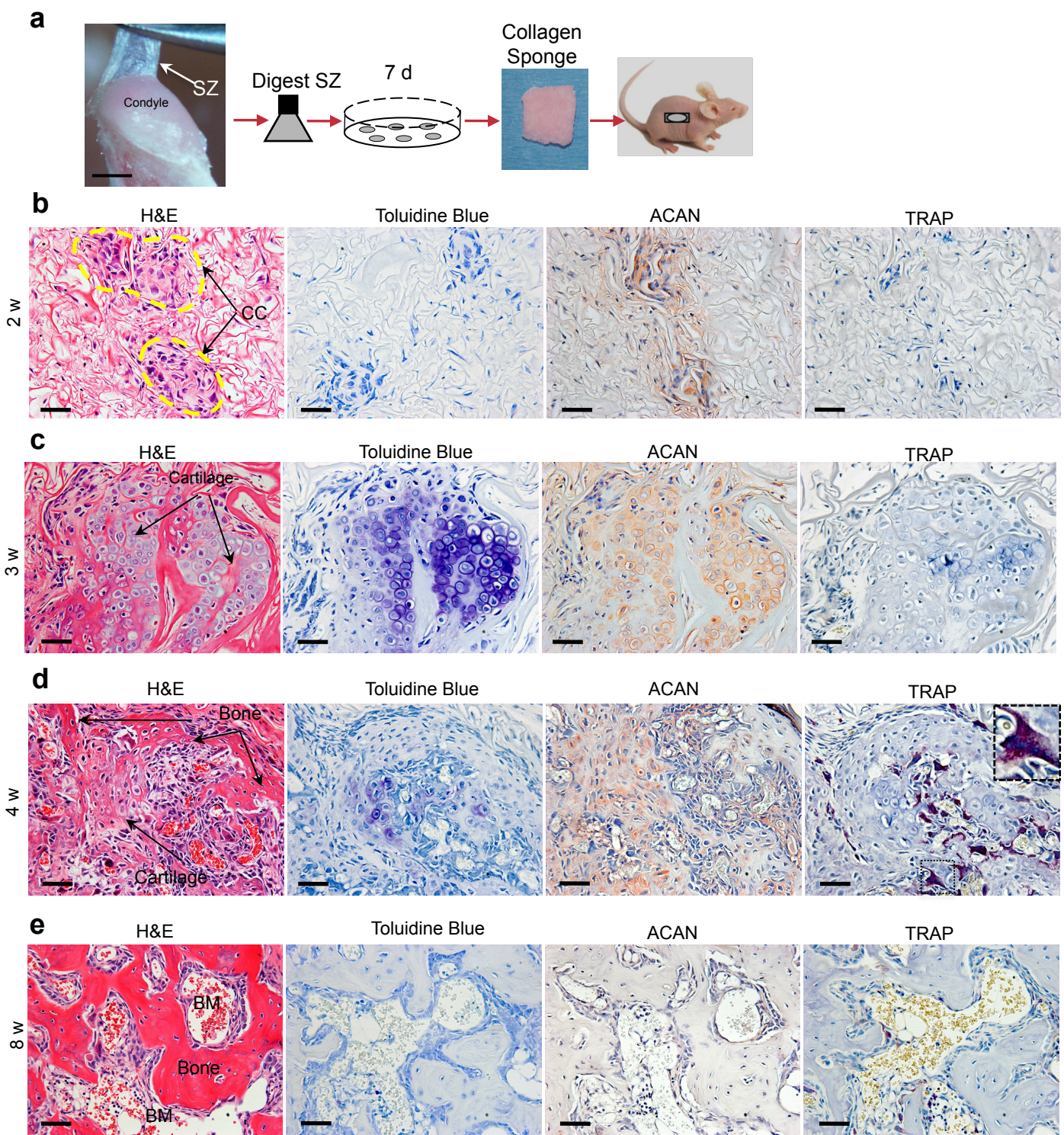


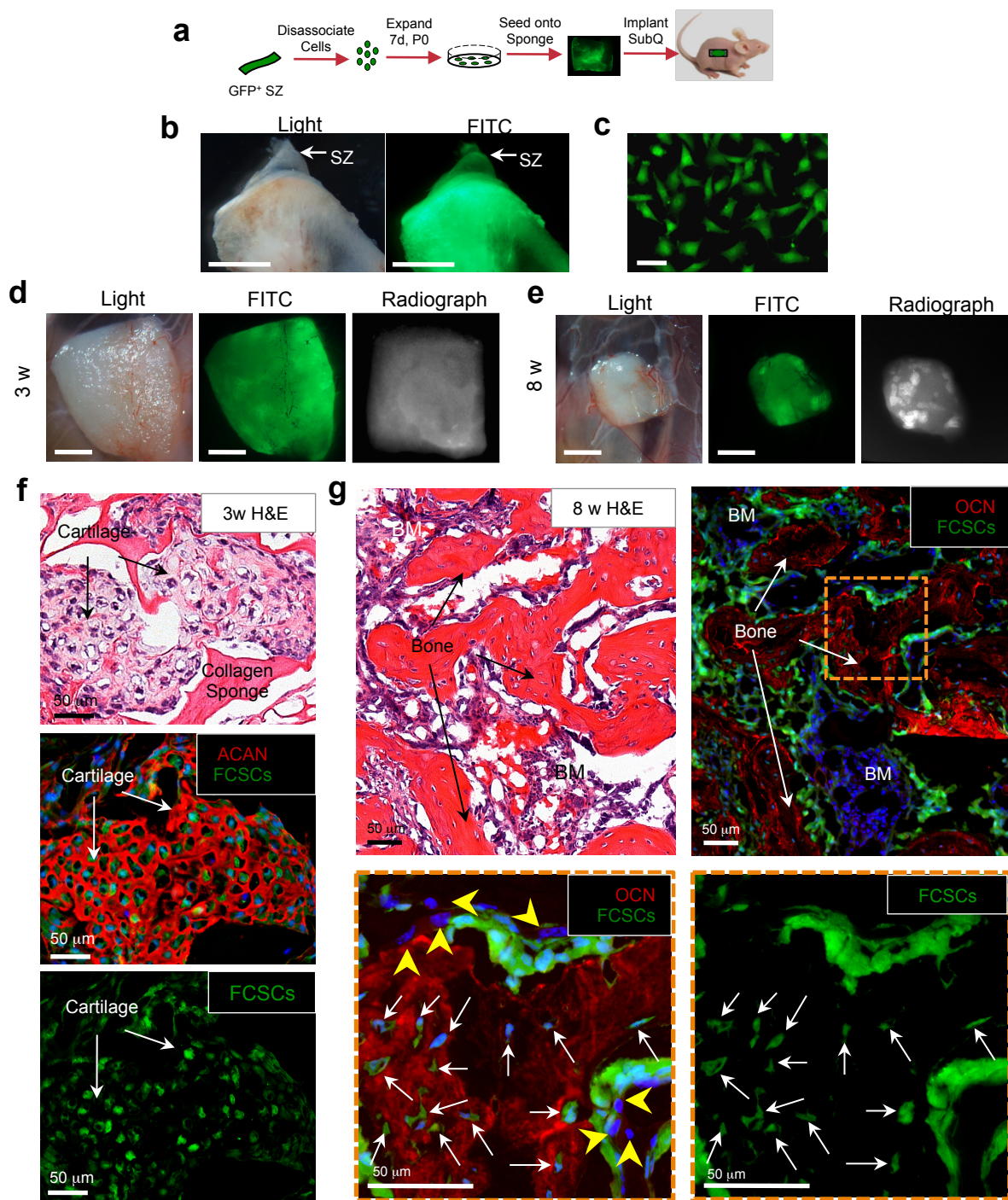
Supplementary Figure 1: FCSCs cell surface markers

(a) Flow cytometric analysis of FCSCs and donor-matched BMSCs. Data are mean \pm S.D.; n=4 experiments. **(b)** Representative gating strategy