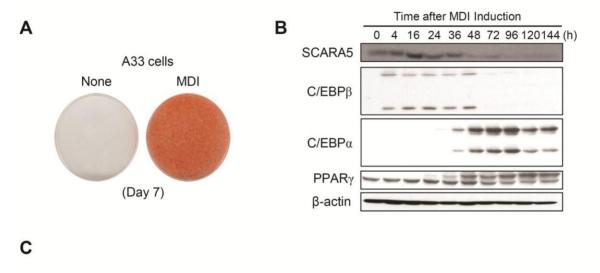
Supplementary Materials

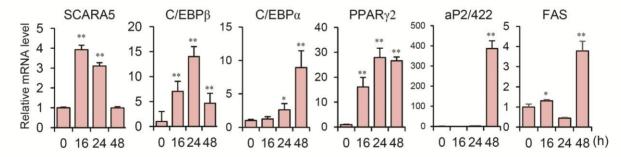
SCARA5 plays a critical role in the commitment of mesenchymal stem cells to adipogenesis

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Contents : Supplementary Figures and Legends (Figures S1-S5), Supplementary Dataset 1 (available in the separate excel file), and raw data for western blots.

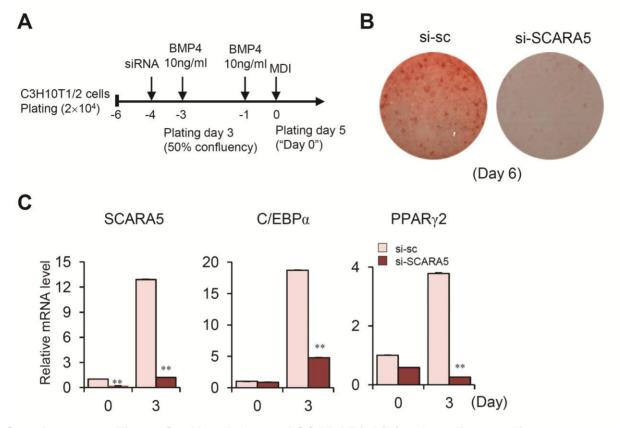
Supplementary Figure S1





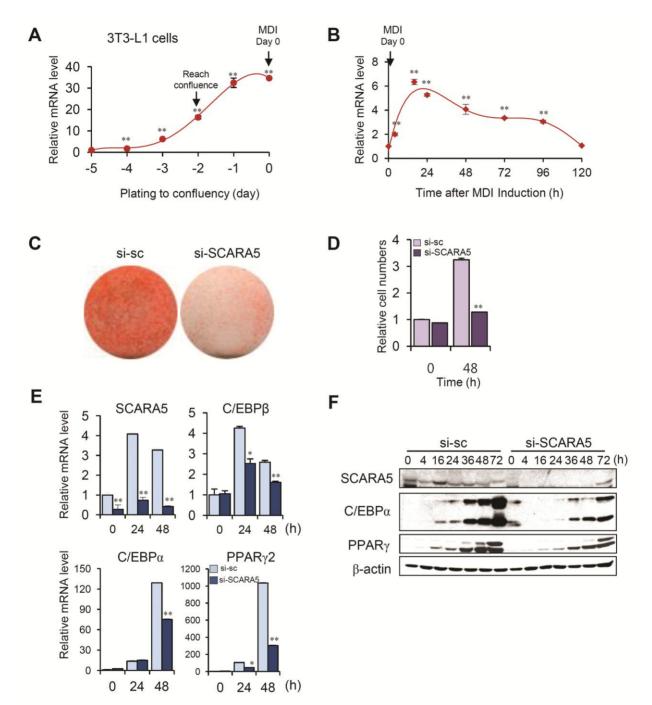
Supplementary Figure S1. The effects of SCARA5 on the differentiation of A33 cells. (A) Post-confluence A33 preadipocyte cells were treated with or without differentiation medium. The cells were then stained with Oil red-O on day 7. (B) SCARA5 expression during adipogenesis was validated by western blotting. (C) SCARA5 expression during adipogenesis was validated using real-time PCR. C/EBP β , C/EBP α , PPAR γ 2, aP2/422, and FAS were used as adipocyte differentiation markers. Quantitative data are presented as the mean ± SD (n = 3). **P* < 0.05; ***P* < 0.01 compared with control.

