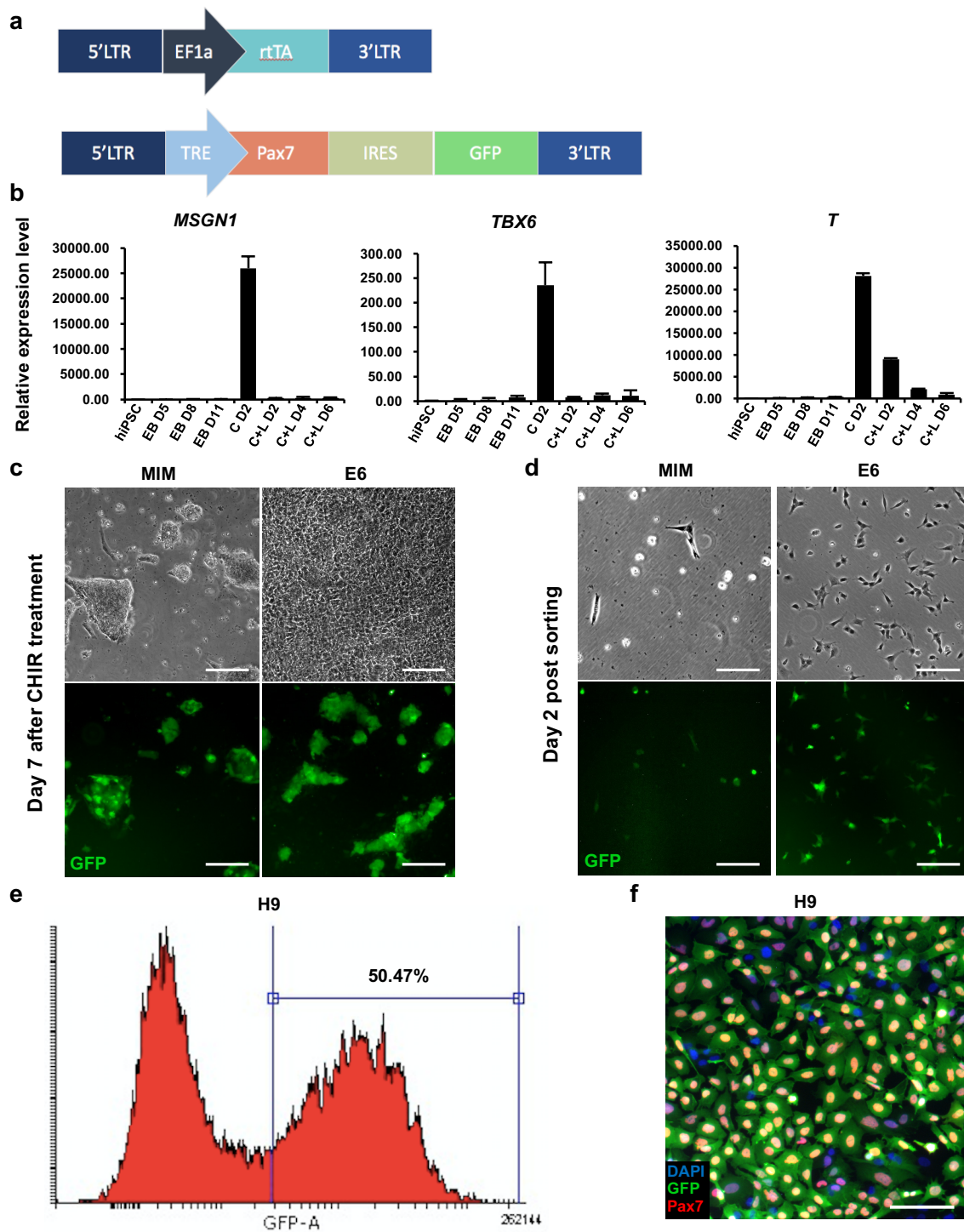
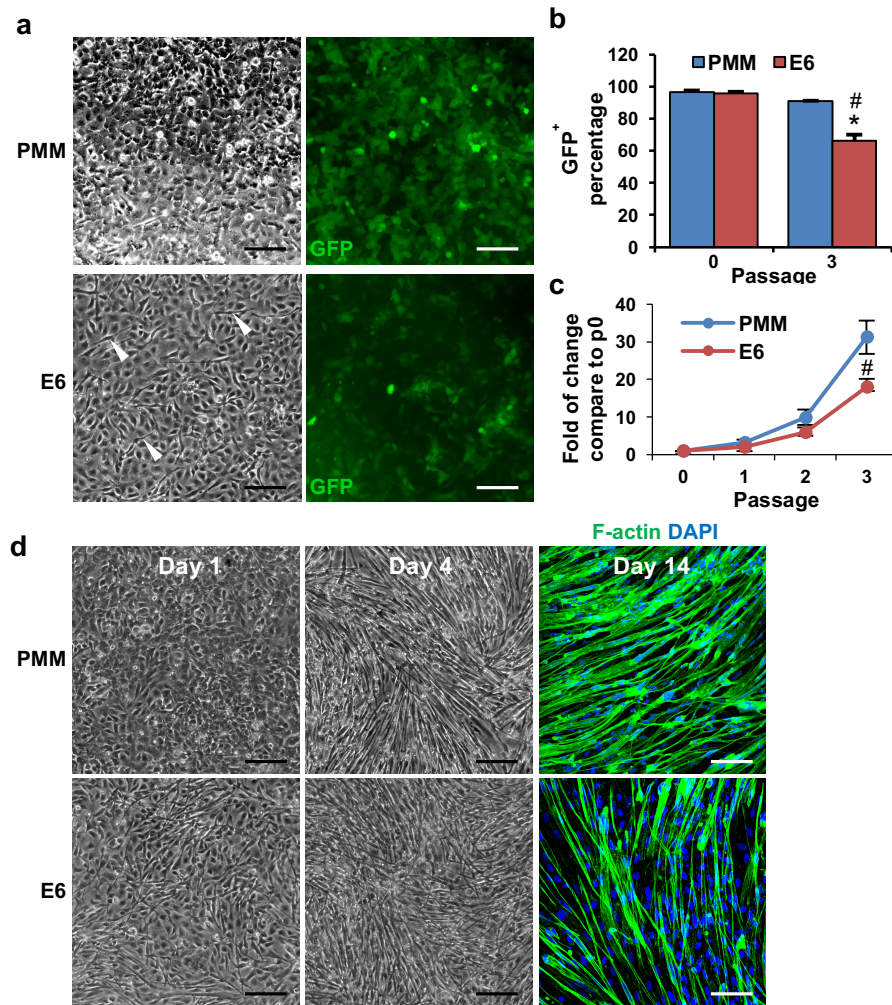


