Supplemental material

40%

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Bipolar Polar Astral Flat YFP Actin DNA В 50% Polar ** 45% YFP

Bipolar 35% Percentage of cells 30% Flat Tubulin 25% 20% 15% Astral 10% YFP Tubulin 5% 0% Rounded Fzd7-YFP YFP YFP Fzd7-YFP &Veh. &Wnt7a &Veh. &Wnt7a Fzd7-YFP DNA **Tubulin DNA** Merge Human myoblast Fzd7-YFP DNA Mouse primary myoblast ubulin DNA Fzd7-YFP DNA Pax7 DNA Merge Mouse satellite cell Merge Merge **Tubulin DNA** Fzd3-YFP DNA C2C12

Figure S1. Wnt7a and Fzd7 polarize myogenic cells. (A) Morphological quantification of C2C12 cells transfected with the indicated constructs and treated with either Wnt7a or vehicle. Representative pictures of morphologies are depicted with YFP-transfected cells above. Bars represent means \pm SEM; $n \ge 4$. **, P < 0.01; *, P < 0.05. (B) Subcellular localization of YFP with respect to the tubulin cytoskeleton. Bar, 2 µm. (C) Localization of Fzd7-YFP in primary myoblasts. Note the accumulation of Fzd7 in the cellular periphery. Bar, 10 µm. (D) Localization of Fzd7-YFP in satellite cells that were transfected on single muscle fibers. Accumulation of Fzd7 in the cellular periphery can be observed. Bar, 10 µm. (E) A human primary myoblast that was transfected with Fzd7-YFP. Similar to other myogenic cell types, Fzd7 shows accumulation in the periphery of the cell. Bar, 10 µm. (F) Little to no peripheral localization could be observed for Fzd3-YFP in C2C12 cells. Bar, 10 µm.

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Rounded



Figure S2. **Fzd7 mRNA levels in Fzd7 knockout muscle tissue and dose dependency of Wnt7a-mediated cell migration.** (A) qPCR comparing Fzd7 expression in muscles of Fzd7^{+/-} and Fzd7^{-/-} mice. Bars represent means \pm SEM; n = 3. *, P < 0.05. (B) Scratch assay using mouse primary myoblasts that were exposed to different concentrations of Wnt7a. Bars represent means \pm SEM; n = 3. *, P < 0.05. (B) Scratch assay using mouse primary myoblasts that were exposed to different concentrations of Wnt7a. Bars represent means \pm SEM; n = 3. *, P < 0.05; **, P < 0.01. n.s., no significant difference.



Figure S3. **Gray value quantification of Western blots, Dvl2 knockdown, and Wnt7a endocytosis in C2C12 cells.** (A) Gray value quantification of Western blots for active Rac1 as shown in Fig. 3 B. Bars represent means \pm SEM; n = 3. **, P < 0.01. n.s., no significant difference. (B) qPCR comparing Dvl2 expression upon siDvl2 or siSCR treatment in primary myoblasts. Bars represent means \pm SEM; n = 3. ***, P < 0.001. (C) Gray value quantification from Western blots of Rac1 bound to Dvl2 as shown in Fig. 3 C. Bars represent means \pm SEM; n = 5. *, P < 0.001. (C) Gray value quantification from Western blots of Rac1 bound to Dvl2 as shown in Fig. 3 C. Bars represent means \pm SEM; n = 5. *, P < 0.05. (D) Scratch assay using Wnt7a-stimulated mouse primary myoblasts expressing either EV, RhoA-DN, or Cdc42-DN. Bars represent means \pm SEM; n = 3. **, P < 0.01. (E) A C2C12 cell that was loaded for 3 h with Wnt7a-HA from conditioned medium produced in COS-1 cells, washed extensively, and then cultured for >72 h. Wnt7a was detected by staining for the HA epitope. Bar, 10 µm.



Figure S4. Wnt7a loading does not influence the cell cycle, increases engraftment, and does not alter the number of endogenous satellite cells. (A and B) The proliferation of equal numbers of primary myoblasts in the presence of different concentrations of Wnt7a and Wnt3a was assayed over five days. Data points represent means \pm SEM; n = 3. *, P < 0.05. (C) Immunostaining for engrafted zsGreen-positive satellite cells (full arrowheads) upon Wnt7a and vehicle treatment. Host-derived satellite cells are negative for zsGreen (empty arrowhead). Bar, 50 µm. (D) Pax7⁺/zsGreen⁺ engrafted satellite cells that were treated with Wnt7a do not show more Ki67 staining than vehicle-treated cells. Bars represent means \pm SEM; n = 3. (E) The number of endogenous Pax7⁺/zsGreen⁻ satellite cells is not significantly changed by transplantation of Wnt7a or vehicle-treated satellite cells. Bars represent means \pm SEM; n = 3. n.s., no significant difference.



Figure S5. Wnt7a loading improves the engraftment of mouse primary myoblasts. (A) Strategy used for Wnt7a or vehicle treatment and subsequent transplantation of mouse primary myoblasts into *mdx* mice. (B) Number of dystrophin-positive fibers upon transplantation of Wnt7a or vehicle-treated mouse primary myoblasts. Bars represent means \pm SEM; n = 3. *, P < 0.05. (C) Minimal fiber feret of dystrophin-positive fibers generated from fusion with Wnt7a or vehicle-treated mouse primary myoblasts. Bars represent means \pm SEM; n = 3. *, P < 0.01. (D) Mean maximum cluster distance in muscles transplanted with Wnt7a or vehicle-treated mouse primary myoblasts. Bars represent means \pm SEM; n = 3. *, P < 0.01. (D) Mean maximum cluster distance in muscles transplanted with Wnt7a or vehicle-treated mouse primary myoblasts. Bars represent means \pm SEM; n = 3. *, P < 0.05. (E and F) Three weeks after transplantation of Wnt3a-and Wnt5a-treated zsGreen* myoblasts, no difference in the number of dystrophin-positive fibers or in the mean maximal cluster distance is observed when compared with vehicle. Bars represent means \pm SEM; n = 3. n.s., no significant difference.



Video 1. Localization of Fzd7-YFP in migrating C2C12 cells. C2C12 cells were transfected with Fzd7-YFP. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (LSM710; Carl Zeiss). Frames were taken every 15 s for 29.5 min.



Video 2. Localization of YFP in migrating C2C12 cells. C2C12 cells were transfected with YFP. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (LSM710; Carl Zeiss). Frames were taken every 15 s for14.75 min.



Video 3. Effect of Wnt7a on satellite cell migration. Satellite cell time-lapse microscopy was performed using an inverted microscope (DMI 5100; Leica) and MetaMorph 7.6.1 software (Molecular Devices) with time points being acquired every 7 min in a stage-top incubator (LiveCell Imaging) for 24 h.