for FCSCs. To determine positive cells compensation control including single fluorochrome and gating control excluding single fluorochrome were used.



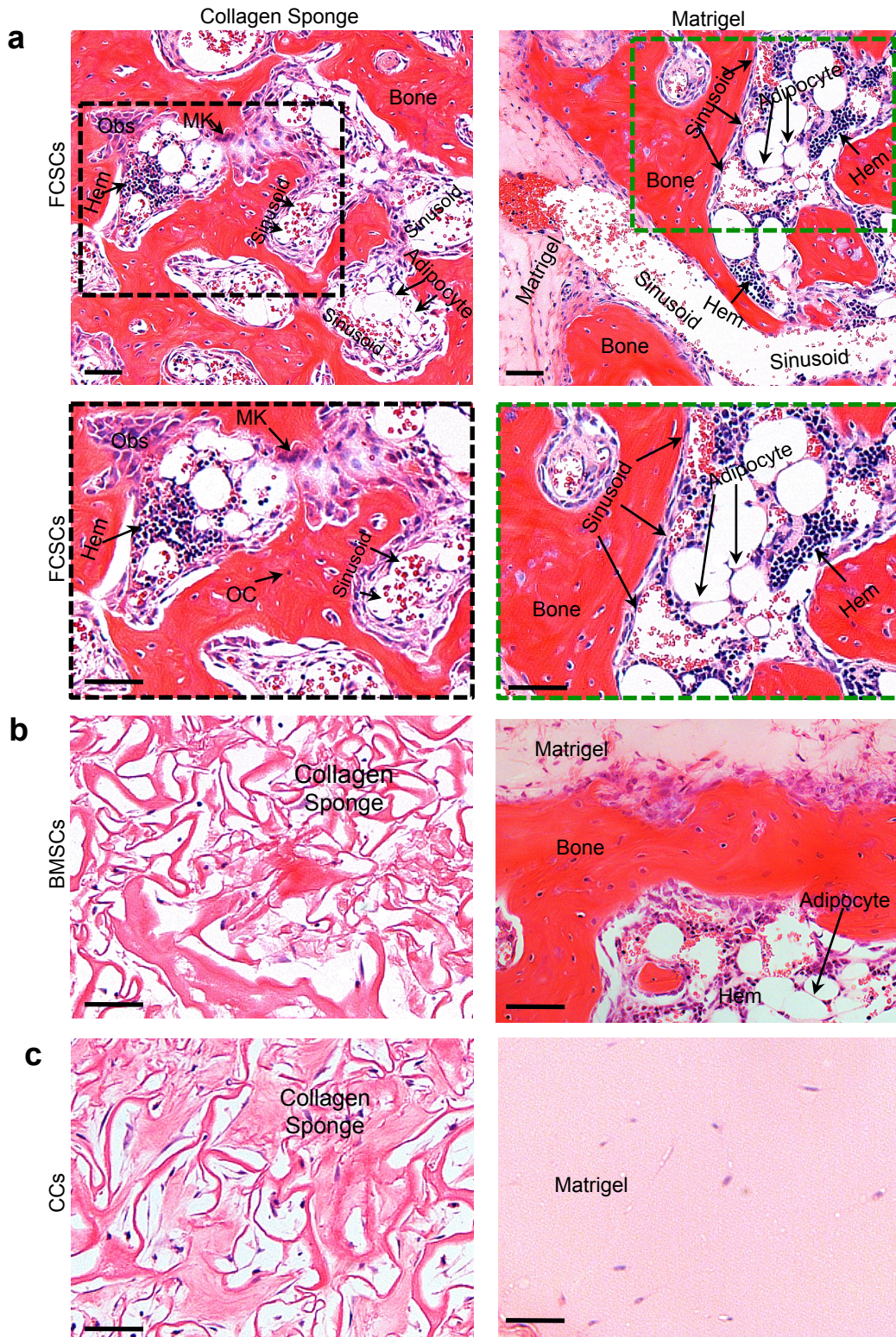
Supplementary Figure 2: FCSCs recapitulate endochondral ossification *in vivo*