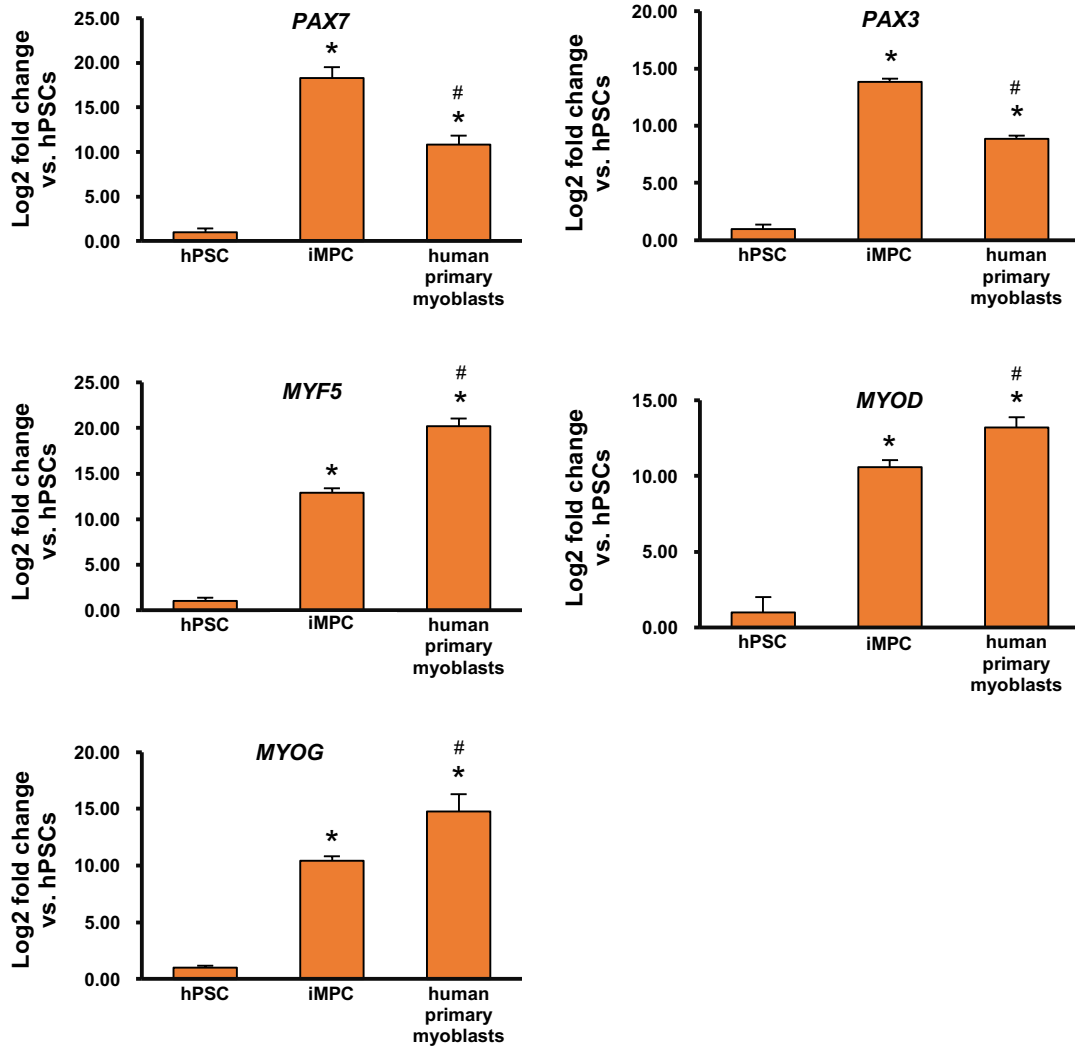
Supplementary Figure 1. Expression of pluripotency markers and karyotype analysis for TRiPSC line. (a) Representative co-staining of Oct4 and DAPI in undifferentiated TRiPSCs. **(b)** Co-staining of Tra-1-81 and DAPI in undifferentiated TRiPSCs. **(c)** Live cell FACS analysis for Tra-1-81⁺ undifferentiated TRiPSCs. Iso, isotype antibody control. **(d)** Alkaline Phosphatase (APS) live staining in undifferentiated TRiPSCs. **(e)** TRiPSCs exhibit normal karyotype. Scale bar, 200µm.



