

Supplementary Materials

MicroRNA-24-3p promotes skeletal muscle differentiation and regeneration by regulating HMGA1

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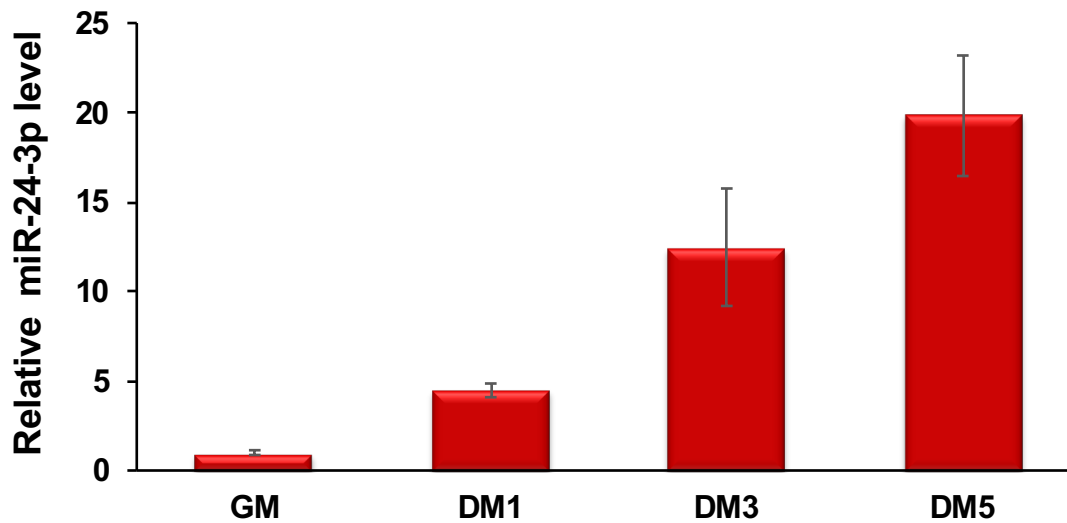
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Transfection	MYOG-positive cell number			MHC-positive cell number		
	Total counts (10 fields)	Percentage (%)		Total counts (10 fields)	Percentage (%)	
		Mean	SD		Mean	SD
NC	291	100	24.65	256	100	11.82
miR-24	554	197.78	41.19	542	213.83	27.72
Ant-NC	259	100	13.78	330	100	22.04
Ant-24	123	48.22	8.28	179	55.38	10.52

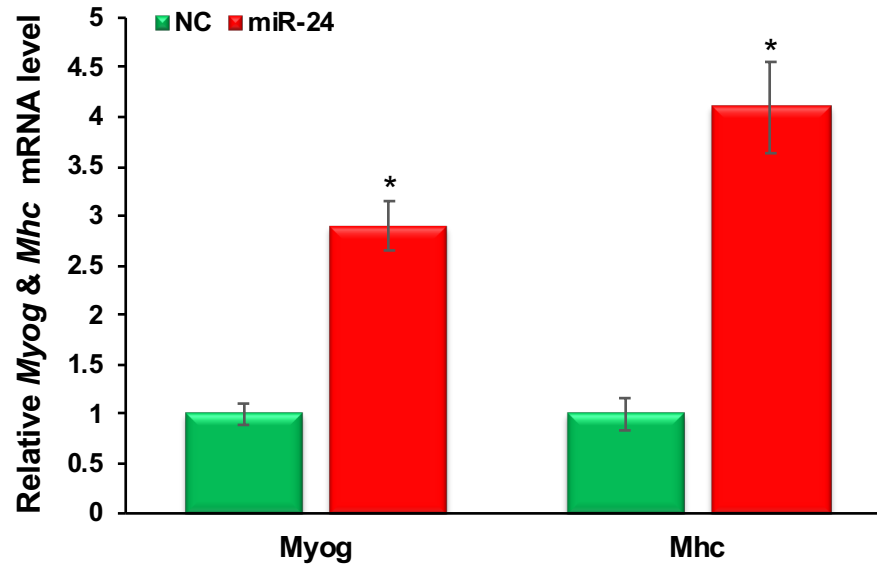
Suppl Table 1: miR-24-3p increases and Ant-24 decreases number of MYOG- and MHC-positive cells. Primary myoblast cells were transfected with mature miR-24-3p or GL2 as an NC twice at 24-hour intervals in GM, then replaced the GM with DM. Immunostaining on these cells was carried out for an early myogenic marker, myogenin (MYOG), and a late myogenic marker, myosin heavy chain (MHC) 24 and 48 hours after adding the DM, respectively. In a reciprocal experiment, primary myoblast cells were transfected twice with antagomirs specific to miR-24-3p (Ant-24) or GL2 (Ant-NC) at 24-hour intervals in GM, then replaced the GM with DM. Immunostaining on these cells was carried out for MYOG and MHC 24 and 48 hours after adding the DM, respectively. Fractions of MYOG- and MHC-positive cells were determined from 10 random fields, and data are presented relative to the NC or Ant-NC control as 100%.

