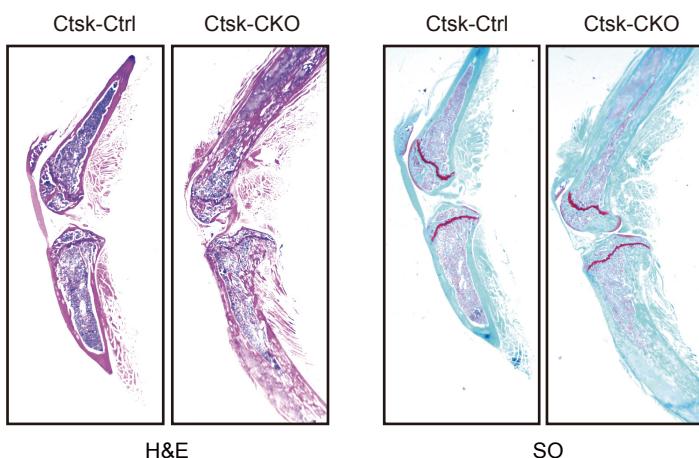
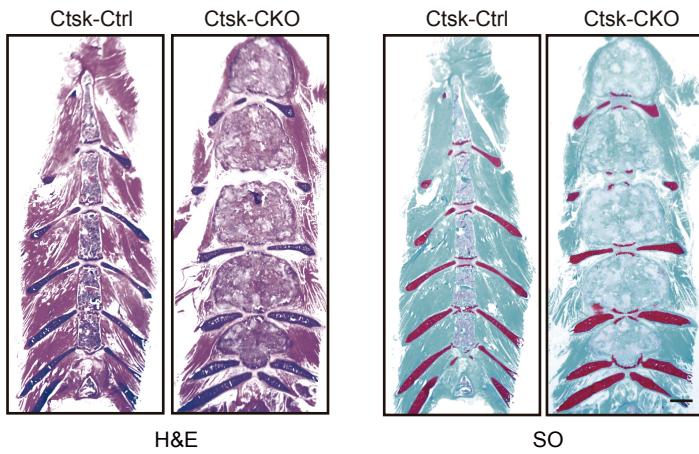


