

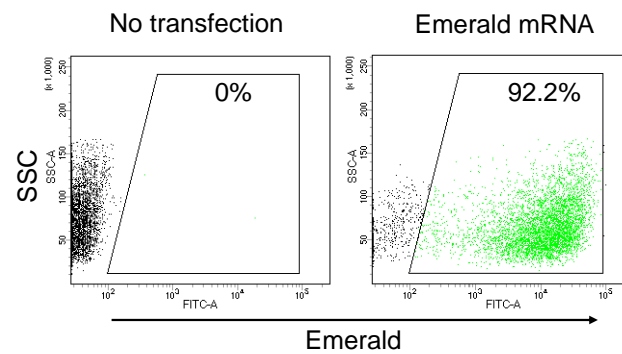
**Efficient differentiation of human pluripotent stem cells into skeletal muscle cells by combining RNA-based MYOD1-expression and POU5F1-silencing**

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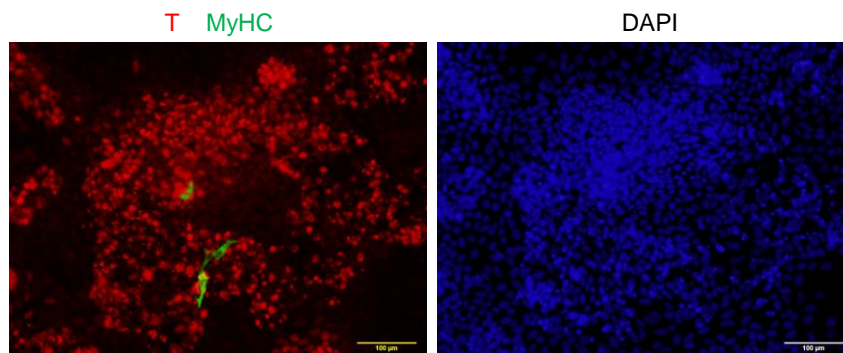
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## Supplementary Figure S1 (Akiyama et al)



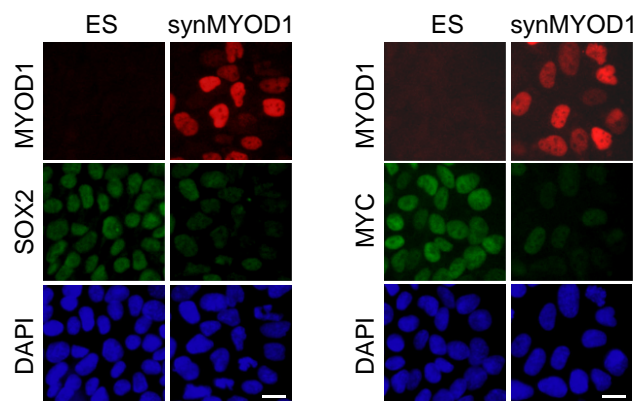
**Supplementary Figure S1. Synthetic mRNA transfection in hiPSCs.** The percentage of mRNA transfection in 409B7-hiPSCs was tested using synthetic mRNA encoding Emerald GFP by FACS analysis.

Supplementary Figure S2 (Akiyama et al)



**Supplementary Figure S2. Expression of T and MyHC in the synMYOD1-transfected cells.** Immunostaining analysis for Brachyury T (a mesoderm marker) and MyHC in the synMYOD1-transfected cells. Nuclei were stained with DAPI.

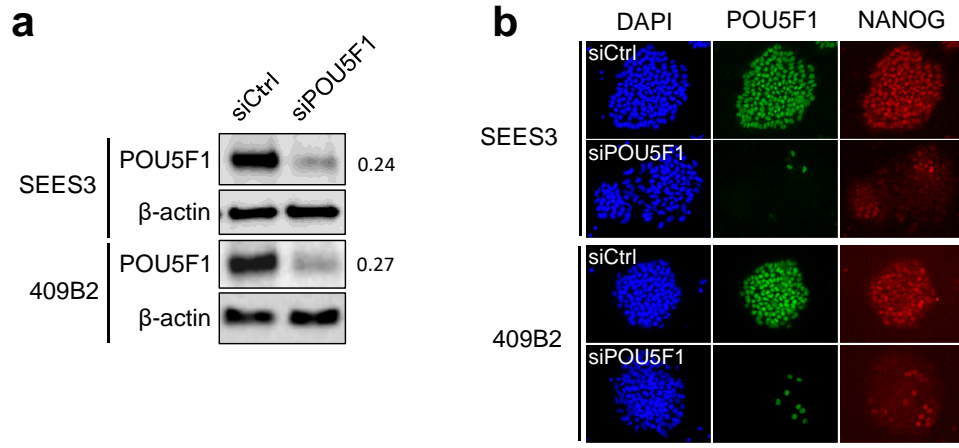
Supplementary Figure S3 (Akiyama et al)