Oligos	Sequence
U6sn primer	CTGCGCAAGGATGACACGCA
miR-24 primer	TGGCTCAGTTCAGCAGGAACAG
miR-192 primer	CTGACCTATGAATTGACAGCC
Gapdh Forward primer	ATGACATCAAGAAGGTGGTGAAGC
Gapdh Reverse primer	GAAGAGTGGGAGTTGCTGTTGAAG
Rps13 Forward primer	TGACGACGTGAAGGAACAGATT
Rps13 Reverse primer	ATTTCCAGTCACAAAACGGACCT
Myog Forward primer	AGCGCAGGCTCAAGAAAGTGAATG
Myog Reverse primer	CTGTAGGCGCTCAATGTACTGGAT
Mhc Forward primer	TCCAAACCGTCTCTGCACTGTT
Mhc Reverse primer	AGCGTACAAAGTGTGGGTGTGT
Hmga1 Forward primer	CGACCAAAGGGAAGCAAGAATAA
Hmga1 Reverse primer	TCCTCTTCCTCCTTCTCCAGTTTC
Id3 Forward primer	CTCTTAGCCTCTTGGACGACATGA
Id3 Reverse primer	TGTAGTCTATGACACGCTGCAGGA
miR-24	UGGCUCAGUUCAGCAGGAACAG
miR-24Mut	UCGCACUGUUCUGCAGGAACAG
GL2	UCGAAGUAUJCCGCGUACG
Ant-24	mU(*)mG(*)mGmCmUmCmAmGmUmUmCmAmGmCmAmGmGmA mAmCmA(*)mG(*) (3'-Chl)
Ant-NC	mU(*)mA(*)mUmCmGmCmGmAmGmUmAmCmGmUmCmGmAmG (*)mG(*)mC(*)mC(*) (3'-Chl)

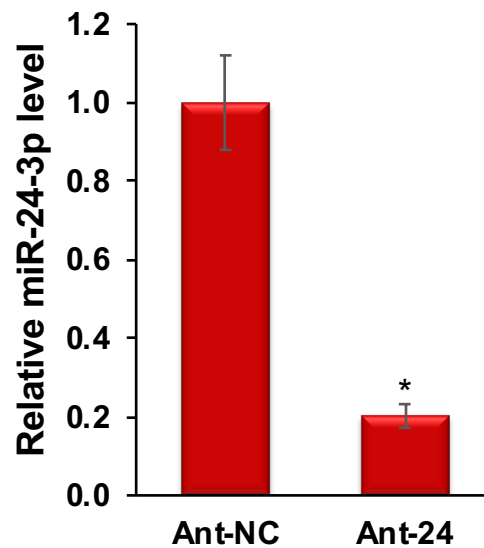
Suppl Table 2: List of oligos used in this study.



