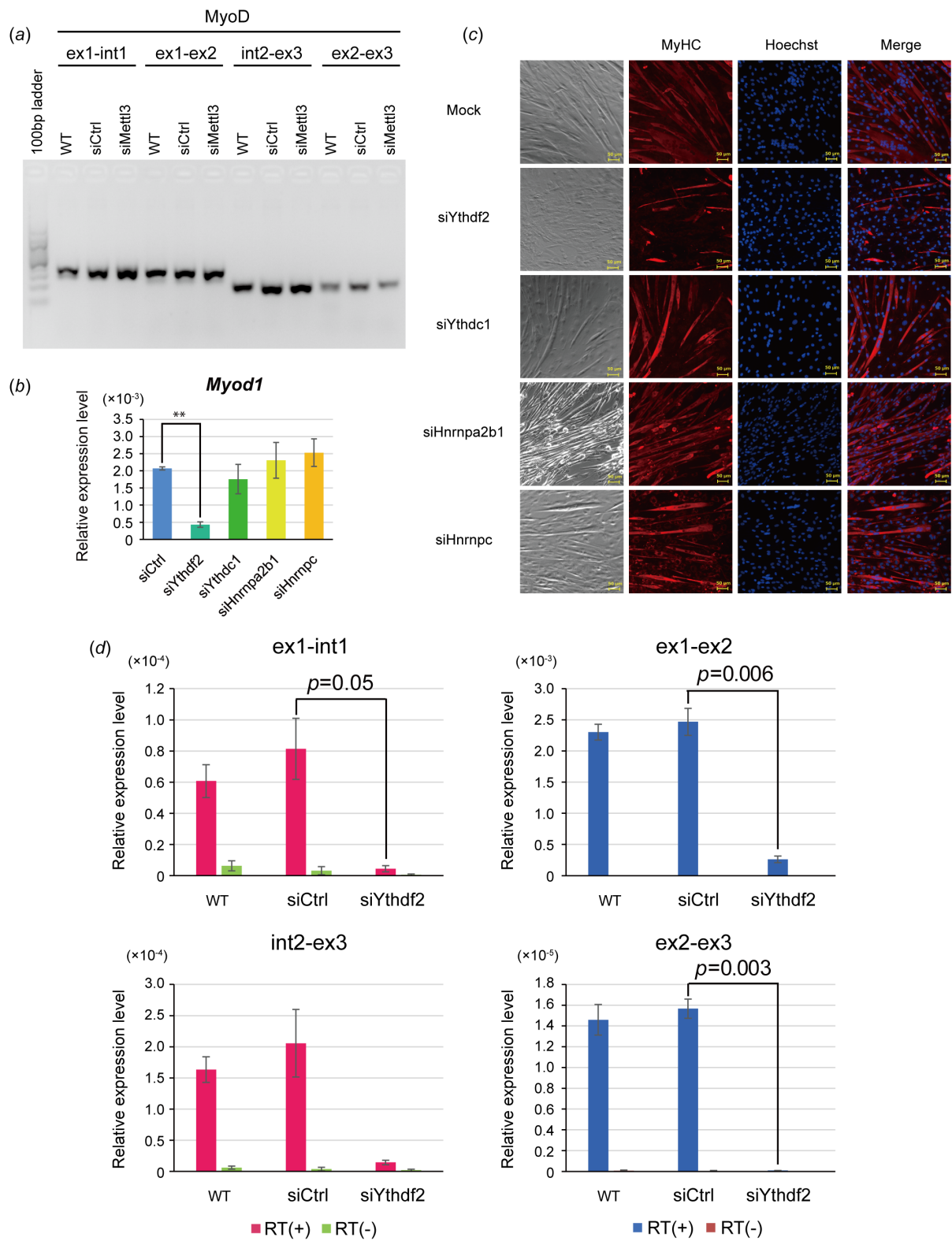


Figure S3. Knockdown of Ythdf2 leads to a decrease of both processed and unprocessed *MyoD* RNA. (a) Agarose gel electrophoresis of PCR products obtained using the four *MyoD* primer sets described in figure 3a. ex: exon, int: intron. (b) qRT-PCR analysis of *Myod1* in undifferentiated C2C12 cells after knockdown of Ythdf2, Ythdc1, Hnrnpa2b1, or Hnrnpc. Data are presented as the mean \pm SD (n=3). (c) Morphology of C2C12 cells after knockdown of Ythdf2, Ythdc1, Hnrnpa2b1, or Hnrnpc. Cells were immunostained for anti-MyHC and Hoechst at 72 h after the induction of differentiation. (d) qRT-PCR analysis of *Myod1* in Ythdf2 knockdown C2C12 cells using the four primers described in figure 3a. To confirm the absence of DNA contamination, each sample was also analysed by qRT-PCR without reverse transcriptase. RT: reverse transcriptase. Error bars show \pm SD (n=3). *P*-value versus control siRNA.

