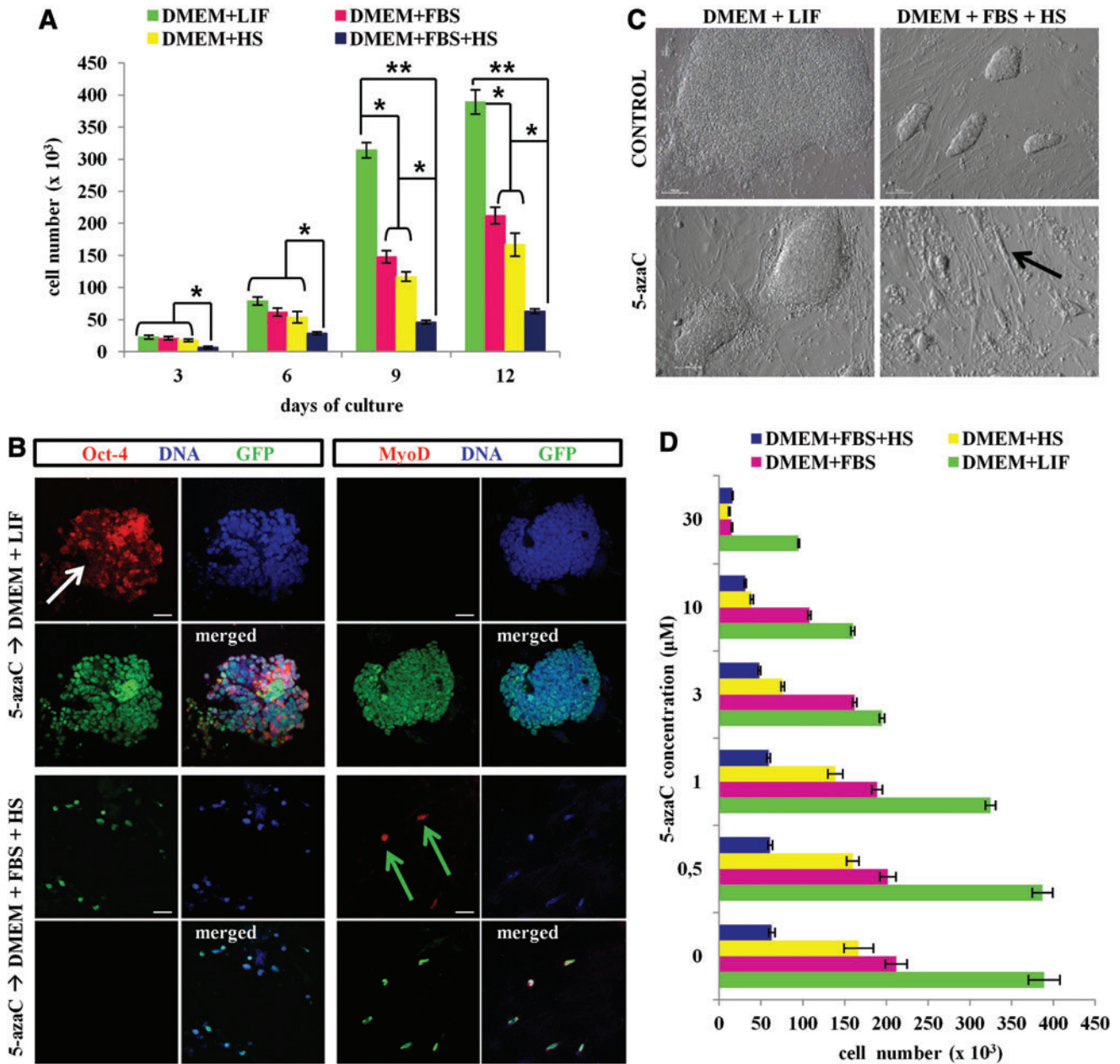
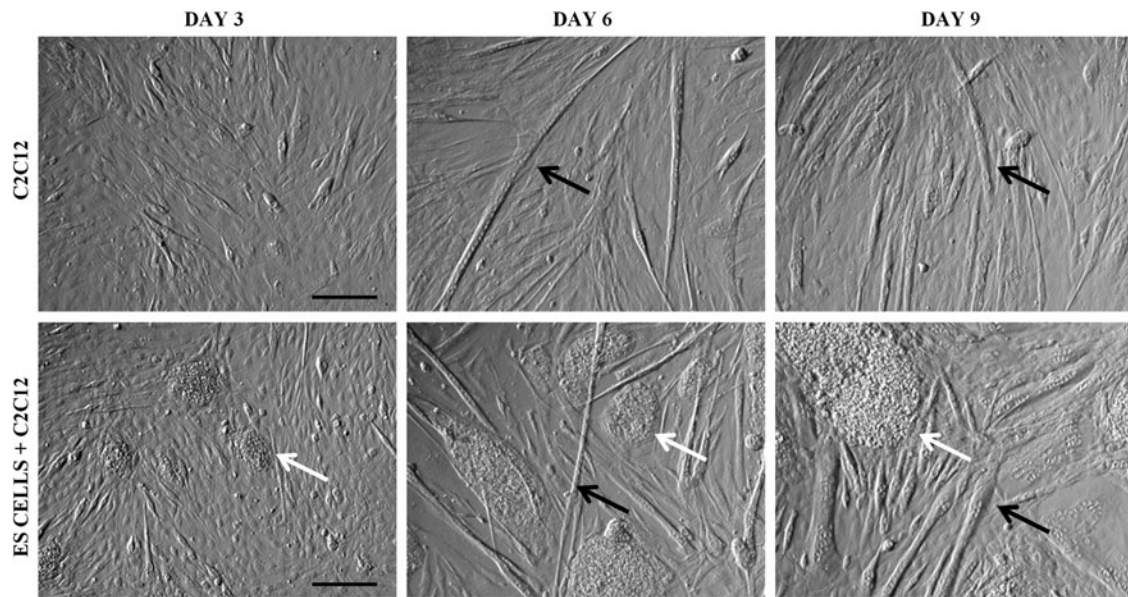


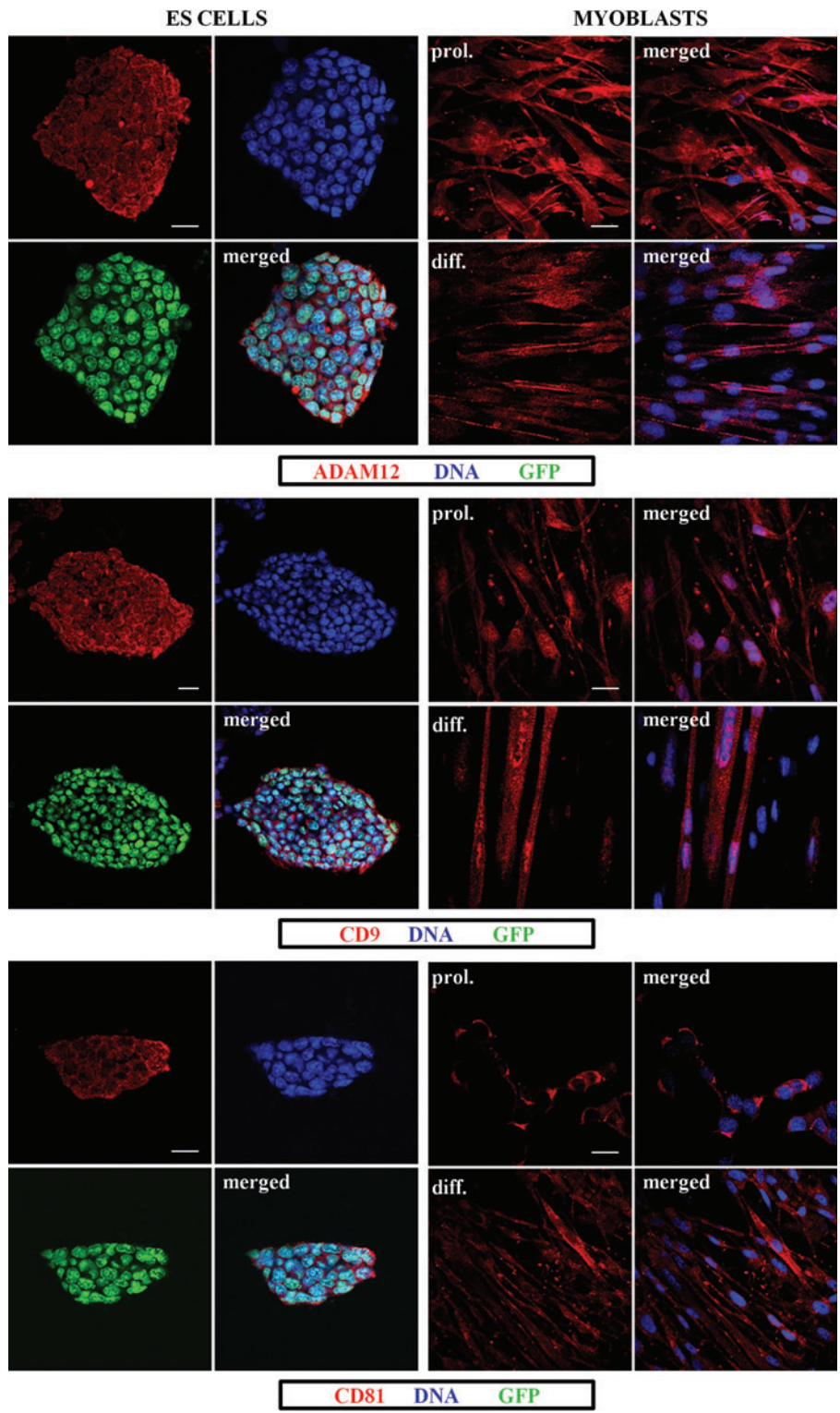
## Supplementary Data



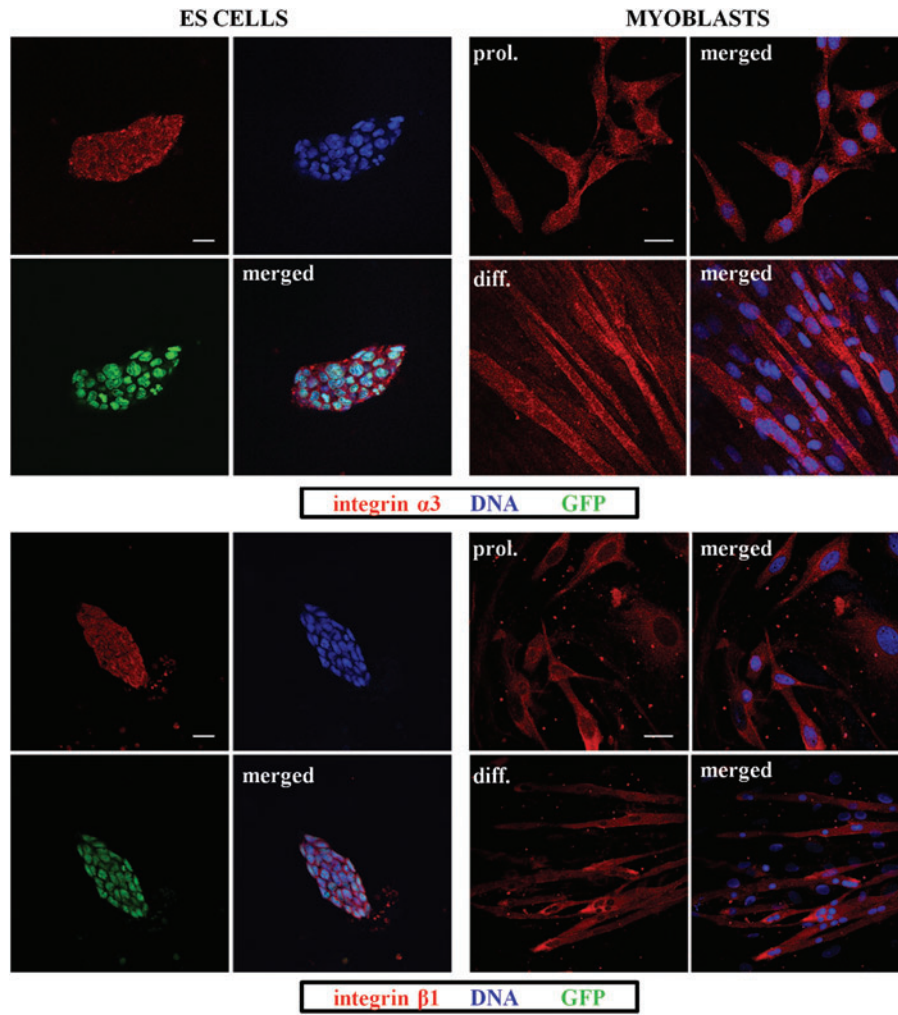
**SUPPLEMENTARY FIG. S1.** Behavior of embryonic stem (ES)-green fluorescent protein (GFP) cells in different culture conditions. (A) Number of ES-GFP cells after 3, 6, 9, and 12 days of culture in different media; \* $P < 0.05$ ; \*\* $P < 0.01$ . Three independent experiments were performed. (B) Immunofluorescent detection of Oct-4 and MyoD in ES-GFP cells pretreated with 10  $\mu\text{M}$  