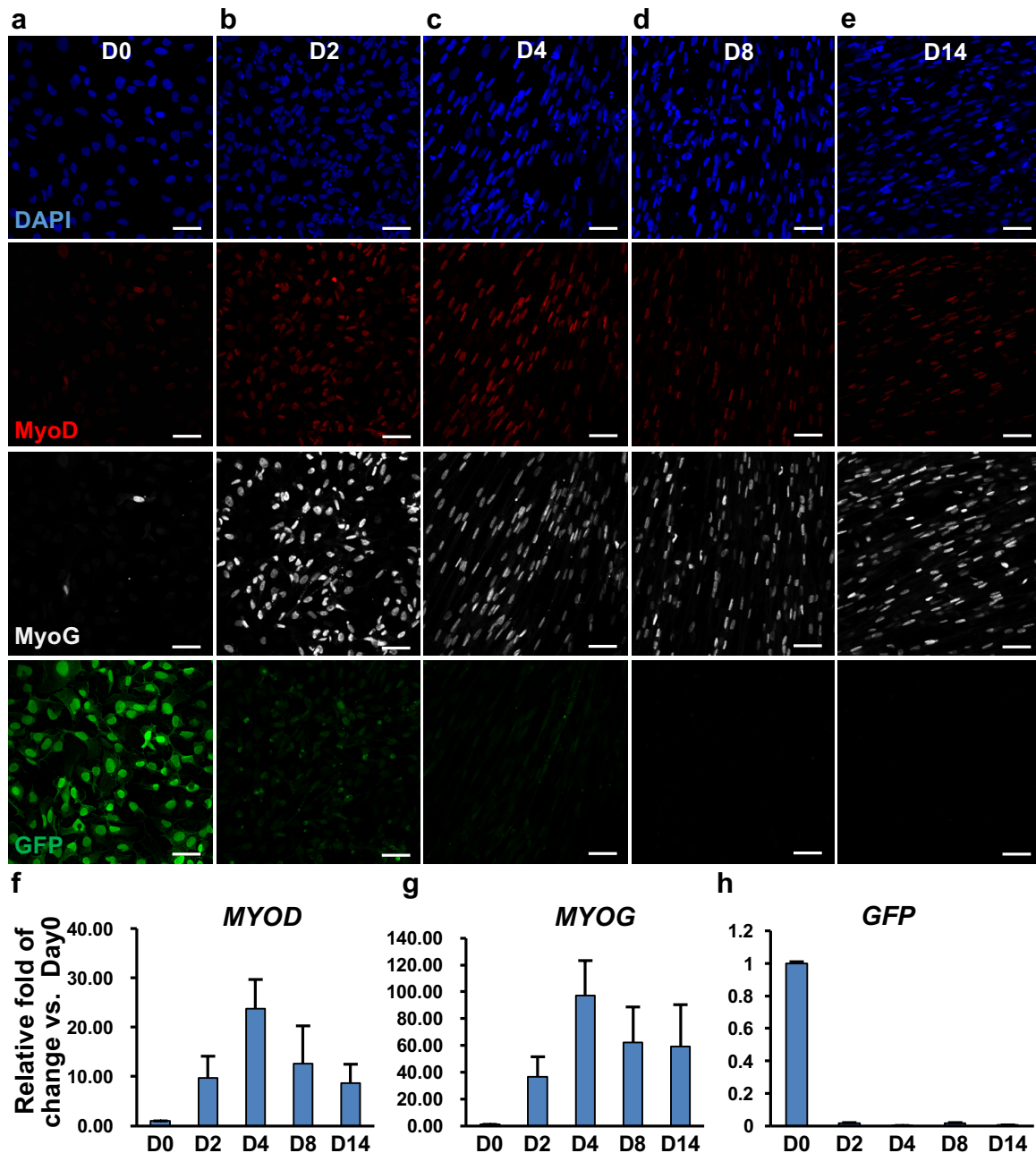
Supplementary Figure 2. Optimizing myogenic progenitor cell derivation. (a) Tet-on lentiviral plasmids used for transient Pax7 induction. (b) mRNA expression level of paraxial mesoderm markers, *MSGN1* and *TBX6*, and early mesoderm marker, *T*, under various treatments for mesoderm induction. EB D5/8/11: 5/8/11 days in embryoid body (EB) suspension culture; C D2: 2 days in CHIR99021 (10 μ M); C+L D2/4/6: 2/4/6 days in CHIR99021 (3 μ M) + LDN193189 (0.5 μ M). Expression shown as fold change relative to undifferentiated hPSCs (n=4 samples for hPSC ctrl, n=6 samples from 3 differentiations for other data points). (c-d) Comparison of MIM and E6 induction media before (c) and after (d) sorting for GFP. (e) Representative FACS analysis of GFP⁺ cells from H9 at sorting. (f) Representative immunostaining of GFP and Pax7 in H9 derived progenitors at day 4 post sorting. Scale bar, 50 μ m. Data are presented as mean \pm SEM.



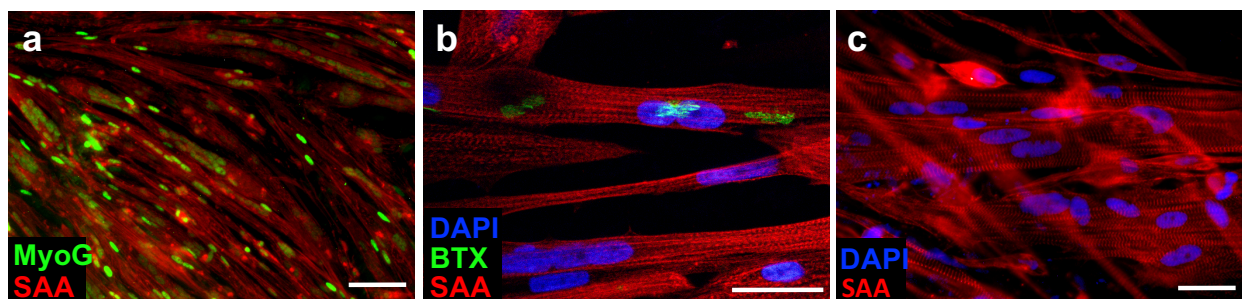
Supplementary Figure 3. Optimizing myogenic progenitor cell expansion. (a) Representative live images of post-sorted GFP⁺ cells in PMM and E6 expansion media. Arrows indicate cells with spindle-like shape. Scale bar, 100 μ m. (b-c) Quantification of GFP⁺ cell fractions (b) and total cell numbers (c) during passaging in different expansion media (*P<0.05 vs. E6 passage 0, #P<0.05 vs. PMM passage 3, Tukey-Kramer HSD test; n = 6 samples from 2 expansions for each condition, data are presented as mean \pm SEM). (d) Representative images of expanded cells differentiated in monolayers for 1, 4 and 14 days showing higher myotube density in PMM-expanded cells. Scale bar, 100 μ m.