Figure S4

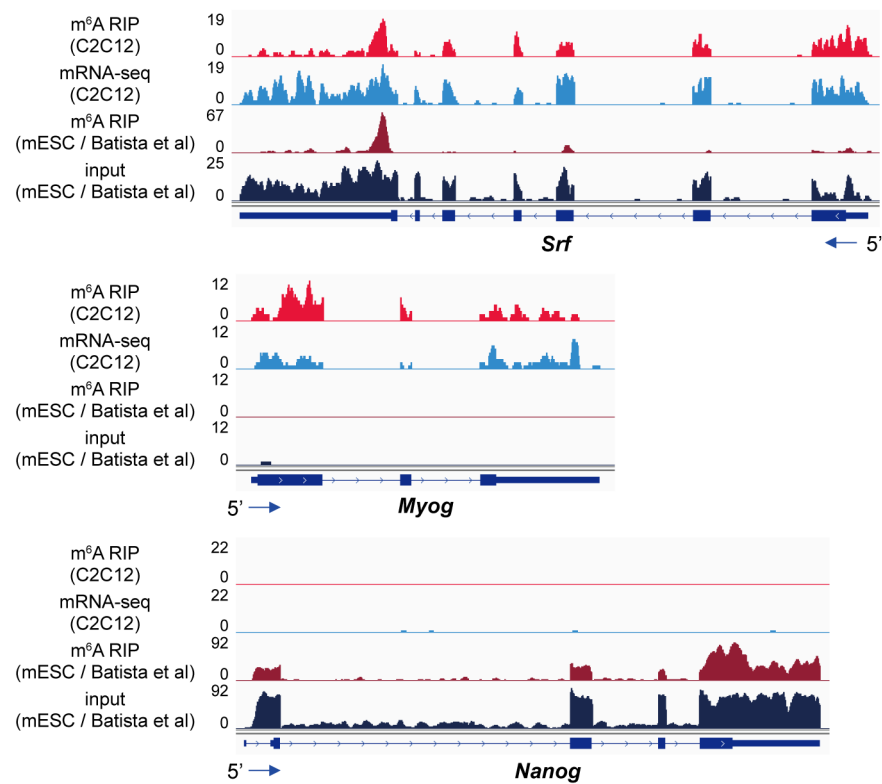


Figure S4. m⁶A-seq analysis using C2C12 and mouse embryonic stem cells. Integrative Genomics Viewer tracks displaying reads coverage of *Myog*, *Srf*, and *Nanog* in m⁶A-seq of total RNA and mRNA-seq data of C2C12 cells and mouse embryonic stem cells (mESC) (SRR1207291 (input), SRR1207292 (m⁶A RIP), Batista et al. 2014).

List of primers for qRT-PCR	
Myod1_F	CAACCCACAGAACCTTTGTCATTG
Myod1_R	GTCGAAACACGGGTCATCATAGAA
Gapdh_F	TTCCTACCCCCAATGTGTCCGTGG
Gapdh_R	GTGGGAGTTGCTGTTGAAGTCGCA
Eef1a1_F	CTCTGACTACCCTCCACTTGGTCG
Eef1a1_R	ATTAAGACTGGGGTGGCAGGTGTT
Myog_F	AAAGCCATCACTTCTGTAGCAGGG
Myog_R	TCTCTGGACTCCATCTTTCTCTCC
Acta1_F	TCCTCAGGACGACAATCGACAATC
Acta1_R	TTCCTTTCCACAGGGCTTTGTTTG
Mettl3_F	CTCTGGGCACTTGGATTTAAGGAA
Mettl3_R	AGCAACTTCTTCTCTAACTCAGGG
Elavl1_F	AGAACATGACCCAAGAGGAACTAC
Elavl1_R	GACACCTTAATGGTTTTGGACTGG
Srf_F	ATGAGTATTAGCTGACCCGATGGG
Srf_R	GGCAGACACAAAGCTAACCAGAGG
Pax7_F	TCTCGGGCCAGACAAAATTGCTGC
Pax7_R	ATGGGGTGGCTTAGAAACAGTGCC
Myod1_ex1_Fw	AGCAAAGTGAATGAGGCCTTCGAG
Myod1_int1_Rv	ATCTAGGTATAAGGGACACCCCA
Myod1_ex2_Rv	TCACTGTAGTAGGCGGTGTCGTAG
Myod1_int2_Fw	ATTACTAACCTTCCACTCCCCTCA
Myod1_ex2_Fw	GCTACGACACCGCCTACTACAGTG
Myod1_ex3_Rv	CCTGTTCTGTGTCGCTTAGGGATG
Myod1_exon1_CDS_F	CCGCCTGAGCAAAGTGAATGAGGC
Myod1_exon1_CDS_R	CCTTCGATGTAGCGGATGGCGTTG
List of primers for cloning	
Myod1_full length_F	CGGAATCCCCTACTACACTCCTATTGGCTTGAGGC
Myod1_full length_R	CGCAATTGCCCTTCTTCATTACCCAGCACTACCCAG
Myod1_5'UTR_deletion_F	CGGAATTCATGGAGCTTCTATCGCCGCCAC
Myod1_5'UTR_deletion_R	CGGAATTCGCCACTTTGTATAAATTAGCGTC
Myod1_Mut1_F	TGACAGGCCAGGCCAGGGAGGAGGGGTAGA
Myod1_Mut1_R	TCCCTGGCCTGGCCTGTCAGAGGTGTGGTG
Myod1_Mut2_F	GTAGAGGCCAGCCGGTGTGCATTCCAACCC
Myod1_Mut2_R	CGGCTGGCCTCTACCCCTCCTCCCTGTCCT
Myod1_6bp-del_F	CACACCTCTGACAGGCAGGGAGGAGGGGTA
Myod1_6bp-del_R	TACCCCTCCTCCCTGCCTGTCAGAGGTGTG
Myod1_3'UTR_deletion_F	CGGAATCCCCTACTACACTCCTATTGGCTTGAGGC
Myod1_3'UTR_deletion_R	CGGAATTCTCAAAGCACCTGATAAATCGCATTGGGG
Myod1_CDS_F	CGTCTAGAATGGAGCTTCTATCGCCGCCACTCC
Myod1_CDS_R	CGGAATTCTCAAAGCACCTGATAAATCGCATTG

Table S1. List of primers used for RT-PCR and cloning.