**Supplementary Figure S3. Expression of SOX2 and MYC1 in the synMYOD1-transfected cells.**

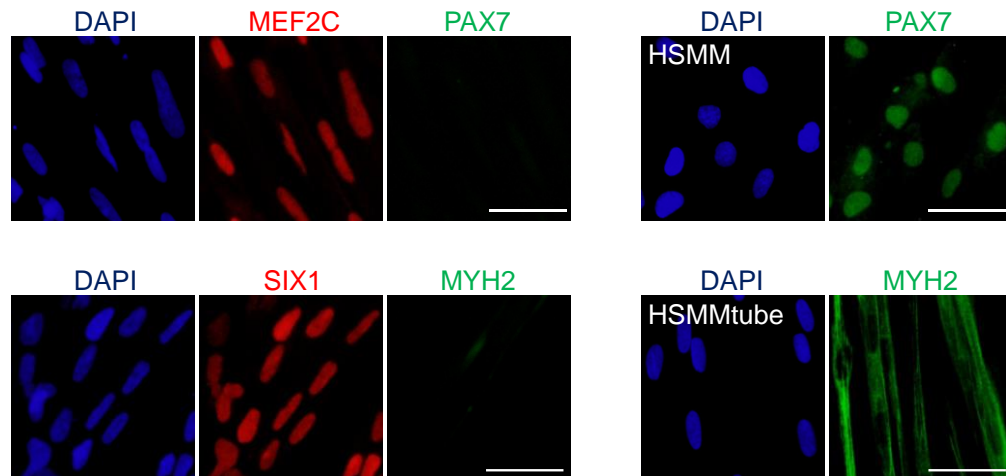
Immunostaining analysis for SOX2 and MYC in the synMYOD1-transfected cells at day 1 post transfection. MYOD1 was detected by the specific antibody. Nuclei were stained with DAPI. Scale bar: 10  $\mu$ m.

## Supplementary Figure S4 (Akiyama et al)



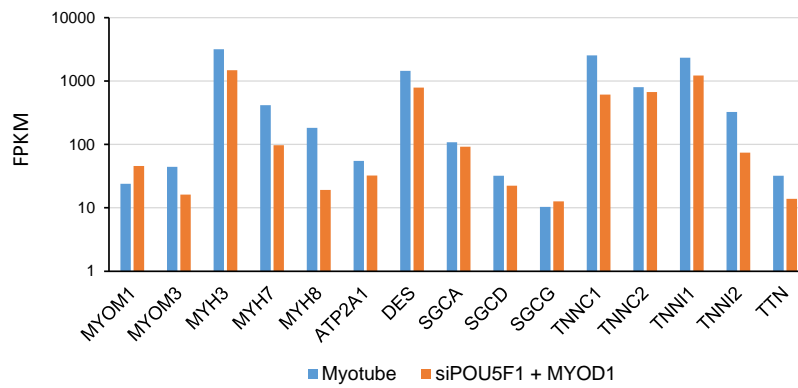
**Supplementary Figure S4. POU5F1 knockdown with siPOU5F1 treatment.** (a) Immunoblotting to detect POU5F1 in siCtrl- and siPOU5F1-treated hESCs (SEES3) and hiPSCs (409B2). The average intensities of siPOU5F1 relative to siCtrl are shown ( $n = 3$ ).  $*p < 0.05$  by t test. (b) Immunostaining to detect POU5F1 and NANOG in siCtrl- and siPOU5F1-treated ESCs (SEES3), and iPSCs (409B2). Scale bar, 100  $\mu\text{m}$ .

Supplementary Figure S5 (Akiyama et al)



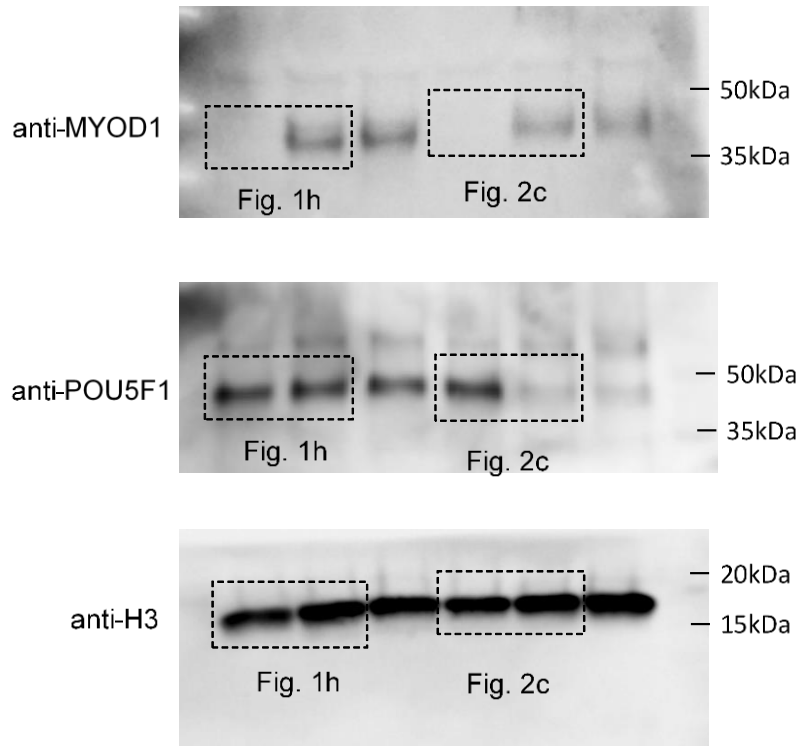
**Supplementary Figure S5. Expression of PAX7 and MYH2 in the siPOU5F1/synMYOD1-transfected cells.** Immunostaining analysis showing no or faint signals of PAX7 (a myoblast precursor marker) and MYH2 (an adult myotube marker) in the siPOU5F1/synMYOD1-treated cells. The HSMM myoblasts and the differentiated myotubes (HSMMtube) were used as positive controls for PAX7 and MYH2 staining, respectively. Nuclei were stained with DAPI. Scale bar: 50  $\mu$ m.

## Supplementary Figure S6 (Akiyama et al)



**Supplementary Figure S6. Comparison of myogenic gene expression between cultured myotubes and siPOU5F1/synMYOD1-induced muscle.** FPKM scores of indicated genes are shown. RNA-seq data of myotubes differentiated from myoblast cells (ref. 9) was used.

Supplementary Figure S7 (Akiyama et al)



**Supplementary Figure S7. Original images of immune blots for indicated figures.**