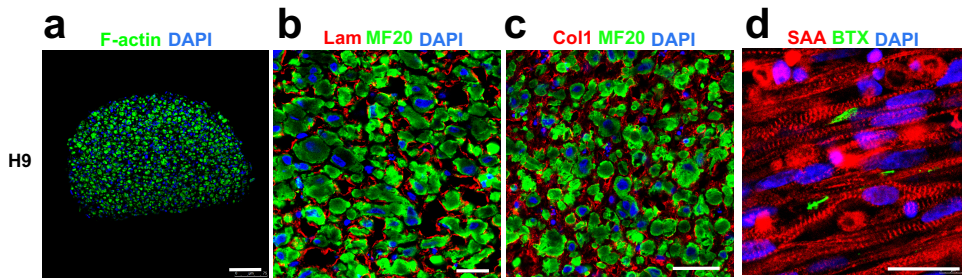
Supplementary Figure 4. Relative mRNA expression levels of *PAX7*, *PAX3*, *MYF5*, *MYOD*, and *MYOG* in undifferentiated hPSCs, iMPCs, and early passaged primary human myoblasts. (* $P < 0.05$ vs. hPSC, # $P < 0.05$ vs. iMPC, Tukey-Kramer HSD test; for hPSCs, $n = 4$ samples from 2 hPSC lines, for iMPCs, $n = 6$ samples from 3 differentiations, for primary human myoblasts, $n = 4$ samples from 2 donors, data are presented as mean \pm SEM).



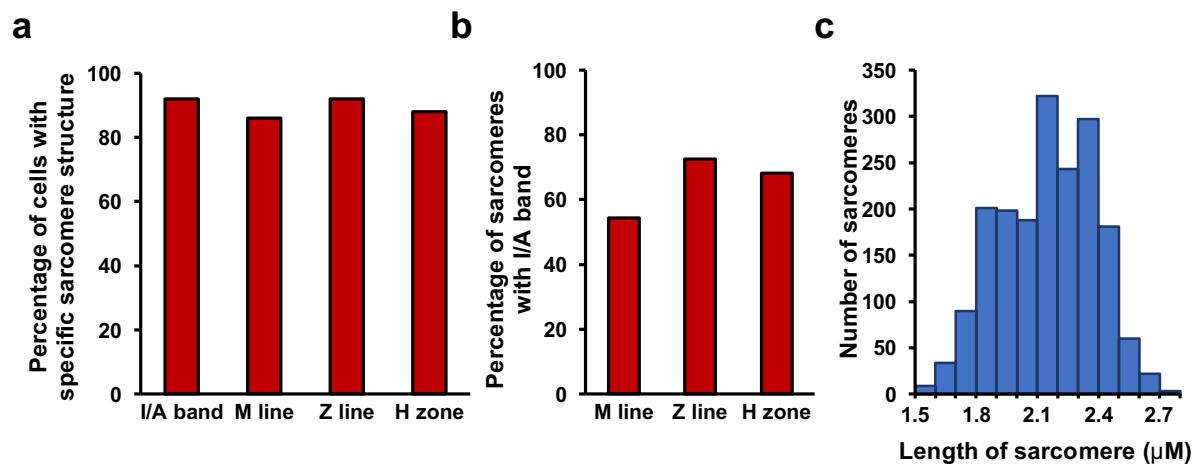
Supplementary Figure 5. Dynamic expression of myogenic markers during 2-week monolayer differentiation of iPSCs. (a-e) Representative immunostaining of MyoD, MyoG, and GFP fluorescence at day (D) 0, 2, 4, 8, and 14 of differentiation. Scale bar, 25 μ m; (f-h) Relative mRNA expression levels of *MYOD*, *MYOG* and *GFP* during 2-week differentiation (n=4 samples from 2 differentiations, data are presented as mean \pm SEM).



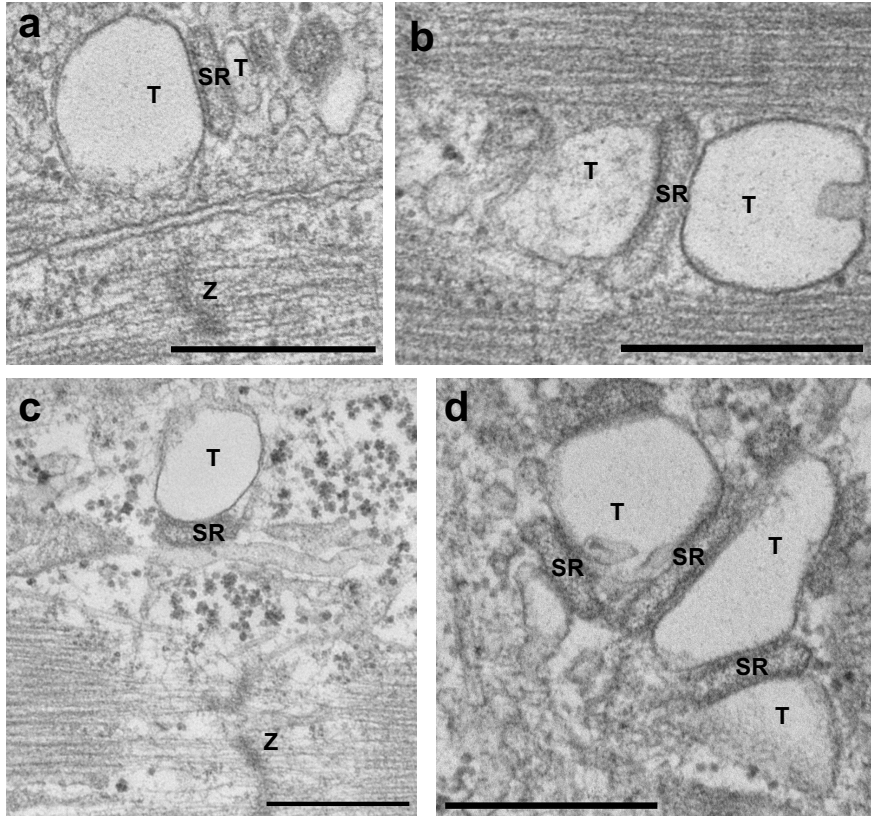
Supplementary Figure 6. Monolayer differentiation of iMPCs. Representative immunostaining of SAA, MyoG, and α -bungarotoxin (BTX) after 2 weeks of monolayer differentiation of iMPCs showing high efficiency of fusion and myogenesis (a, scale bar, 100 μ m), formation of acetylcholine receptors (b, scale bar, 25 μ m), and cross-striations (c, scale bar, 25 μ m).