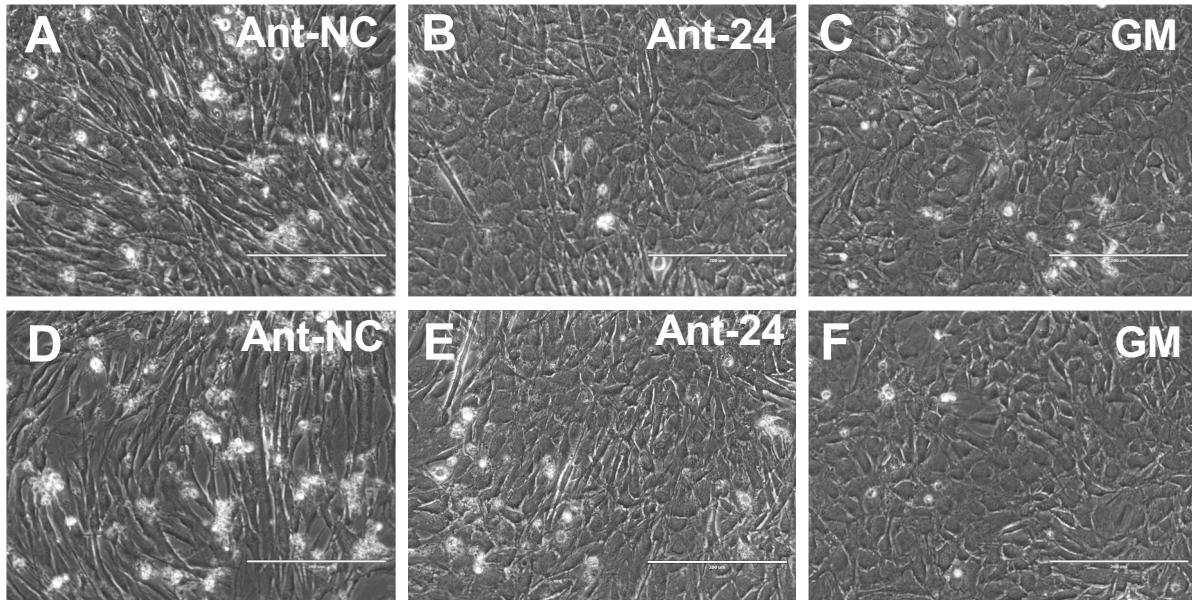
Suppl Fig 1: miR-24-3p is upregulated during C2C12 myoblast differentiation. qRT-PCR analyses show that miR-24-3p is upregulated during C2C12 myoblast differentiation. miR-24-3p values were first normalized to the U6sn values, and fold-change of miR-24-3p was determined relative to the undifferentiated myoblasts (GM). DM1, DM3, and DM5 indicates the number of days C2C12 myoblasts cultured in differentiation medium (DM). The values are expressed as Mean \pm SD of biological triplicates.



Suppl Fig 2: miR-24-3p promotes C2C12 myoblast differentiation. C2C12 myoblast cells were transfected twice at 24-hour intervals with GL2 negative control (NC) or miR-24-3p in GM. The cells were harvested after another 48 hours. qRT-PCR was carried out for *Myog* and *Mhc*, and the values were normalized to *Gapdh*, then again to the values of the NC samples. The values are expressed as Mean \pm SD of biological triplicates. *P < 0.001.



Suppl Fig 3: miR-24-3p level is downregulated in the Ant-24-transfected primary myoblasts. We transfected primary myoblast cells twice with antagomirs specific to miR-24-3p (Ant-24) or GL2 control (Ant-NC) at 24-hour intervals in GM and harvested the samples after another 48 hours. We performed qRT-PCR analyses for miR-24-3p. miR-24-3p values were normalized to the respective U6sn values. The values are expressed as Mean \pm SD of biological triplicates. *P < 0.001.



Suppl Fig 4: Ant-24 represses primary myoblast differentiation. We transfected primary myoblast cells twice with antagomirs specific to miR-24-3p (Ant-24) or GL2 control (Ant-NC) at 24-hour intervals in GM and replaced the GM with DM. We followed up these cells for 5 days. (A, D) Ant-NC-treated cells differentiate normally and forms multinucleated myotubes. (B, E) Ant-24-treated cells do not differentiate following the normal kinetics, and the majority of the cells looks like (C, F) proliferating myoblasts grown in GM. Scale Bar: 200 μ M.

