

**SUPPLEMENTARY FIG. S1.** (A) Chromatographic separation of noncollagenous proteins (NCPs) extracted from the long bone of rats. To identify the major acidic protein components in the bone extracts, the extracts were subjected to a Q-Sepharose ion-exchange (positive charge beads) chromatography with a gradient ranging 0.1–0.8 M NaCl in 6 M urea solution (pH 7.4). The peaks of the elution profile represent the total protein eluted from the chromatography at specific time points with relative concentration of NaCl. The Q-Sepharose column separated NCPs into 120 fractions. Each fraction contained 0.5 mL of 6 M urea solution. By using our method described in the Materials and Methods section, the NCPs from the bone matrix were mainly eluted into fractions from 37 to 87. (B) Stains-All staining was performed to evaluate the NCPs eluted from the ion-exchange chromatography. All the protein bands pointed by arrows in Stains-All staining figures were confirmed by using western immunoblotting with specific antibodies. The identity of dentin matrix protein 1 (DMP1) fragments, osteopontin (OPN), and bone sialoprotein (BSP) was confirmed by western immunoblotting analyses. Western immunoblotting *et al.*, 2010, Oral Diseases)<sup>S1</sup>.



**SUPPLEMENTARY FIG. S2.** Attenuated total reflectance Fourier Transform Infrared (ATR-FTIR) spectroscopy of the nanofibrous gelatin (NF-Gelatin) before and after surface modified with NCPs. (a) NCPs; (b) NF-Gelatin; (c) NF-Gelatin. NCPs. As marked with the dotted lines, the curve (c) includes the absorption peaks of both (a) and (b), indicating that NCPs have been successfully coupling onto the surfaces of the NF-Gelatin.

## **Supplementary Reference**

 Zhang, B., Sun, Y., Chen, L., Gunn, C., Guo, L., and Qin, C. Expression and distribution of SIBLING proteins in the predentin/dentin and mandible of hyp mice. Oral Dis 16, 453, 2010.