(a) Schematic demonstrating FCSC transplantation experiment. Scale bar = 2mm. (b-e) Time course of representative histological sections of regenerated tissue after 2 (4/6 transplants), 3 (5/6 transplants), 4 (5/6 transplants) and 8 (7/8 transplants) weeks *in vivo*. Serial sections showing hemotoxylin and eosin (H&E) stain, toluidine blue stain, aggrecan immunohistochemistry (ACAN) and tartrate-resistant acid phosphatase (TRAP) staining. BM=bone marrow. CC=cell condensation. Sections are representative images. Scale bar= 50 μm.



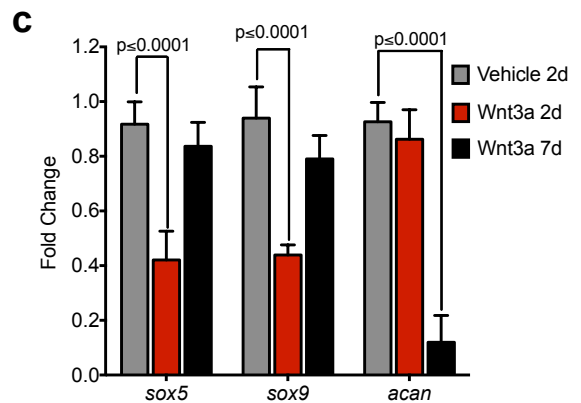
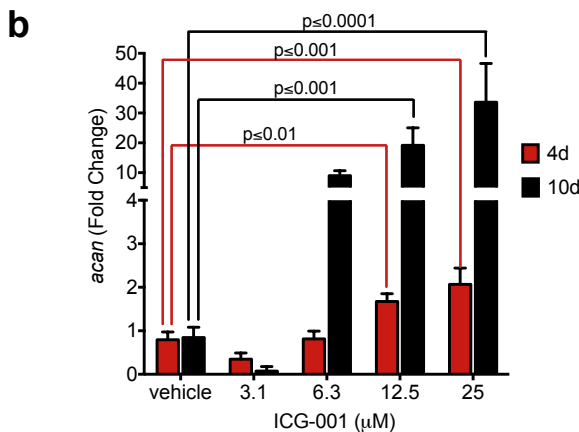
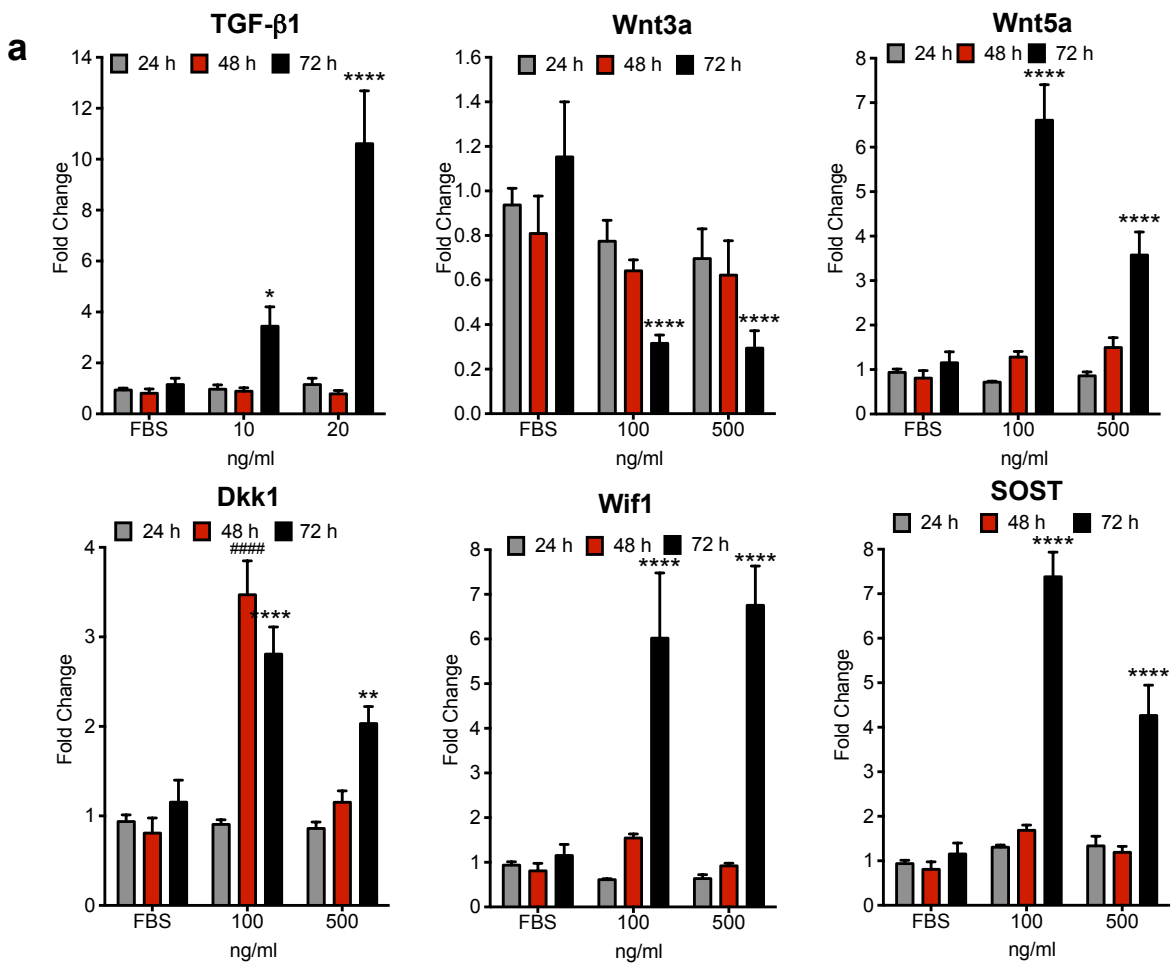
Supplementary Figure 3: GFP⁺ FCSCs spontaneously form cartilage and bone

