## Supplementary figures

Supplementary Figure 1. (A) Immunofluorescence staining comparison of CXCL8 (green) in healthy cartilage and preserved areas of OA cartilage. Nuclei are stained using propidium iodide. Scale bar, 100 $\mu$ m. (B) Safranin orange staining of healthy articular cartilage and preserved areas of OA articular cartilage. Scale bar, 100 $\mu$ m. (C) Immunofluorescence staining of mouse CXCL1 (green) in articular cartilage from sham and DMM operated mouse articular cartilage. Nuclei are stained using DAPI (blue). Scale bar, 100 $\mu$ m. (D) Real time RT-PCR for CXCL8 mRNA in early and late passage human articular chondrocytes (n = 3), \*\*P < 0.01 by paired t-test.



Supplementary Figure 2. (A) Histological comparison of 10 week old CXCR2- /- mouse articular cartilage to wild type controls stained using Safranin orange. Scale bar, 200µm. (B, C) Immunofluorescence staining and quantification of type II collagen and type X collagen (green) in unchallenged wild type and CXCR2-/- articular cartilage. Nuclei are stained using DAPI (blue). Scale bar, 100µm.



Supplementary Figure 3. Immunofluorescence staining of phospho-AKT (473) (red) in wild type sham operated control and DMM operated knees 8 weeks following surgery. Nuclei are stained using DAPI (blue). Scale bar,  $100\mu m$ .



Supplementary Figure 4. (A) Quantification of TUNEL postive chondrocytes in superfical and deep zones of articular cartilage of unchallenged wild type and CXCR2-/- mice (n = 4). (B) Quantification of TUNEL positive in vitro cultured chondrocytes from wild type and CXCR2-/- mice (n = 4).



Supplementary Figure 5. (A) Immunohistochemical staining for Ly-6B.2 in hind paws of hTNFtg mutant mice. Scale bar,  $100\mu m$  (B) Immunohistochemical staining for Ly-6B.2 in knee joints of wild type and CXCR2-/- mice 8 weeks following DMM surgery. Scale bar,  $100\mu m$ .