position 311
 target 5' U C UUC U 3'
 CUG CCUGCUG CUG GCU
 GAC GGACGAC GAC CGG
 miRNA 3' AA UU U U 5'

position 182
 target 5' C GCCC C C ACACCU C 3'
 CUGU UCCU C GAAC GCC
 GACA AGGA G CUUG CGG
 miRNA 3' C A ACU U 5'

position 1087
 target 5' U GUGGG GA G G 3'
 UGUUCU G GGG UG GCCA
 ACAAGG C CUU AC CGGU
 miRNA 3' G A GA G U 5'

position 500
 target 5' C CCCACCCU A A U 3'
 UCCUGC AC GAG UCA
 AGGACG UG CUC GGU
 miRNA 3' GACA ACU A 5'

position 897
 target 5' A UCA C U U 3'
 UGU UCCU CUGAAUU GCC
 ACA AGGA GACUUGA CGG
 miRNA 3' G C CU U 5'

position 78
 target 5' N CCCA C 3'
 CCUGC U GGCC
 GGACGA UCGG
 miRNA 3' GACAA CUUGAC U 5'

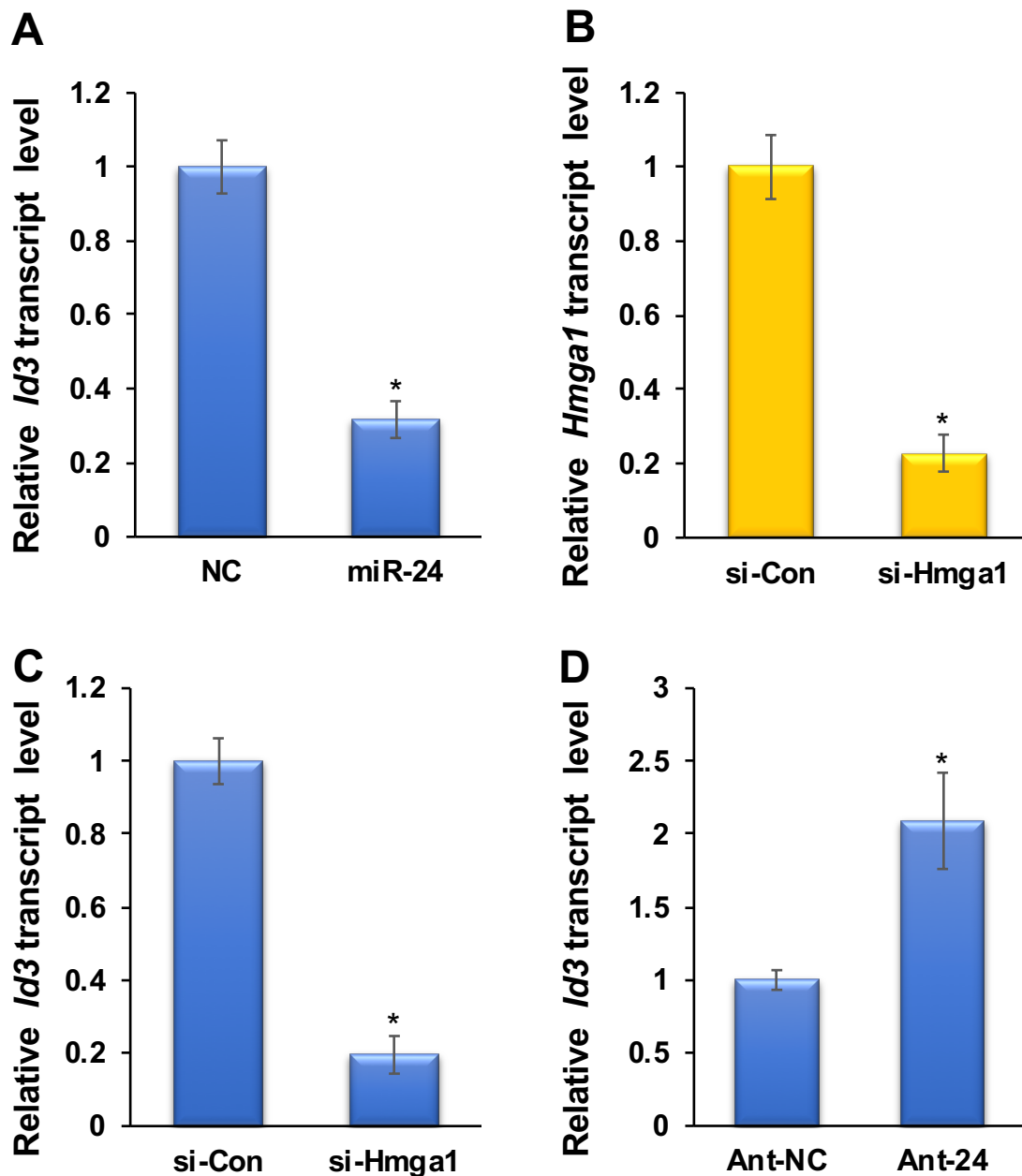
position 1018
 target 5' C GGA C A G G 3'
 GU UGCUG AGC GAGU A
 CA ACGAC UUG CUCG U
 miRNA 3' GA AGG A G 5'