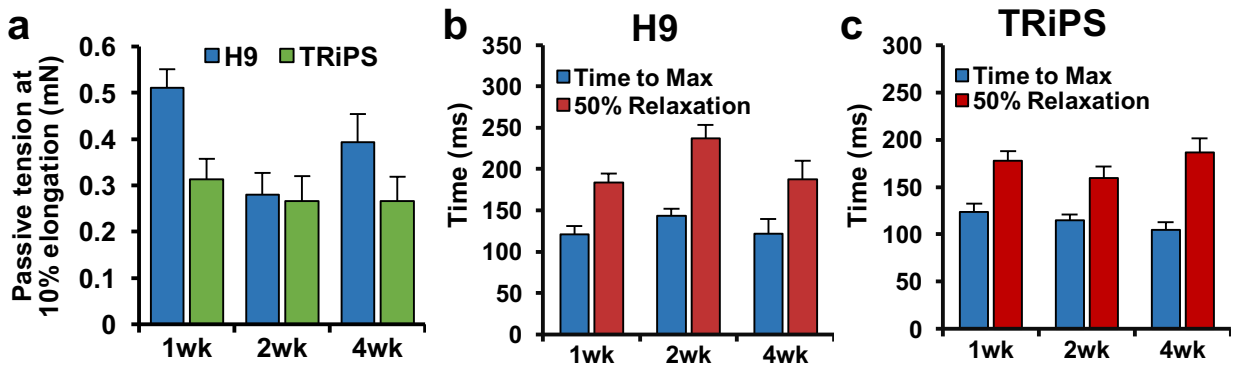
Supplementary Figure 7. Structure of iSKM bundles made from H9 line. (a-c) Representative immunostaining of dense, uniformly distributed, myotubes in bundle cross-section. Scale bars: **a**, 75 μ m; **b**, **c**, 25 μ m. Col 1, collagen I. (d) Representative longitudinal section of 2-week bundles showing aligned, cross-striated myotubes with BTX labeled acetylcholine receptors. Scale bars, 25 μ m.



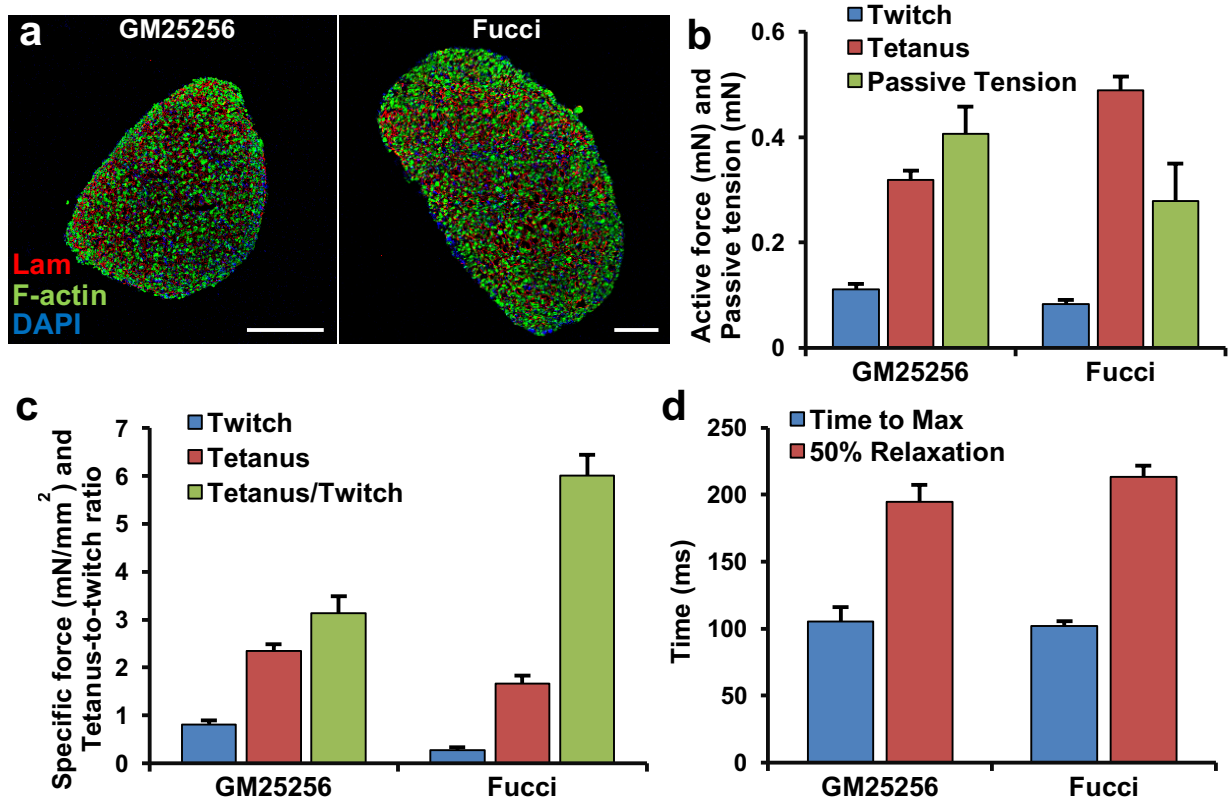
Supplementary Figure 8. Quantification of sarcomeric structures in 4-week differentiated iSKM bundles. (a) Percentage of cells with I/A-band, M-line, Z-line and H-zone (n=50 cells). (b) Percentage of sarcomeres containing I/A bands that also contain M-line, Z-line, and H-zone (n=1367 sarcomeres). (c) Distribution of sarcomere lengths (n=1848 sarcomeres).



