Supplementary Information

Monocarboxylate transporter-1 promotes osteoblast differentiation via suppression of p53, a negative regulator of osteoblast differentiation

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Supplementary Figure S1. Silencing of *Mct1* **gene by siRNA in C2C12 cells.** (a) C2C12 cells were cultured for the indicated periods. Expression of mRNAs for *Mct1*, *Mct2*, *Mct3*, and *Mct4* were analyzed by real-time RT-PCR. Amplification signals from these genes were normalized against that of *Gapdh*. Relative values are indicated as the mean \pm SD (n=3). (b) C2C12 cells were introduced with *Mct1* siRNA and cultured for 48 hours in the presence or absence of BMP-2 (300 ng/mL). *Mct1* gene expression in C2C12 cells was quantitatively assessed by real-time RT-PCR, then normalized against that of *Gapdh*, with the results indicated as relative values. (c) Intracellular lactic acid in C2C12 cells introduced with control or *Mct1* siRNA was determined after culturing for 24 hours. (d) C2C12 cells introduced with control or *Mct1* siRNA were cultured for the indicated periods in the presence or absence of BMP-2 (300 ng/mL). Cell viability was assessed based on the reduction of MTS. (b-d) Values are expressed as the mean \pm SD (n=3). **Significantly different from control group (*p* <0.01). n.s., not significant

Supplementary Figure S2. Suppression of mineralization by *Mct1* knockdown in C2C12-cell cultures. C2C12 cells introduced with control or *Mct1* siRNA were cultured for 4 days in the presence or absence of 300 ng/mL BMP-2 in DMEM plus 2.5% FBS, followed by additional 3-day culture in DMEM supplemented with 50 μ g/mL ascorbic acid, 10 mmol/L β -glycerophosphate, and 10 nmol/L dexamethasone. (a) Cells were stained with 1% alizarin red. (b) Alizarin red bound to cultured cells was dissolved in 10% (w/v) cetylpyridinium chloride and quantified by reading the absorbance at 570 nm. Data are expressed as the mean ± SD for 12 experiments. **Significantly different from control group (p < 0.01)

Supplementary Figure S3. Phosphorylation of smad1/5, ERK1/2, and JNK in C2C12 cells after addition of BMP-2. Western blotting for (a) phosphorylated and un-phosphorylated smad1/5, (b) ERK1/2, and (c) JNK of C2C12 cells was performed after incubation with BMP-2 (300 ng/mL) for the indicated periods. Intensity of each band was quantified using Versa Doc 500 MP (Bio-Rad Laboratories). Data are expressed as the mean ± SD for 3 independent experiments. Representative photographs of the membranes are shown in Figure 3.

Supplementary Figure S4. Effects of *Mct1* knockdown on phosphorylation of AKT in C2C12 cells after stimulation with BMP-2. (a) C2C12 cells harboring control or *Mct1* siRNA were exposed to BMP-2 (300 ng/mL). AKT and phosphorylated AKT (p-AKT) in cell lysates obtained at the indicated times after addition of BMP-2 were analyzed by western blotting. (b) Quantitative data are expressed as the mean \pm SD for 3 independent experiments. *Significant difference between groups (p < 0.05)

а

b

MyoD

С

Supplementary Figure S5. Effect of Mct1 siRNA on myogenic differentiation of C2C12 cells. C2C12 cells introduced with control or Mct1 siRNA in combination with Trp53 siRNA or Klf4 siRNA were cultured for 6 days without addition of BMP-2. (a) Representative results of immunocytochemical analysis of the cells for myosin heavy chain are shown. (b, c) Expression of mRNAs for MyoD (b) and Myogenin (c) were analyzed by real-time RT-PCR. Amplification signals from these genes were normalized against that of *Gapdh*. Relative values are indicated as the mean \pm SD (n=3). *Significant difference between groups (p < 0.05)

Mct1

Mct1

Trp53

Mct1 +

Klf4

0.5

0

siRNA: Control

а

Supplementary Figure S6. Effects of Mct1 siRNA on expression and nuclear translocation of HIF-1a in C2C12 cells. C2C12 cells introduced with control or Mct1 siRNA were cultured for 2 days in the absence of BMP-2. (a,b) Representative images of immunofluorescence staining of cells using anti-HIFa (green) and nuclear staining with DAPI (blue) are shown. (c) Quantitative data are expressed as the mean ± SD for 3 independent experiments.

Supplementary Figure S7. Reduction of glucose uptake by Mct1 siRNA.

Control (Control) or *Mct1* siRNA was introduced into C2C12 cells. After incubation for 2 hours in DMEM supplemented with 2.5% FBS, 2-deoxyglucose uptake was analyzed using a Glucose Uptake-GloTM Assay kit (Promega), with luminescence (vertical axis) proportional to the concentration of 2-deoxyglucose. *Significantly different from control group (p < 0.05)

Supplementary Figure S8. Promotion of osteoblast differentiation by glucose in C2C12 cells. ALP activity staining of C2C12 cells cultured in DMEM containing glucose at a concentration of 4500 mg/L (high glucose) or 1000 mg/L (low glucose), and supplemented with 2.5% FBS with (+) or without (-) BMP-2 (300 ng/mL).

Supplementary Figure S9. Original images of immunoblots with molecular weight standards.