(a) Schematic of GFP⁺ FCSC transplantation experiment. (b) GFP⁺ rat superficial zone tissue (SZ) dissection shown in light and fluorescent microscopy. Scale bar=4 mm. (c) GFP⁺ FCSCs shown at 5 days. Scale bar=50 μm. (d,e) GFP⁺ FCSC collagen sponge transplants on dorsum of nude mice after 3 and 8 weeks in vivo, respectively. Scale bar=3mm. (f) Representative hematoxylin and eosin staining of GFP FCSC transplant after 3 weeks. Serial section showing GFP⁺ FCSCs under fluorescent light (green) and immunohistochemistry for aggrecan (acan, red). Dapi blue staining=nuclei. Scale bar=50 μm. (g) Representative hematoxylin and eosin staining of GFP FCSC transplant after 8 weeks. Serial section showing GFP⁺ FCSC under fluorescent light (green) and immunohistochemistry for osteocalcin (OCN, red). Dapi blue staining=nuclei. Scale bar=50 μm.



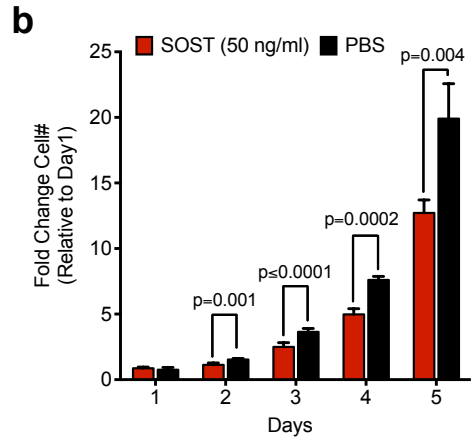
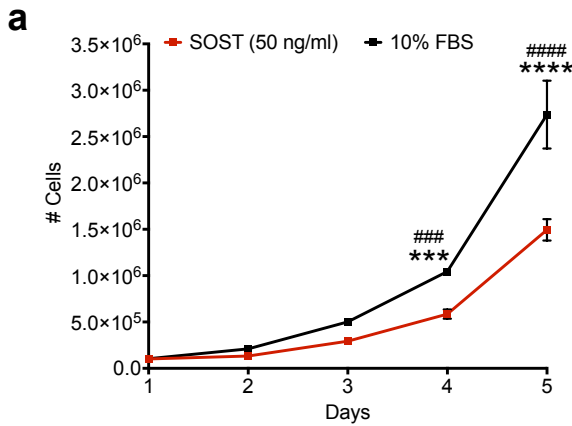
Supplementary Figure 4: FCSCs spontaneously generate organized bone

H&E staining of transplants show FCSCs generate bone with a hematopoietic microenvironment in multiple carriers after 8 weeks, but donor-matched condyle cells are unable to generate bone. Donor-matched FCSCs, bone marrow stromal cells (BMSCs), and condyle cells (CCs) were seeded onto (a) collagen sponge, (b) Matrigel or (c) gelfoam and transplanted on the dorsum on 8 week-old male nude mouse for 8 weeks. FCSCs form bone with mature osteocytes (OC) with an organized hematopoietic microenvironment including sinusoids, hematopoietic cell clusters (Hem), adipocytes, and osteoblasts (Obs). Scale bar = 50 μ m.