position 955
 target 5' A GG G C 3'
 G GGGCUGGG CG
 C CUUGACUC GU
 miRNA 3' GACAAGGA GA G 5'

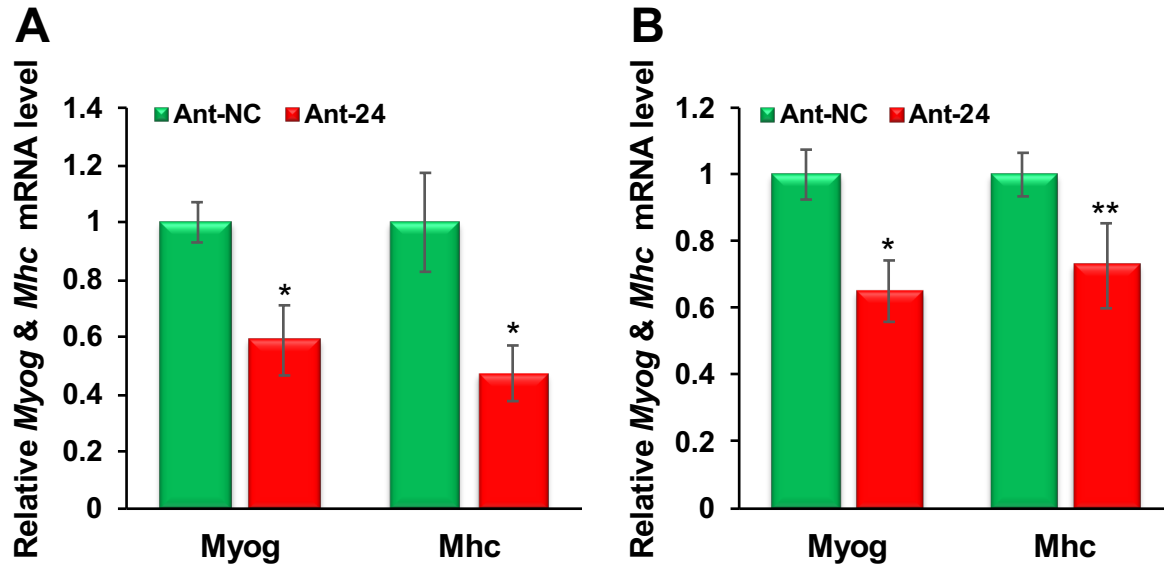
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 UCCUG GAGC GGCC
 AGGAC CUUG UCGG
 miRNA 3' GACA GA AC U 5'

position 141
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 GAC AGG CG C UG CUCG
 miRNA 3' A A A U A GU 5'

