

# Supplementary Materials

## Fluorescence Spectroscopy of Low-Level Endogenous $\beta$ -adrenergic Receptor Expression at the Plasma Membrane of Differentiating Human iPSC-Derived Cardiomyocytes

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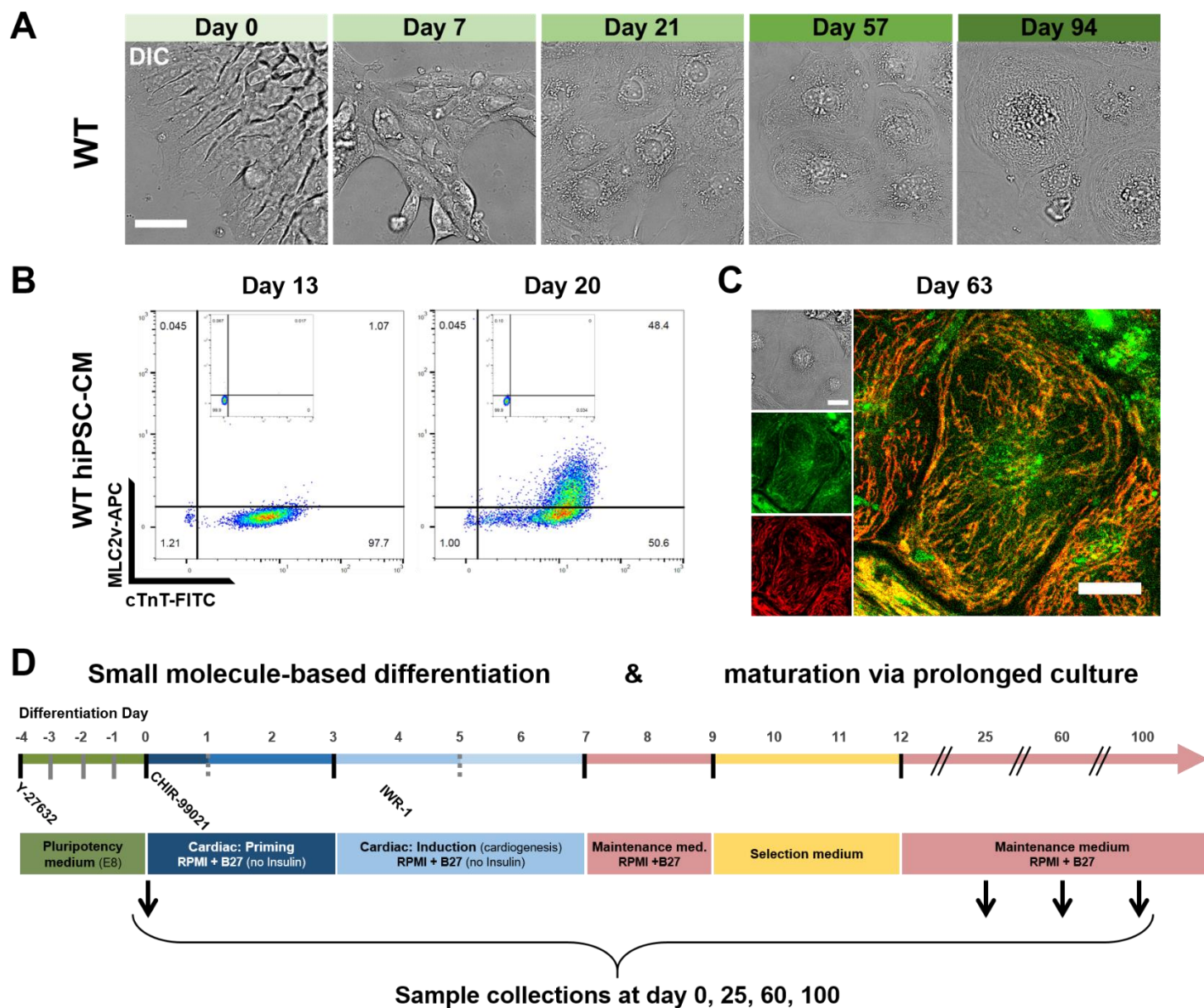
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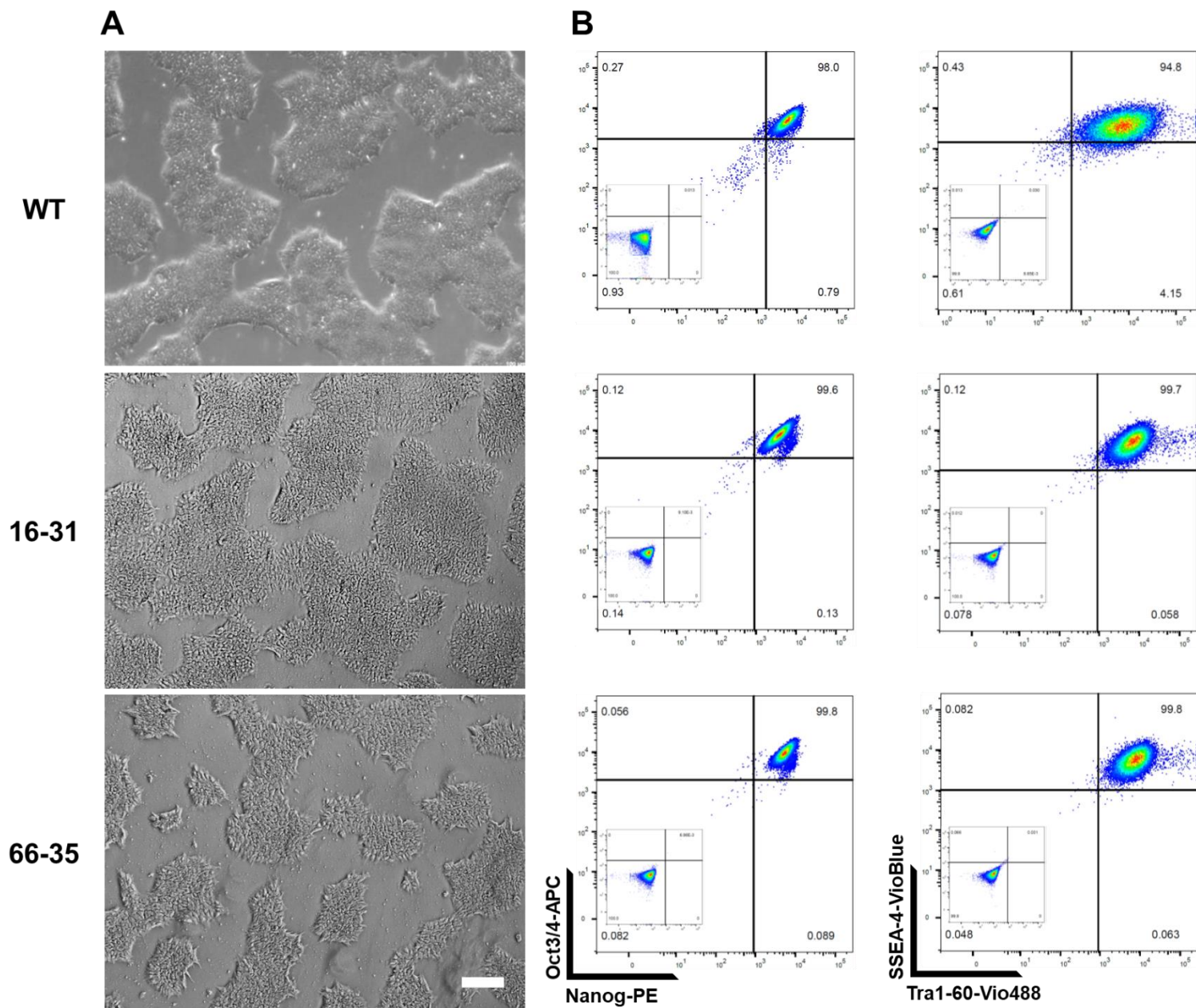
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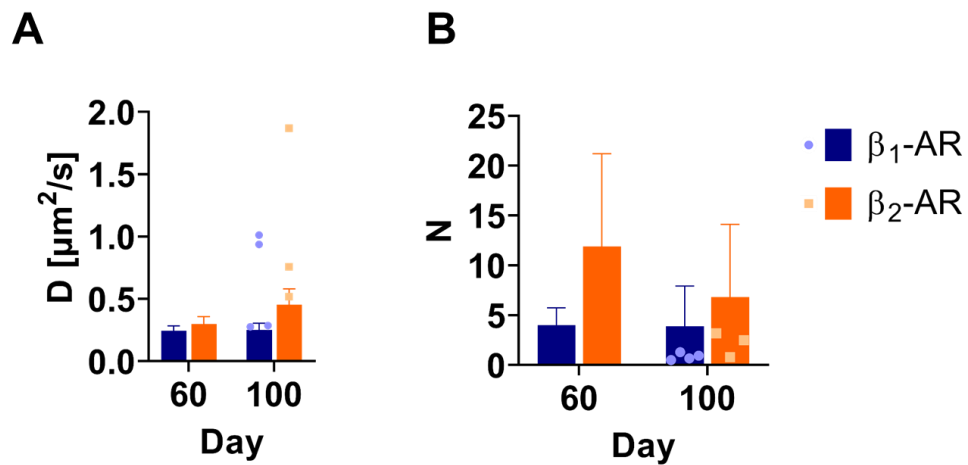