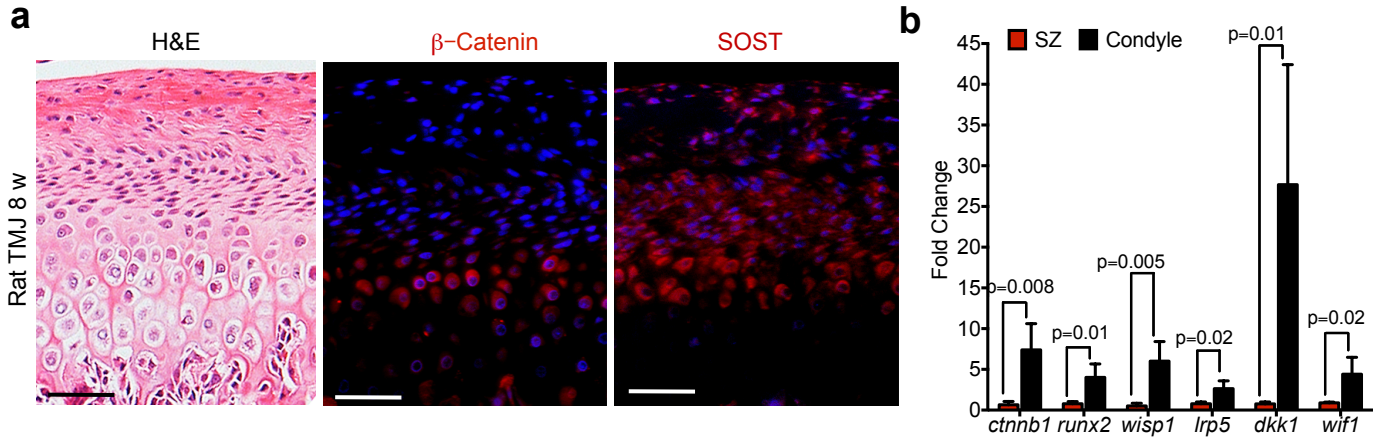
Supplementary Figure 5: FCSCs to differentiate into chondrocytes through inhibited Wnt

(a) FCSCs were treated with TGF- β 1, Wnt3a, Wnt5a and canonical Wnt inhibitors (Dkk1, Wif1, SOST). qRT-PCR was used to determine aggrecan expression in FCSCs after 24h, 48h and 72h. Data are mean fold changed normalized to GAPDH and relative to FBS or vehicle control \pm S.D.; n=3 independent experiments; two-way ANOVA followed by Tukey's post hoc test. (b) qRT-PCR of aggrecan (*acan*) expression in FCSCs with ICG-001 after 4d and 10d. Data are normalized to GAPDH and mean fold change relative to FBS/vehicle control \pm SD; n=3 independent experiments; one-way ANOVA followed by Tukey's post hoc test. (c) qRT-PCR was used to *sox5*, *sox9* and *acan* expression in FCSCs treated with Wnt3a after 2d and 7d. Data are normalized to GAPDH and mean fold change relative to FBS/vehicle control 2 days \pm S.D.; n=5 independent experiments; one-way ANOVA followed by Tukey's post hoc test.



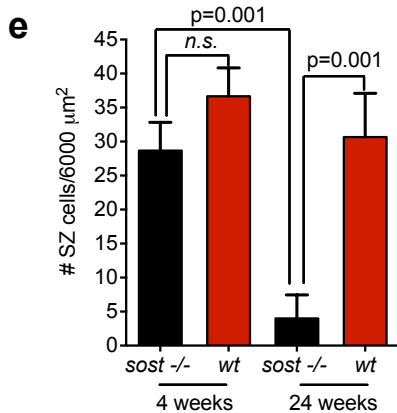
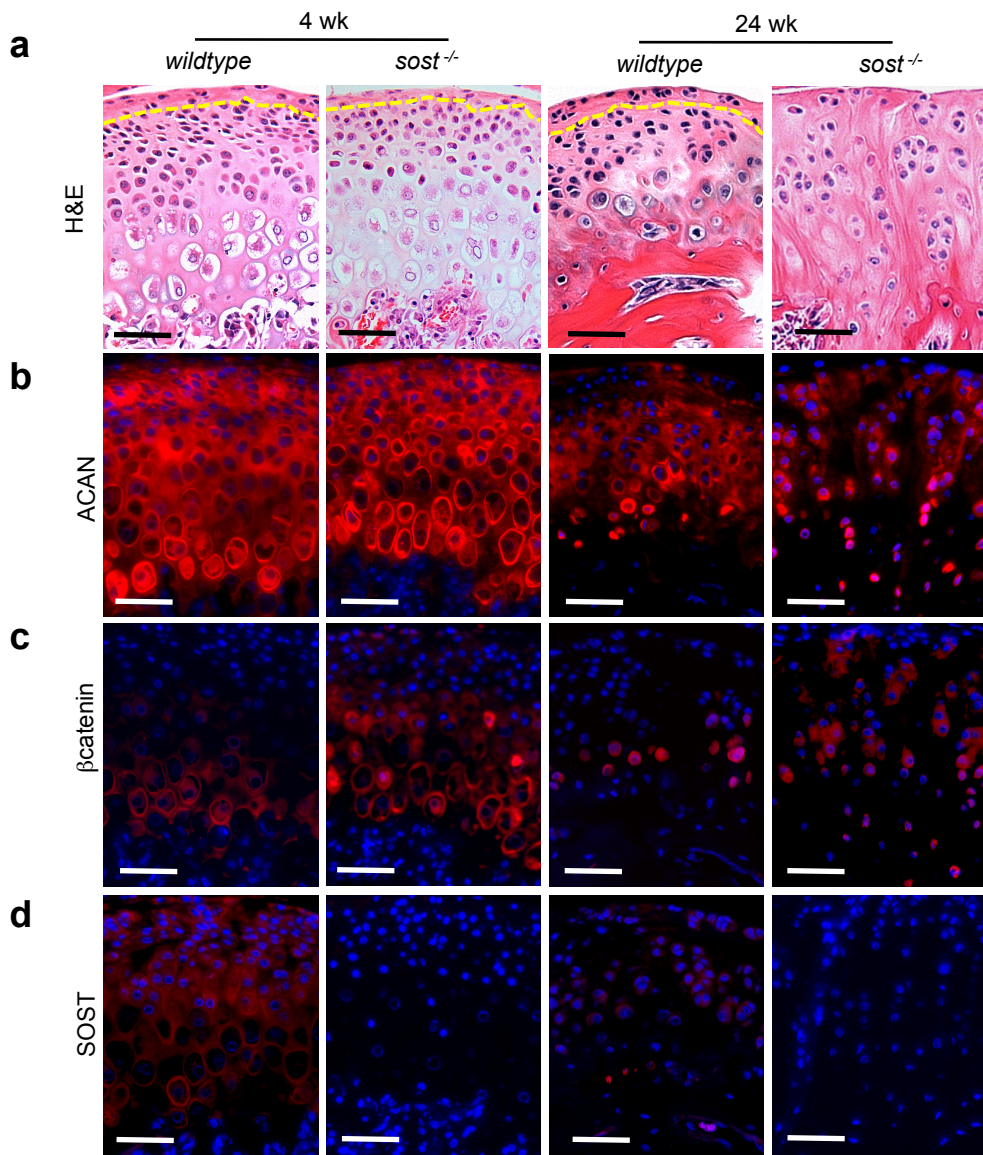
Supplementary Figure 6: Sclerostin inhibits FCSC proliferation

