Description of additional supplementary files

Supplementary Data 1. Enriched gene list by cluster for whole calvaria dataset (related to Fig. 1B).

Supplementary Data 2. Enriched gene list by cluster for connective and skeletogenic subset (related to Fig. 1C).

Supplementary Data 3. Enriched gene list by cluster for osteoblast and prrx1a+ mesenchyme subset (related to Fig. 1E).

Supplementary Data 4. CellChat Signaling pairs for the prrx1a+ mesenchyme subset.

Supplementary Data 5. Genotyping primers for mutant alleles.