5-azaC and then cultured in DMEM+LIF or DMEM+FBS+HS for 12 days. White arrow indicates cells devoid of pluripotency marker Oct-4; green arrows indicate nuclei positive for MyoD. Scale bar, 20  $\mu\text{m}$ . (C) Morphology of ES-GFP cell colonies pretreated with 5-azaC and then cultured in DMEM+LIF or DMEM+FBS+HS for 12 days. In control cultures, incubation with 5-azaC was omitted. Black arrow indicates one of the elongated cells observed among ES cells treated with 5-azaC and cultured in DMEM+FBS+HS (or DMEM+HS). Scale bar, 100  $\mu\text{m}$ . (D) The number of ES-GFP cells pretreated with different concentrations of 5-azaC and then cultured in different types of media for 12 days. Three independent experiments were performed.



**SUPPLEMENTARY FIG. S2.** Co-culture of ES cells and myoblasts. Morphology of ES-GFP cells and C2C12 myoblasts after 3, 6, and 9 days of co-culture (*lower row*). *White arrows* indicate ES cells colonies; *black arrows* indicate myotubes. A control myoblast culture is shown in the *upper row*. Scale bar, 100  $\mu\text{m}$ .



**SUPPLEMENTARY FIG. S3.** Immunolocalization of ADAM12, CD9, and CD81 in ES-GFP cells and myoblasts. “prol.” indicates images showing localization of analyzed proteins in proliferating C2C12 or SC-derived myoblasts; “diff.” indicates images showing presence of proteins in differentiating myoblasts. Scale bar, 20  $\mu$ m.



**SUPPLEMENTARY FIG. S4.** Immunolocalization of integrin  $\alpha$ 3 and  $\beta$ 1 in ES-GFP cells and myoblasts. “prol.,” marks images showing localization of analyzed proteins in proliferating C2C12 or SC-derived myoblasts; “diff.,” indicates images showing presence of proteins in differentiating myoblasts. Scale bar, 20  $\mu$ m.