(a) Cell growth curve of FCSCs with SOST or vehicle over 5 days. Data are mean \pm S.D.; n=6 independent experiments; *** $p \leq 0.001$ SOST 4d vs vehicle 4d; ### $p \leq 0.001$ SOST 1d vs SOST 4d; **** $p \leq 0.0001$ SOST 5d vs vehicle 5d; ##### $p \leq 0.0001$ SOST 1d vs SOST 5d; two-way ANOVA followed by Tukey's post hoc test. (b) Data are mean fold change in cell number relative to 1d of SOST treated FCSCs and vehicle \pm S.D.; paired Student's t-test.



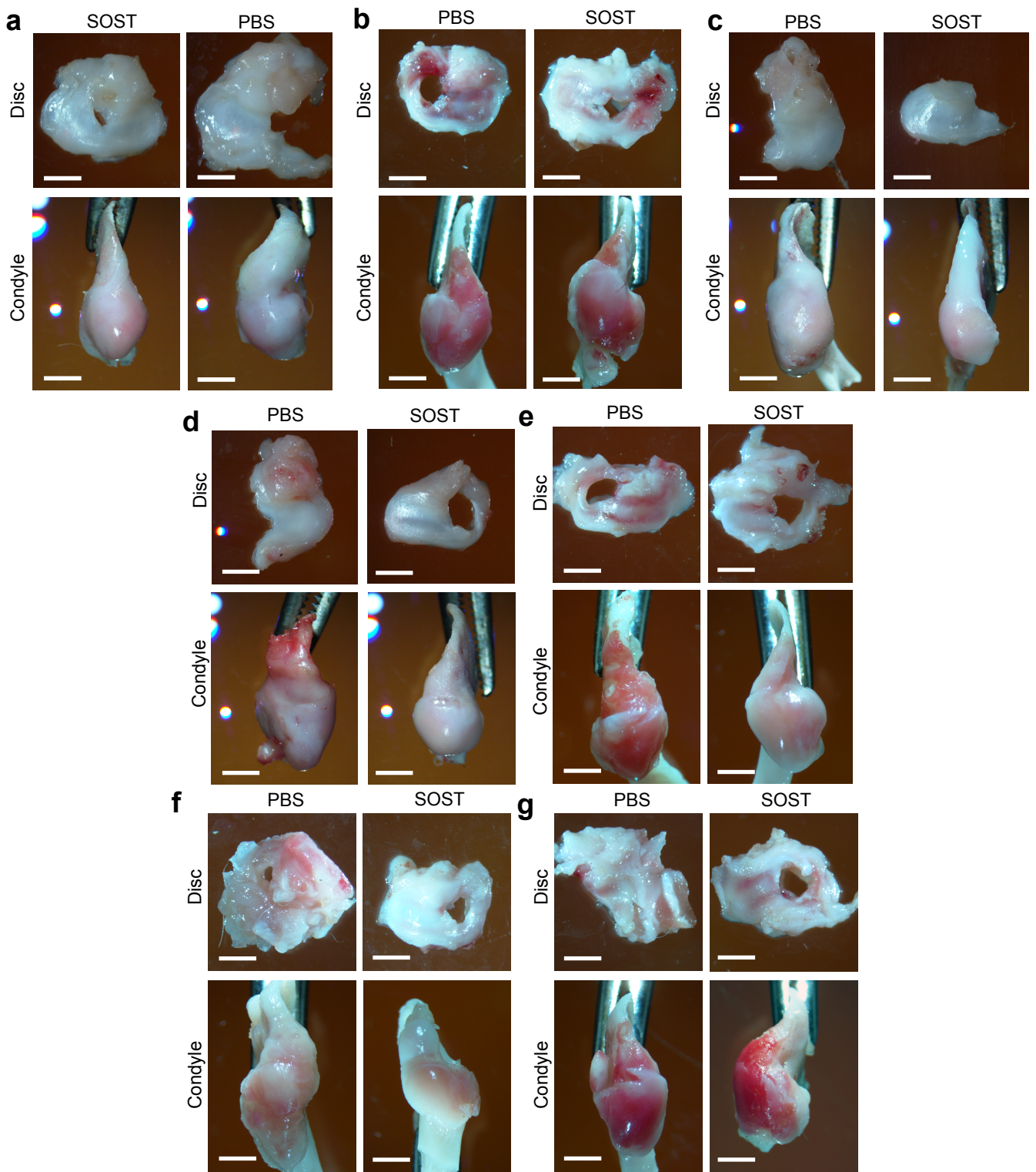
Supplementary Figure 7: Wnt signaling is temporally and spatially regulated in TMJ