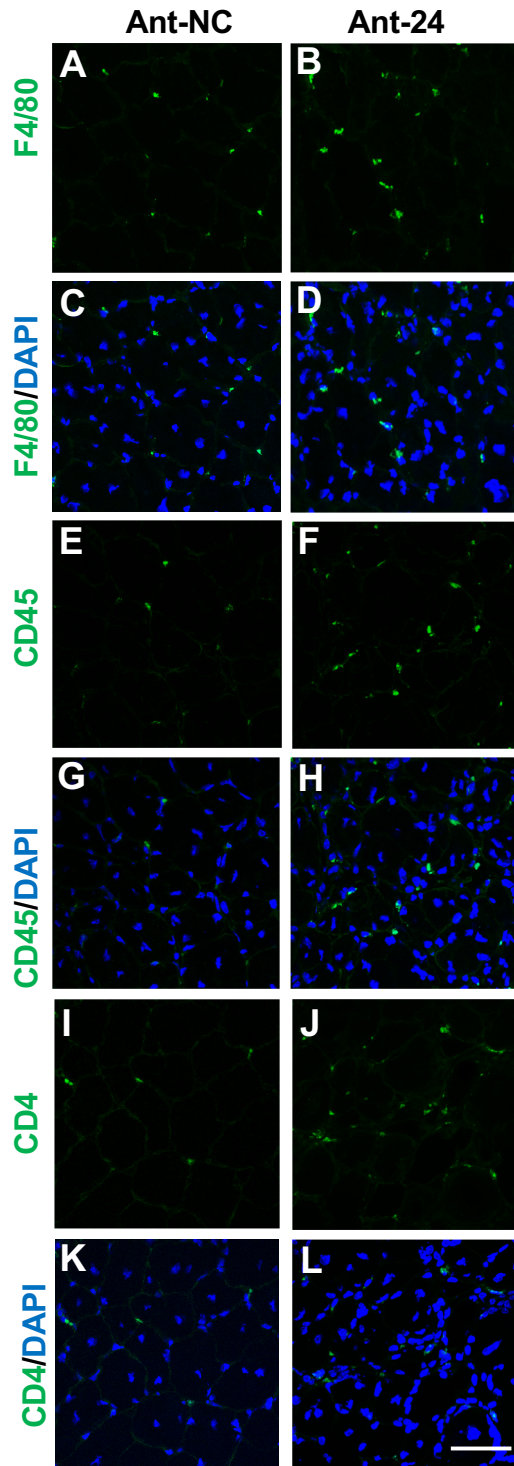
Suppl Fig 5: RNA hybrid microRNA target prediction algorithm identifies non-seed match miR-24-3p target sites in the 3'UTR of *Hmga1*. We have identified eighteen non-seed match miR-24-3p target sites in the 3'UTR of *Hmga1*. Here, we have shown the top ten such miR-24-3p target sites.