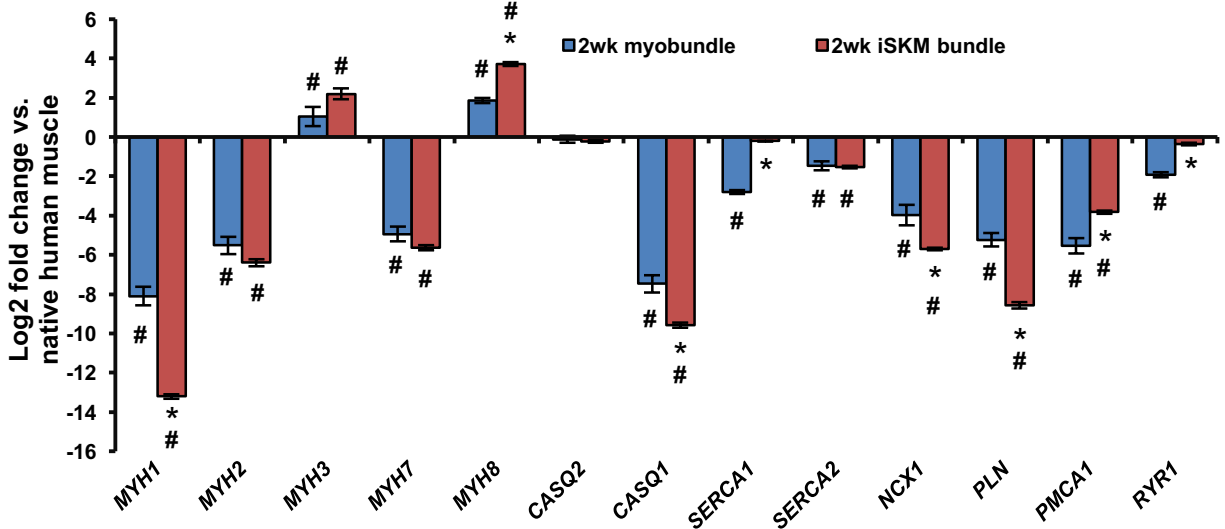
Supplementary Figure 9. Examples of T-SR structures observed in myotubes from 4-week differentiated iSKM bundles. (a,b) Inverted triads. (c) Diads. (d) multi-junction structure. Z: Z-line; T: T-tubule; SR: sarcoplasmic reticulum. Scale bars, 500nm.



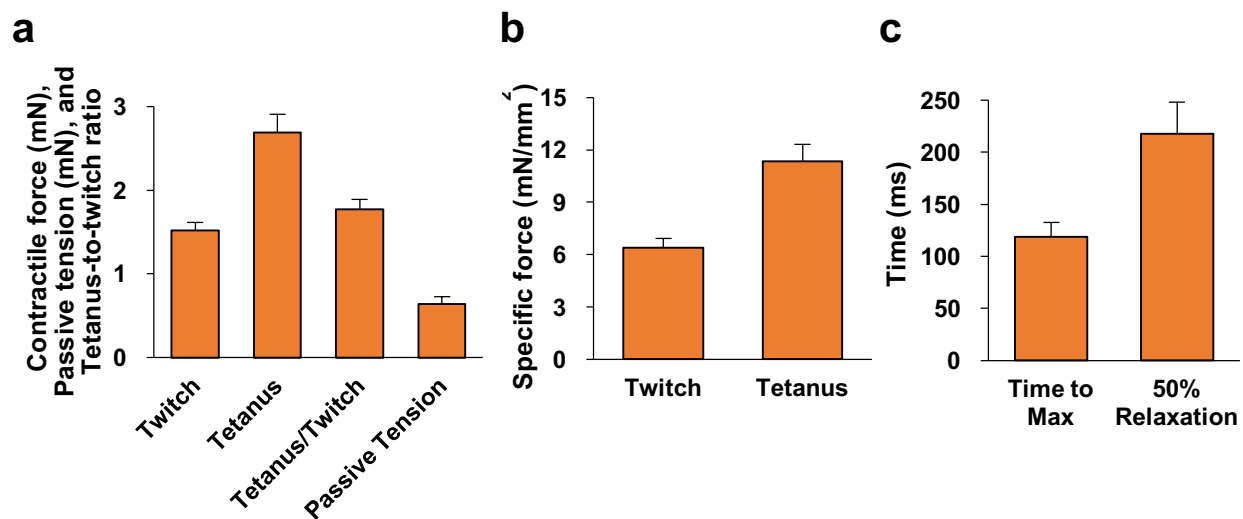
Supplementary Figure 10. Passive mechanical properties and twitch kinetics of iSKM bundles. (a) Passive tension of H9 and TRiPS bundles during 4-week culture. (b, c) Twitch kinetics of H9 (b) and TRiPS (c) derived iSKM bundles during 4-week culture (n=6-8 bundles per group at each time point, data are presented as mean \pm SEM).