Supplementary Figure S2



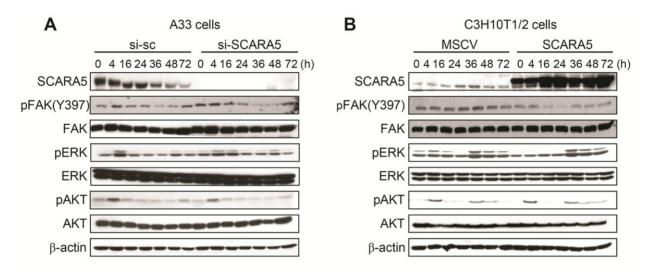
Supplementary Figure S2. Knockdown of SCARA5 inhibits the adipocyte lineage commitment of C3H10T1/2 cells. (**A**) C3H10T1/2 cells were incubated with SCARA5 siRNA or scramble for 24h prior to treatment with BMP4 (10 ng/ml). Cells were grown until postconfluence, and then cell differentiation was induced (MDI). (**B**) The effects of SCARA5 silencing on commitment to the adipocyte lineage and subsequent adipocyte differentiation were assessed on day 6 by Oil red-O staining. (**C**) SCARA5, C/EBPα and PPARγ2 mRNA levels were measured in the scramble (sc) or si-SCARA5 cells using real-time PCR.

Supplementary Figure S3



Supplementary Figure S3. Knockdown of SCARA5 inhibits adipocyte differentiation in **3T3-L1 cells.** (A) 3T3-L1 cells were grown until post-confluence, total RNA was extracted, and real-time PCR was conducted to detect SCARA5 expression. (B) 3T3-L1 cells were differentiated, total RNA was isolated, and real-time PCR was conducted to detect SCARA5

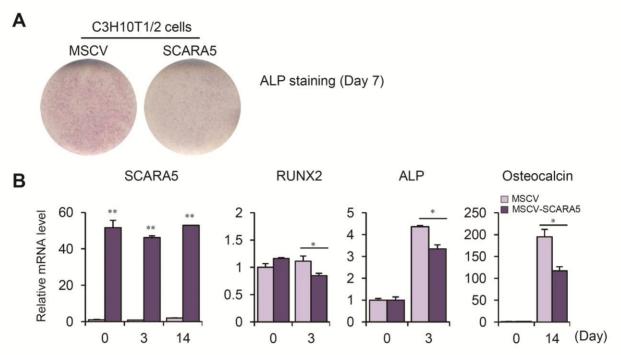
expression at the indicated time points. (**C**) The effects of SCARA5 silencing on commitment to the adipocyte lineage and subsequent adipocyte differentiation were assessed on day 5 by Oil red-O staining. (**D**) Cell counts were determined at 0 and 48 h after induction. (**E**) SCARA5, C/EBP α , C/EBP β and PPAR γ 2 mRNA levels were measured in the scramble (sc) or si-SCARA5 cells using real-time PCR (**F**) Proteins collected at the indicated times after induction were used to assay SCARA5, C/EBP α , and PPAR γ 2 levels with β -actin as the loading control. Quantitative data are presented as the mean ± SD (n = 3). **P* < 0.05; ***P* < 0.01 compared with the control.



Supplementary Figure S4

Supplementary Figure S4. The function of SCARA5 in adipocyte lineage commitment is dependent on FAK-ERK. (A) A33 cells were transfected with si-scramble and si-SCARA5 and differentiated using IBMX, insulin, and dexamethasone (MDI). SCARA5, pFAK (Tyr397), FAK, pERK, ERK, pAKT, and AKT were detected at the indicated time points by western blotting with β -actin as a loading control. (B) C3H10T1/2 cells were infected with MSCV or MSCV-SCARA5. After reaching post-confluence, cells were induced to differentiate using an adipocyte differentiation protocol. SCARA5, pFAK (Tyr397), FAK, pERK, ERK, pAKT, and AKT were detected at the indicated time points by western blotting with β -actin as a loading control.