SUPPLEMENTARY TABLE S1. PRIMERS USED IN REVERSE TRANSCRIPTION-POLYMERASE CHAIN REACTION

<i>Gene</i>	<i>Sequence</i>	<i>Product size (bp)</i>	<i>Annealing temperature (°C)</i>	<i>References</i>
<i>Oct-4</i>	5' GAAGTTGGAGAAGGTGGAACC 3'	450	55	1
	5' AACCATCCTTCTCTAGCCC 3'			
	5' GCGTTCCTTTGGAAAGGTGTT 3'	312	55	2
	5' CTCGAACCACATCCTTCTCT 3'			
<i>Nanog</i>	5' AGGGTCTGCTACTGAGATGCTCTG 3'	363	59	3
	5' CAACCACTGGTTTTTCTGCCACCG 3'			
	5' CAGGTGTTTGAGGGTAGCTC 3'	222	52	4
	5' CGGTTCATCATGGTACAGTC 3'			
<i>Sox2</i>	5' GGCGGCAACCAGAAGAACAG 3'	414	55	5
	5' GTTGCTCCAGCCGTTTCATGTG 3'			
<i>Pax3</i>	5' GCTGTCTGTGATCGGAACACT 3'	417	55	6
	5' CTCCAGCTTGTTTCTCCATC 3'			
	5' GCTGTCTGTGATCGGAACACTG 3'	509	62	7
	5' GTCTCCGACAGCTGGTATGTTG 3'			
<i>Pax7</i>	5' CTGGATGAGGGCTCAGATGT 3'	243	52	6
	5' GGTTAGCTCCTGCCTGCTTA 3'			
	5' CAAGAGGTTTATCCAGCCGAC 3'	498	57	7
	5' GAGGGCACCGTGCTTCGGTC 3'			
<i>MyoD</i>	5' ACATAGACTTGACAGGCCCCGA 3'	450	52	8
	5' AGACCTTCGATGTAGCGGATGG 3'			
	5' GCCCGCGCTCCAAGTCTCTGAT 3'	397	59	9
	5' CCTACGGTGGTGCGCCCTCTGC 3'			
<i>Myf-5</i>	5' GAGCCAAGAGTAGCAGCCTTCG 3'	440	54	8
	5' GTTCTTTCGGGACCAGACAGGG 3'			
	5' TGCCATCCGCTACATTGAGAG 3'	353	59	9
	5' CCGGGTAGCAGGCTGTGAGTTG 3'			
<i>Myog</i>	5' CCATCCAGTACATTGAGCGCCTA 3'	550	55	8
	5' GGGGCTCTCTGGACTCCATCTT 3'			
	5' GGGCCCCTGGAAGAAAAG 3'	364	55	9
	5' AGGAGGCGCTGTGGGAGTT 3'			
<i>Mrf4</i>	5' CTGCGCGAAAGGAGGAGACTAAAG 3'	367	55	10
	5' ATGGAAGAAAGGCGCTGAAGACTG 3'			
	5' CTACATTGAGCGTCTACAGGACC 3'	234	55	11
	5' CTGAAGACTGCTGGAGGCTG 3'			
<i>Ncam</i>	5' TGTC AAGTGGCAGGAGATGC 3'	137	52	12
	5' GCGGTTGTAGATGGTGAGGGT 3'			
<i>Vcam-1</i>	5' AACTCTTACCTGTGCGCTGT 3'	304	57	13
	5' ATTTCCCGGTATCTTCAATGG 3'			
<i>Itga3</i>	5' AAGCCAAATCTGAGACTGTG 3'	660	47	14
	5' GTAGTATCGGTCCCAATCT 3'			
<i>Itgb1</i>	5' TGTGGAGACTCCAGACTGTCCTACT 3'	247	57	14
	5' TCATTTCCCTCATACTCGGATT 3'			
<i>Mcad</i>	5' CCACAAACGCCTCCCCTACCCACTT 3'	446	58	10
	5' TCGTCGATGCTGAAGAACTCAGGGC 3'			
<i>Adam12</i>	5' CACGAATCGCTGCTGTAACGCTA 3'	396	49	15
	5' CTCTCAGCTCACATTTGGCGAAGGC 3'			
<i>Cd9</i>	5' GAGCATGCCGGTCAAAGGAGGTAG 3'	685	58	16
	5' TCAGCACATTTCTCGGCTCC 3'			
<i>Cd81</i>	5' AGTACACGGAGCTGTTCCGG 3'	304	52	17
	5' ATGGGAGTGGAGGGCTGCAC 3'			
<i>GAPDH</i>	5' ACTCCACTCACGGCAAATTC 3'	385	59	7]
	5' ACTGTGGTCATGAGCCCTTC 3'			

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