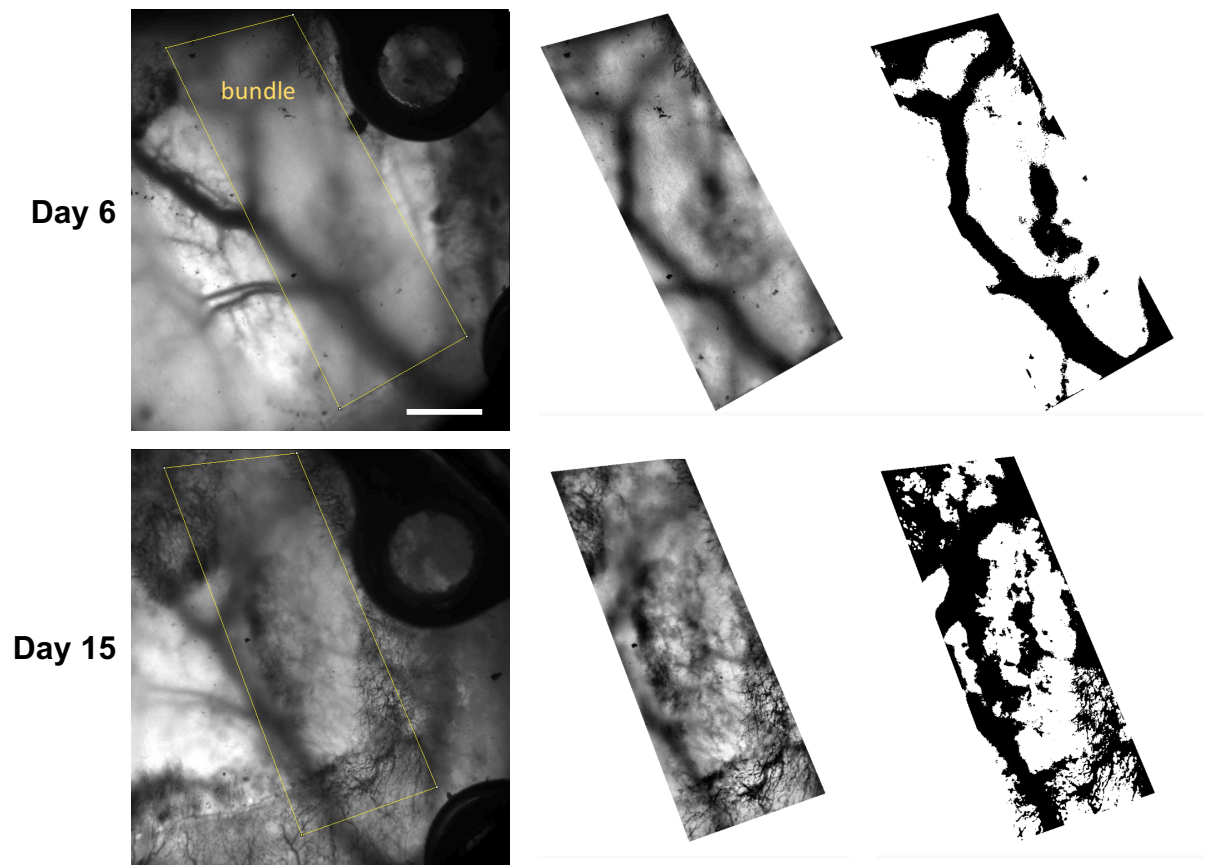
Supplementary Figure 11. iSKM bundles derived from GM25256 and Fucci hPSCs. (a) Representative cross-sections of 2-week differentiated iSKM bundles showing dense F-actin⁺ myofibers surrounded by Laminin (Lam). Scale bar, 100 μ m. (b) Amplitudes of twitch and tetanus (at 40 Hz) and passive tension at 10% elongation generated by GM25256 and Fucci derived iSKM bundles at 2 weeks of differentiation (n=8 bundles for each line). (c) Specific forces and tetanus-to-twitch ratios of 2-week differentiated GM25256 and Fucci derived bundles (n=8 bundles for each line). (d) Twitch kinetics of 2-week differentiated GM25256 and Fucci derived bundles (n=8 bundles for each line). Data are presented as mean \pm SEM.



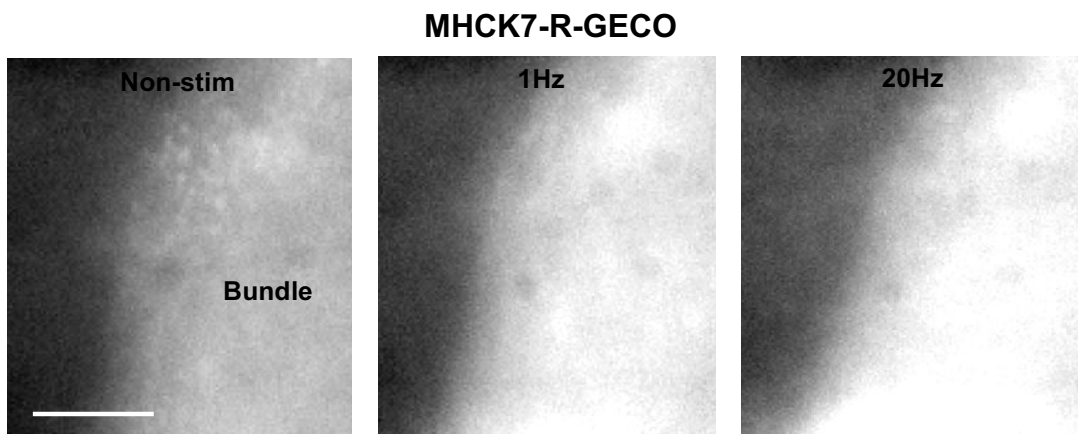
Supplementary Figure 12. Relative mRNA expression levels of different myosin isoforms and calcium handling related genes in 2-week differentiated iSKM bundles, primary human myobundles, and native human muscle. (* $P < 0.05$ vs. 2wk myobundle group, # $P < 0.05$ vs. native human muscle, Tukey-Kramer HSD test; for iSKM bundles, $n = 8$ bundles from 3 differentiations, for primary human myobundles, $n = 6$ bundles from 3 differentiations, for native human muscle, $n = 3$ samples from 3 donors with 4 technical replicates each, data are presented as mean \pm SEM).



Supplementary Figure 13. Contractile properties of primary human myobundles differentiated for 2 weeks in serum-free iSKM differentiation media. (a) Contractile force, passive tension at 10% elongation and tetanus-to-twitch ratio of 2-week differentiated primary human myobundles. **(b)** Specific force of 2-week primary human myobundles. **(c)** Twitch kinetics of 2-week primary human myobundles (n=6 bundles from 2 differentiations). Data are presented as mean \pm SEM.



Supplementary Figure 14. Processing and quantification of blood vessel density in iSKM bundles implanted in dorsal skin-fold window chambers in immunocompromised mice. Representative raw (left), cropped/contrast enhanced (middle), and thresholded (right) intravital images used for quantification of blood vessel density in bundle area 6 and 15 days post-implantation. Note increased transparency of the bundle area with time PI. Scale bar, 1mm.