Supplementary Figure S5



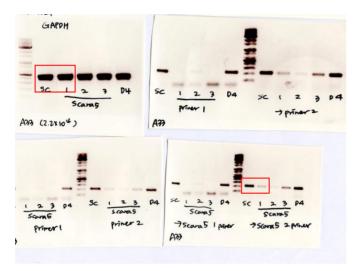
Supplementary Figure S5.The Effects of SCARA5 on osteoblast differentiation in C3H10T1/2 cells. Cells were cultured in an osteogenic medium. (A) On day 7, cells were fixed and stained with Alkaline phosphatase (ALP) staining. (B) Cells were prepared and mRNA of osteoblastic marker genes was analyzed by real-time PCR. Quantitative data are presented as the mean \pm SD (n = 3). **P* < 0.05 compared with the control.

Supplementary Dataset 1

Supplementary Dataset 1. Microarray analysis on proliferating A33 and C3H10T1/2 cells (Excel 1.96MB)

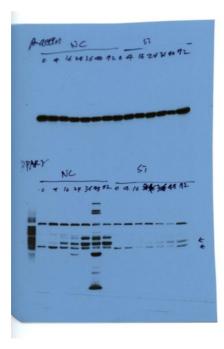
Raw data

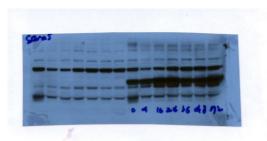
Fig. 2a





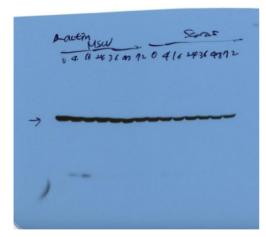






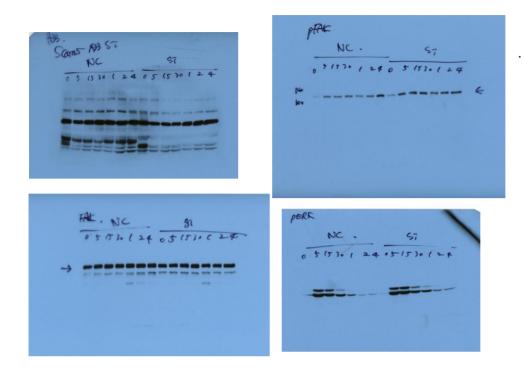


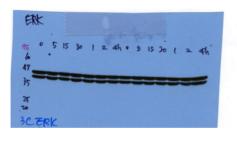
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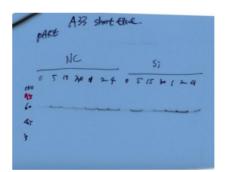












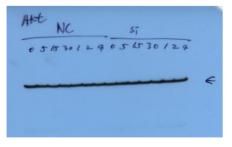
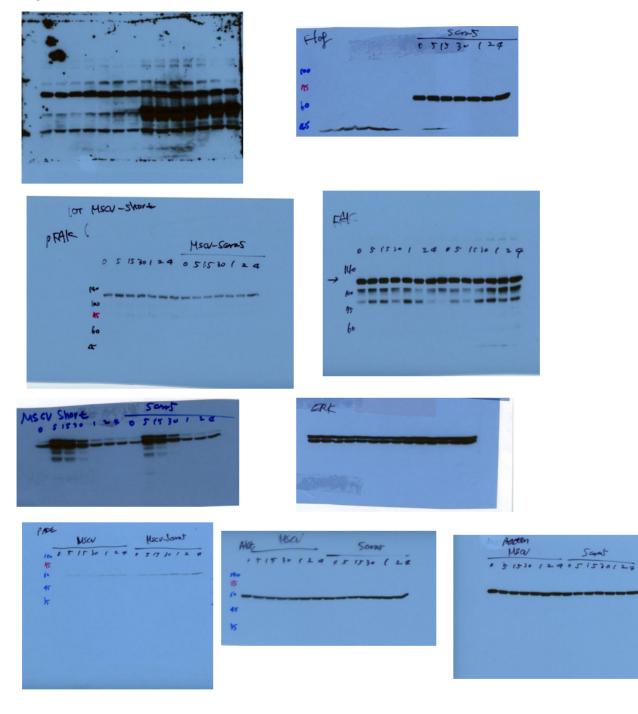
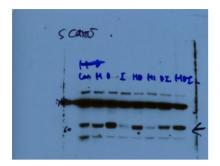




Fig. 3d







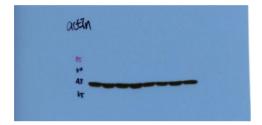


Fig. 4d

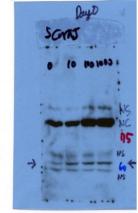




Fig. 4e