**Figure S1.** Timeline displaying differentiation and maturation of hiPSC-CMs. (A) Shows changes of cellular morphology of undifferentiated WT hiPSC (day 0) and WT cells differentiated and matured in 2D monolayers for 7, 21, 57 and 94 days in culture. Scale bar is 30  $\mu$ m. (B) Representative flow cytometry plots indicating the expression of cardiac-specific markers (cardiac Troponin T (cTnT) and the ventricular isoform of the myosin light chain 2 (MLC2v)) at day 13 and day 20 of differentiation of the parental WT hiPSC-CM line (BIHi005-A), differentiated according to the protocol described in the methods, are shown. Insets display isotype control staining. (C) Mitochondrial networks of day 63 WT hiPSC-CMs superposed to autofluorescence signal. Counterclockwise from top left: DIC (gray), autofluorescence at 488nm (green), 50 nM MitoTracker Deep Red FM stain (red) and overlay of autofluorescence together with MitoTracker. Scale bars are 20  $\mu$ m. (D) Differentiation strategy based on small molecules (see Methods) depicting the most important treatment media indicated by day and duration of treatment and days of sample collection (25  $\pm$  4 days; 60  $\pm$  4 days; 100  $\pm$  7 days).



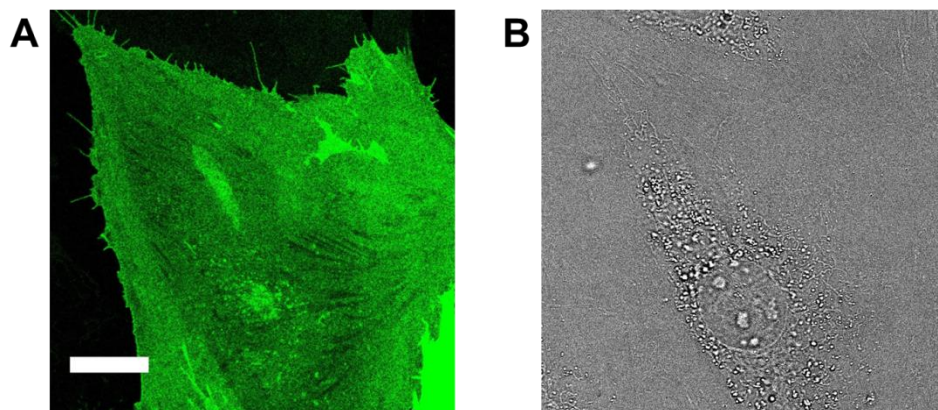


**Figure S3.** Pluripotency characterization of WT hiPSCs and the CRISPR clones. (A) Brightfield images indicate stem cell typical growth in colonies for undifferentiated WT hiPSC and both clones 16-31 and 66-35. Scale is 250  $\mu$ m. (B) Representative flow cytometry plots indicating expression of the 4 pluripotency marker genes Oct3/4 and Nanog (left column) as well as SSEA-4 and Tra1-60 (right column) in all three cell lines used. Insets display isotype control staining.



**Figure S4.** (A) Mean diffusion constants and (B) mean number of particles extracted from the autocorrelation curves in Figure 3. Error bars represent SD values. Dots and small squares indicate diffusion constants and receptor numbers extracted at day 107 (and are not part of the day 100 mean).





**Figure S5.** HA-mEGFP-ADRB2 overexpression in H9c2 cells. (A) Confocal micrograph of the basolateral membrane of an H9c2 cell expressing the construct HA-mEGFP-ADRB2, excited at 488 nm. (B) corresponding DIC image. Scale bar is 20  $\mu\text{m}$ .

**Video S1.** Spontaneous contractions of the hiPSC-CM clone 16-31 (at day 100) before  $\beta_2$ -AR stimulation and after 25 min of 300 nM CGP-20712A addition.

**Video S2.** Spontaneous contractions of the hiPSC-CM clone 16-31 (at day 100) after 5 min of  $\beta_2$ -AR stimulation.