Supplementary Figure 15. Representative images of *ex vivo* recorded R-GECO signals in iSKM bundles implanted for 1 week in mouse TA muscle, shown without stimulation (Non-stim) or during 1Hz and 20Hz stimulation. Scale bar, 50 μ m.

Supplementary Table 1. Cell culture media and solutions

| Media and solutions | Formulation |
|----------------------------|--|
| Neutralizing media | Low glucose DMEM (Thermo) supplemented with DNaseI (20µg/mL, Sigma) and 50% FBS (Hyclone) |
| Flow buffer | PBS supplemented with 1% FBS (Hyclone) and 0.1% Sodium Azide (Sigma) |
| Blocking solution | PBS supplemented with 5% chicken serum (Sigma), 0.2% Triton X 100 (Sigma) |
| Sorting solution | E6 media supplemented with Y27632 (10µM, Tocris), Dox (1µg/mL, Sigma), DNaseI (20µg/mL, Sigma), penicillin G (100 unit/mL, Thermo) and streptomycin (50 µg/mL, Thermo) |
| Collecting solution | FBS (Hyclone) supplemented with Y27632 (10µM, Tocris), Dox (1µg/mL, Sigma), penicillin G (100 unit/mL, Thermo) and streptomycin (50 µg/mL, Thermo) |
| Expansion media (EM) | Low glucose DMEM (Thermo) supplemented with 10% FBS (Hyclone), Fetuin (500x), hEGF (1000x) and Dexamethasone (1000x) from SkGM SingleQuots Kit (Lonza), penicillin G (100 unit/mL, Thermo) and streptomycin (50 µg/mL, Thermo) |
| Differentiation media (DM) | Low glucose DMEM (Thermo) supplemented with N2 supplement (100x, Thermo), penicillin G (100 unit/mL, Thermo) |
| Cell/hydrogel mixture | 12.5 million cells/mL in expansion media, bovine fibrinogen (4 mg/mL, Sigma), growth factor reduced matrigel (20% v/v, Corning), thrombin (0.2 unit/mg, Sigma) |

Supplementary Table 2. Primers used for qRT-PCR.

| | |
|------------------------|---------------------------|
| <i>G3PD F</i> | AGGTCGGAGTCAACGGATTTGG |
| <i>G3PD R</i> | AGGCTGTTGTCATACTTCTCATGG |
| <i>T F</i> | CCCGTCTCCTTCAGCAAAGTC |
| <i>T R</i> | TAAGAGCTGTGATCTCCTCGTTCTG |
| <i>Tbx6 F</i> | CTCCGTGACAGCCTACCAGA |
| <i>Tbx6 R</i> | GGTGTGTCTCCGCTCCATA |
| <i>Msgn1 F</i> | AGGATGTCTGTCCAGCGGAG |
| <i>Msgn1 R</i> | AGGAGGTCTGTGAGTTCCCC |
| <i>Myf5 F</i> | CGCCTGAAGAAGGTCAACCA |
| <i>Myf5 R</i> | ACATTCCGGGCATGCCATCAG |
| <i>Pax3 F</i> | CCTCAGGTAATGGGACTCCTG |
| <i>Pax3 R</i> | CCCCCTAAAAAGTCCAAGGCT |
| <i>Pax7 F</i> | GGAAGCGATTTTTGCCGACT |
| <i>Pax7 R</i> | TTGTGGCGGATGTGGTTAGG |
| <i>MyoD F</i> | CGACGGCATGATGGACTACA |
| <i>MyoD R</i> | TATATCGGGTTGGGGTTCGC |
| <i>MyoG F</i> | GCCAACCCAGGGGATCAT |
| <i>MyoG R</i> | CCCGGCTTGAAGACAATCT |
| <i>MYH1 F</i> | ATCTAACTGCTGAAAGGTGACC |
| <i>MYH1 R</i> | TAAGTAAATGGAGTGACAAAG |
| <i>MYH2 F</i> | GCCGAGTCCCAGGTCAACAAG |
| <i>MYH2 R</i> | TGAGCAGATCAAGATGTGGCAAAG |
| <i>MYH3 F</i> | GGAGCAGGACAGAAGATAT |
| <i>MYH3 R</i> | CCCAGATTGAAACAAAGCA |
| <i>MYH7 F</i> | CTGTCCAAGTTCGCAAGGT |
| <i>MYH7 R</i> | TCATTCAAGCCCTTCGTGCC |
| <i>MYH8 F</i> | ATTTCCACCAAGAACCCA |
| <i>MYH8 R</i> | AAAGGATTCTGCCTCTGG |
| <i>CASQ1 F</i> | ACATTGTGGCCTTCGCAGAG |
| <i>CASQ1 R</i> | CCATACGCTATCCGCATCAGT |
| <i>CASQ2 F</i> | TTGCCATCCCCAACAAACCT |
| <i>CASQ2 R</i> | AGAGTGGGTCTTTGGTGTTC |
| <i>RYR1 F</i> | GCAAAGGAGAGACAGGTGCC |
| <i>RYR1 R</i> | TGGCGATTGATGACAGTGCC |
| <i>SERCA1 F</i> | GAAGGGAGCACAATGGAGGC |
| <i>SERCA1 R</i> | CAGGCCAGCACGAAGGAAAT |
| <i>SERCA2 F</i> | CCAACGTCGGGGAAGTTGTC |
| <i>SERCA2 R</i> | CGCCGACGTAACAGCCAATA |
| <i>NCX1 F</i> | AGGCCAACCTGTCTTCAGGA |
| <i>NCX1 R</i> | AACCACAAGGGCCAGGTTTG |
| <i>PMCA1 F</i> | AATGGGAACATTACTCTGGGGC |
| <i>PMCA1 R</i> | AGCATTCACTACATCCATCTGTGT |
| <i>PLN F</i> | GCTGCCAAGGCTACCTAAAAG |
| <i>PLN R</i> | GACGTGCTTGTGAGGCATTT |