(a) H&E and immunostaining for β Catenin (red) and sclerostin (SOST, red) in 8 week-old rat male TMJ condyle. Dapi=nucleus. Scale bar = 50 μ m. **(b)** qRT-PCR for wnt target genes (*ctnnb1*, *runx2*, *wisp1*, *lrp5*, *dkk1*, and *wif1*) using RNA extracted from SZ (red) and condyle (black) tissues. Data are normalized to GAPDH and mean fold change relative relative to SZ tissue; error bars are S.D.; n=4 rats male rats 8w; paired Student's t-test.



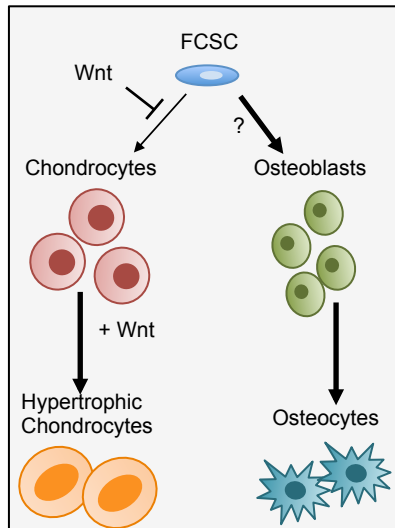
Supplementary Figure 8: Sclerostin deficiency in mice causes loss FCSCs

Mice deficient in sclerostin show decrease FCSCs, degeneration, and abnormal β catenin and aggrecan distribution. Scale bar = 50 μm . **(a)** H&E staining and immunohistochemical staining of **(b)** aggrecan (ACAN), **(c)** β Catenin (β Cat), and **(d)** sclerostin (SOST). **(e)** The number of FCSCs in SZ tissue in the comparable sagittal sections from WT and *sost*^{-/-} mandibular condylar cartilages at 4 and 24 weeks of age. Data are mean; error bars are S.D.; n=3 *sost*^{-/-} male mice 4 w and 24w; two-way ANOVA followed by Tukey's post hoc test.



Supplementary Figure 9: Sclerostin ameliorates TMJ degeneration in rabbits.

(a-g) TMJ pathology was surgically induced in 12w-old male New Zealand white male rabbits and a punch biopsy was used to create a 2.5 mm perforation in the TMJ disc bilaterally. 3 days post-surgery, 0.1 ml SOST (100ng/ml in PBS) was injected into the inferior TMJ intra-articular space unilaterally on the right (a) or left (b-g) once weekly for 8 weeks. Vehicle control (PBS) was injected into the contra-lateral TMJ. Images are photographs that show the superior view of rabbit TMJ disc and condyle 8 weeks following bilateral disc perforation surgery. SOST was injected into the inferior TMJ intra-articular space unilaterally once weekly for 8 weeks, while vehicle control (PBS) was injected into the contra-lateral TMJ within the same animal. Scale bar = 3mm.



Supplementary Figure 10: FCSC fate specification is regulated by Wnt signaling.

A single FCSC can spontaneously commit to bone and cartilage lineages. Canonical Wnt signals inhibit FCSC differentiation into chondrocytes and promote chondrocyte terminal differentiation.

Supplementary Table 1: Single cell clonal multi-differentiation potential analysis

A total of 31 single cell FCSC clonal progenies were isolated and their ability to undergo adipogenesis, chondrogenesis and osteogenesis was tested *in vitro*.

| | |
|--|----------------------|
| Total Number of Clones Isolated | 31 Clones |
| Tri-lineage Differentiation Potential | 22.5% (7/31 clones) |
| Bi-lineage Differentiation Potential | 64.5% (20/31 clones) |
| Single Lineage Differentiation Potential | 12.9% (4/31 clones) |

Supplementary Table 2: Human TMJ fibrocartilage Analyses

12 patient samples were obtained from NIDCR TIRR that underwent TMJ condyle fibrocartilage replacement surgery. The osteoarthritic histopathological score was graded and β -catenin expression was quantified.