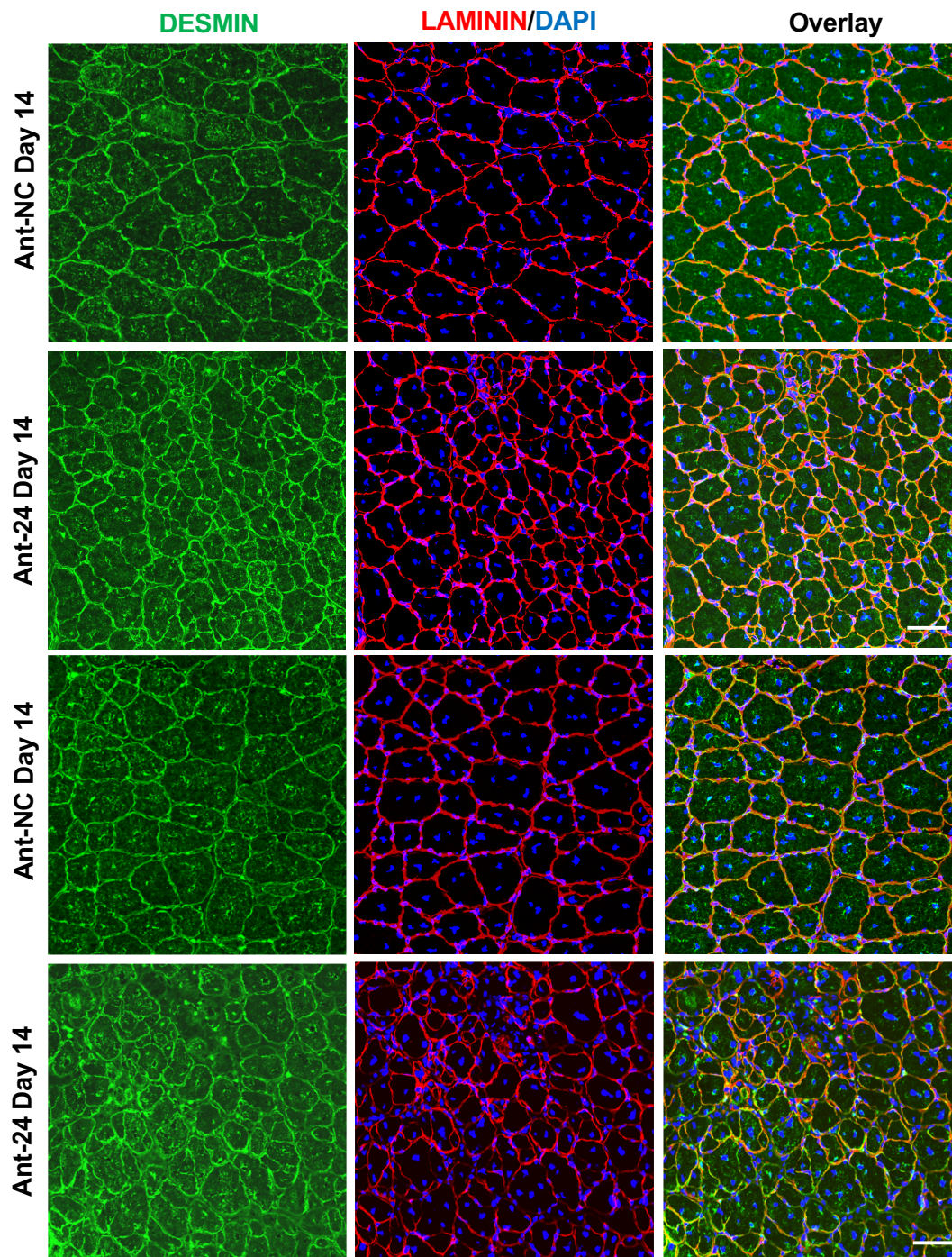
Suppl Fig 6: miR-24-3p regulates HMGA1's direct downstream target *Id3* transcript level during myoblast differentiation. (A) *Id3* transcript level is downregulated in miR-24-3p-transfected myoblasts. (B) *Hmga1* siRNA decreases *Hmga1* transcript level, and subsequently (C) decreases *Id3* transcript level. (D) Ant-24 derepresses *Id3* transcript level during myoblast differentiation. qRT-PCR was carried out for *Id3* or *Hmga1* and the values were normalized to the respective *Gapdh* values, then again to the values of the NC samples. The values are expressed as Mean \pm SD of biological triplicates. *P < 0.001.



Suppl Fig 7: *Myog* and *Mhc* levels are downregulated in the undifferentiated and differentiated MuSCs isolated from Ant-24-injected neonatal mice. (A) *Myog* and *Mhc* mRNA levels are downregulated in the (A) undifferentiated and (B) differentiated (in DM for 48 hours) MuSCs. *Myog* and *Mhc* values were normalized with the respective RSP13 values, then again to the values of the Ant-NC samples. The values are expressed as Mean \pm SD of five neonatal mice. *P < 0.001; **P < 0.01.



Suppl Fig 8: Increased presence of various cells, including inflammatory cells are observed in the Ant-24-injected regenerating skeletal muscle. Representative images of Ant-NC- and Ant-24-injected TA muscle sections immunostained with (A, B) F4/80, (C, D) F4/80 and DAPI, (E, F) CD45, (G, H) CD45 and DAPI, (I, J) CD4, (K, L) CD4 and DAPI. (A, C, E, G, I, K) Anti-NC-injected and (B, D, F, H, J, L) Ant-24-injected TA muscle sections on day 14 post-injury. Scale Bar: 50 μ M.



Suppl Fig 9: Ant-24 impairs skeletal muscle regeneration after injury. Additional representative images of Ant-NC- and Ant-24-injected TA muscle sections stained with DESMIN and LAMININ on day 14 post-injury. Ant-NC and Ant-24-injected samples are indicated. DESMIN (Green), LAMININ (red) and DAPI (Blue). Scale Bar: 50 μ M.