| Research ID | Age DOS | Gender | Race | Stage | Grade | Score | # β catenin cells/ $\mu\text{m}^2 \times 10^6$ |
|-------------|---------|--------|---------|-------|-------|-------|---|
| T06359 | 29 | F | Asian | 4 | 3.5 | 14 | 142.45 |
| T06368 | 33 | F | White | 4 | 4.0 | 16 | 155.40 |
| T06353 | 22 | F | White | 4 | 4.0 | 16 | 366.95 |
| T08620 | 25 | F | White | 4 | 4.0 | 16 | 179.90 |
| T06313 | 52 | F | White | 4 | 4.4 | 18 | 453.03 |
| T06017 | 22 | F | Unknown | 4 | 4.5 | 18 | 449.35 |
| T07616 | 35 | F | White | 4 | 4.5 | 18 | 540.35 |
| T06554 | 41 | F | White | 4 | 5.0 | 20 | 351.44 |
| T09634 | 39 | M | White | 4 | 5.0 | 20 | 366.55 |
| T07610 | 52 | F | White | 4 | 6.0 | 24 | 334.79 |
| T06309 | 38 | M | White | 4 | 6.0 | 24 | 178.19 |
| T06347 | 42 | F | White | 4 | 6.0 | 24 | 100.65 |

Supplementary Table 3: Antibodies used for immunohistochemistry and flow cytometry

| Primary Antibody | Source | Secondary Antibody | Isotype Negative Control |
|--|-------------------------------|---|---|
| Rabbit anti- Aggrecan | Millipore AB1031 (1:100) | Invitrogen A-11010 (1:1000) or Invitrogen A-11008 (1:1000) | Rabbit IgG R&D AB-105-C |
| Rabbit anti- Lubricin | Novus Biologics 19048 (1:100) | Invitrogen A-11010 (1:1000) | Rabbit IgG R&D AB-105-C |
| Mouse anti- Type II Collagen | Millipore MAB8887 (1:50) | Invitrogen A-11003 (1:1000) or Invitrogen A-11001 (1:1000) | Mouse IgG1 R&D MAB002 |
| Rabbit Anti- Osteocalcin | Millipore AB10911 (1:100) | Invitrogen A-11010 (1:1000) | Rabbit IgG R&D AB-105-C |
| Rabbit Anti- β Catenin (for mouse, rat, and human) | Abcam AB6302 (1:50) | Invitrogen A-11010 (1:1000, rodent) (1:2500, human) | Rabbit IgG R&D AB-105-C |
| Mouse Anti- β Catenin (for rabbit) | Millipore Mab2081 (1:100) | Invitrogen A-11001 (1:1000) | Mouse IgG1 R&D MAB002 |
| Rabbit Anti- Sclerostin | Abcam AB63097 (1:50) | Invitrogen A-11010 (1:1000) | Rabbit IgG R&D AB-105-C |
| Mouse Anti- CD90 (PE) | Abcam AB33694 | N/A | Mouse IgG1 (PE) Abcam AB91357 |
| Hamster Anti- CD29 (Pacific Blue) | Biolegend 102224 | N/A | Hamster IgG (Pacific Blue) Biolegend 400925 |
| Mouse anti- CD44 (PE) | Abcam AB23396 | N/A | Mouse IgG1 (PE) Abcam AB91357 |
| Mouse anti- CD105 | Novus Biologics NBP1-42383 | Novus Biologics anti Mouse IgG Antibody (FITC) NB720-F | Mouse IgG1 R&D MAB002 |
| Mouse anti- CD146 (APC) | R&D FAB3250A | N/A | Mouse IgG2a (APC) R&D IC003A |
| Mouse anti- CD45 (PerCp) | Biolegend 202220 | N/A | Mouse IgG1 (PerCP) Biolegend 400150 |
| Mouse anti- CD79a (APC) | Abcam AB188420 | N/A | Mouse IgG1 (PE) Abcam AB67435 |
| Mouse anti- CD11b (FITC) | BD Biosciences 554982 | N/A | Mouse IgA (FITC) 553478 |

Supplementary Table 4: Primers used for qRT-PCR

| Gene Symbol | Taqman Gene® Expression Assay |
|----------------|----------------------------------|
| <i>gapdh</i> | Rn01775763_g1 |
| <i>sox6</i> | Rn00488400_m1 |
| <i>sox9</i> | Rn01751069_mH |
| <i>acan</i> | Rn00573424_m1 |
| <i>col2a1</i> | Rn01637081_g1 |
| <i>col10a1</i> | Rn01408030_m1 |
| <i>bgn</i> | Rn00567229_m1 |
| <i>fmod</i> | Rn00589918_m1 |
| <i>dcn</i> | Rn01503161_m1 |
| <i>scx</i> | Rn01504576_m1 |
| <i>col1a1</i> | Rn01463848_m1 |
| <i>prg4</i> | Rn01490812_m1 |
| <i>sox5</i> | Rn01492418_m1 |
| <i>ppar-γ</i> | Rn00440945_m1 |
| <i>ocn</i> | Rn01455285_g1 |
| <i>twist1</i> | Rn00585470_s1 |
| <i>ccnd1</i> | Rn00432360_m1 |
| <i>ctnnb1</i> | Rn00584431_g1 |
| <i>runx2</i> | Rn01512298_m1 |
| <i>wisp1</i> | Rn01505161_m1 |
| <i>lrp5</i> | Rn01451428_m1 |
| <i>dkk1</i> | Rn01501537_m1 |
| <i>wif1</i> | Rn00586968_m1 |