**A**

H&amp;E

SO

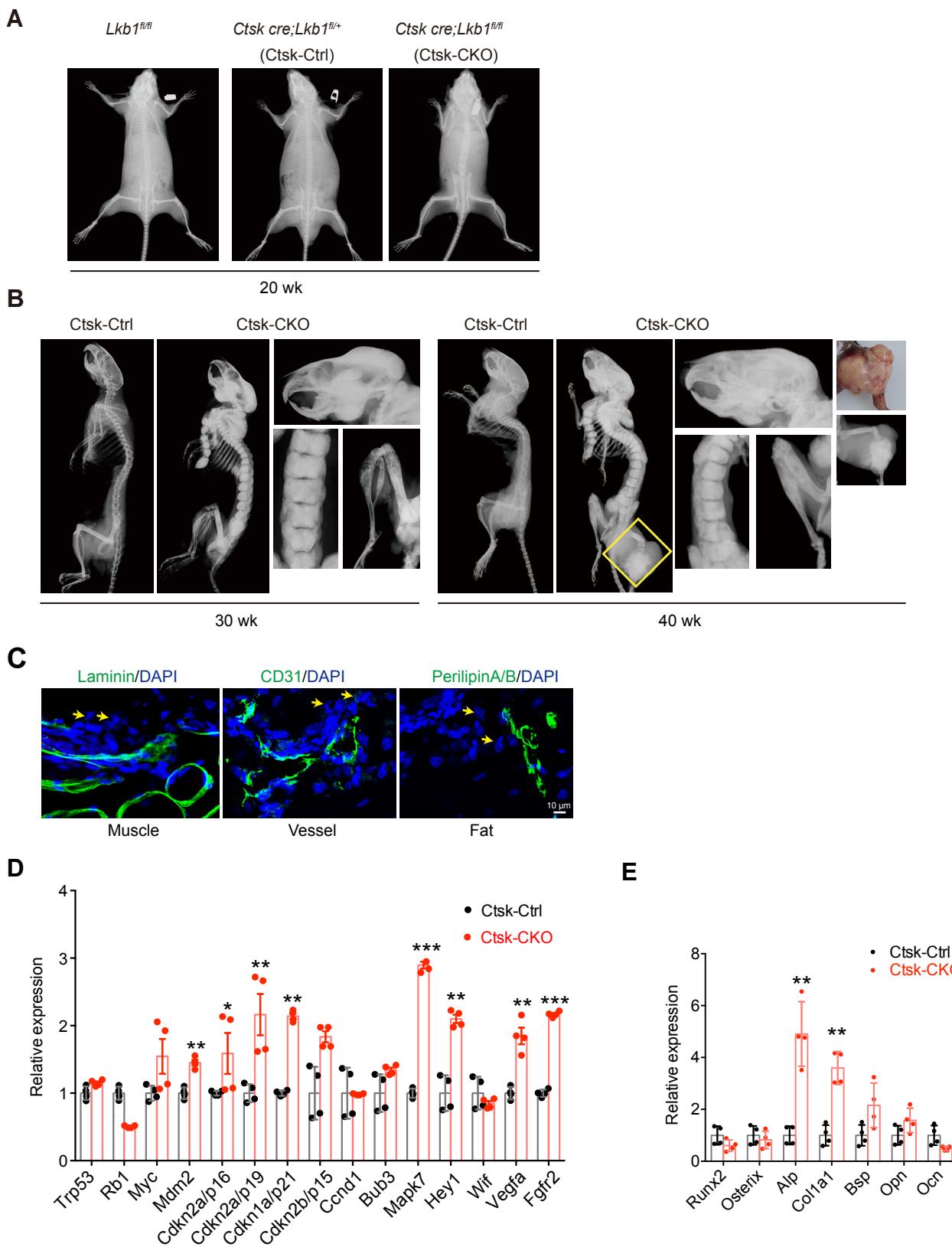
**B**

H&amp;E

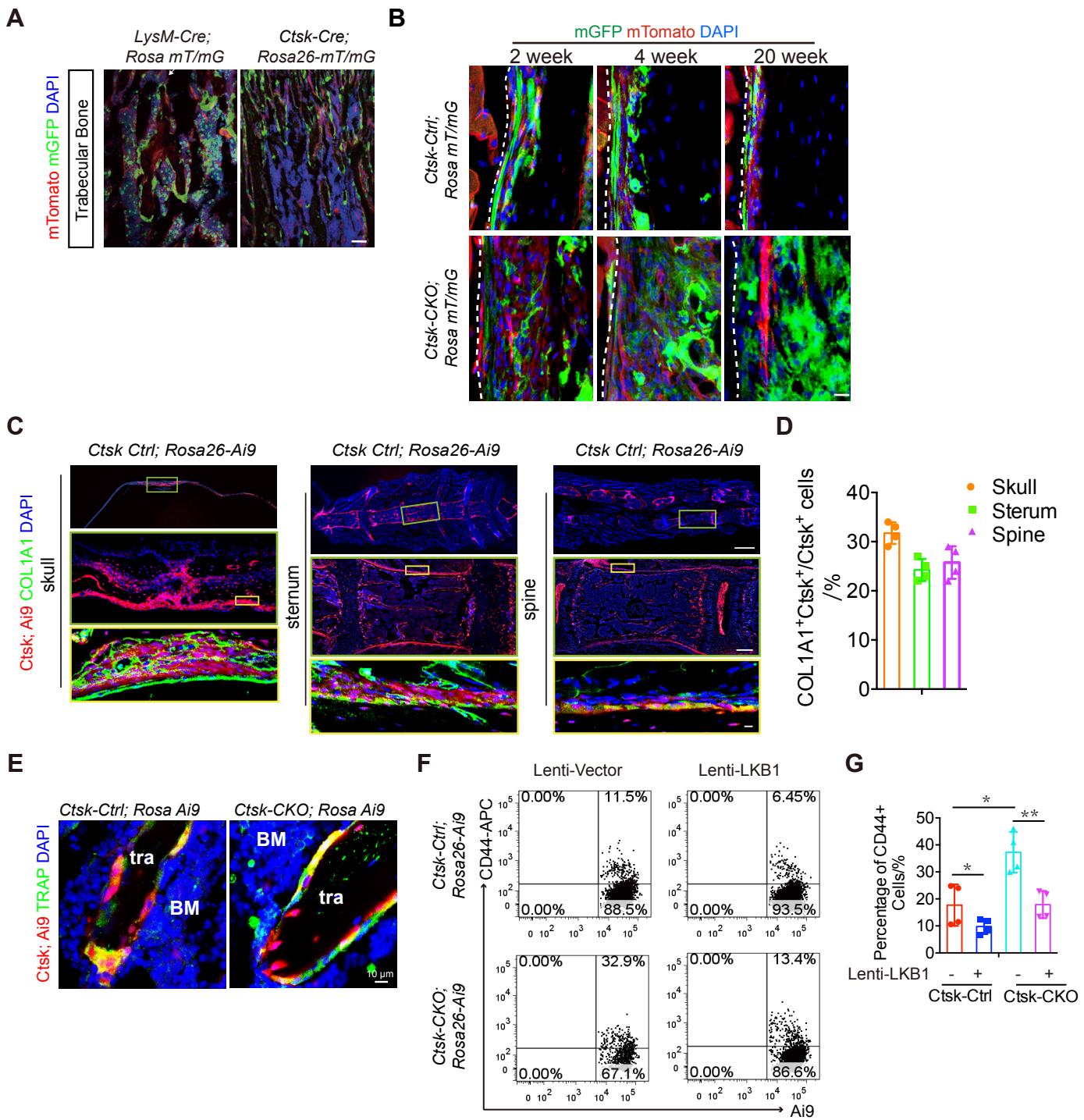
SO

**Supplemental Figure 1. *Lkb1* deletion in *Ctsk*-Cre expressing cells dose not show cartilage tumors.**

(A and B) H&E staining and Safranin O (SO) staining of the tibiae (A) and sternums (B) of 20-week-old Ctsk-Ctrl and Ctsk-CKO mice showed that Ctsk-CKO mice did not display cartilage tumor. Scale bars, 300  $\mu$ m. Similar results were obtained from analyses of at least 3 mice of each genotype.



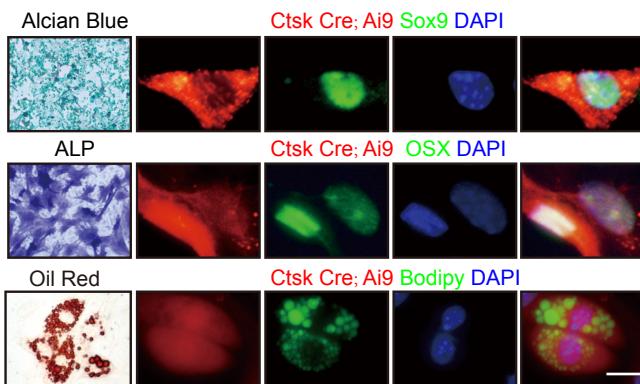
**Supplemental Figure 2. *Lkb1* deletion in *Ctsk*-Cre expressing cells causes osteogenic tumor-like phenotype.**  
**(A)** X-ray images of the whole-body of 20-week-old *Lkb1<sup>fl/fl</sup>*, *Ctsk Cre; Lkb1<sup>fl/+</sup>* and *Ctsk Cre; Lkb1<sup>fl/fl</sup>* mice. **(B)** X-ray images of the whole-body skeletons, spines, tibiae, and skulls of the 30-week-old and 40-week-old *Ctsk*-Ctrl and *Ctsk*-CKO mice. **(C)** Immunostaining of markers of fat (Perilipin A/B), vessels (CD31) and muscles (Laminin) in the tumor region of 40-week-old mice. **(D and E)** qPCR analysis of osteosarcoma-related genes at 5-week-old (**D**) and osteoblast markers at 2-week-old (**E**). Data represent means  $\pm$  SEM. \* $P$  < 0.05, \*\* $P$  < 0.01, and \*\*\* $P$  < 0.001, by unpaired Student's *t* test (**D** and **E**).



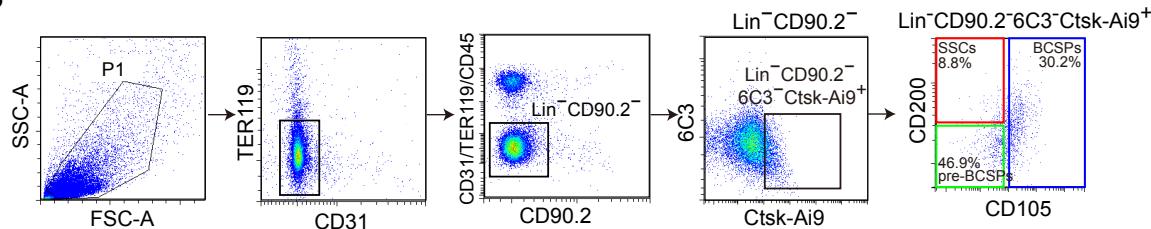
**Supplemental Figure 3. Ctsk<sup>+</sup> periosteal mesenchymal stem cells are novel potential sources for osteogenic tumor.**

(A) Confocal images of the tibiae of 4-week-old male *LysM-Cre; Rosa26-mT/mG* and *Ctsk-Cre; Rosa26-mT/mG* mice showing both *Ctsk-Cre* and *LysM-Cre* were expressed in osteoclasts on the surface of trabecular bone of 4-week-old mice, and *LysM-Cre* were expressed in macrophages in the bone marrow. Scale bar, 20  $\mu$ m. (B) *Ctsk<sup>+</sup>* cells in the periosteum of 2-, 4-, 20- week-old male *Ctsk-Ctrl; Rosa mT/mG* and *Ctsk-CKO; Rosa mT/mG* mice. Scale bar, 20  $\mu$ m. (C and D) Images of the skull, sternum and spine of *Ctsk-Cre; Rosa26-Ai9* mice showing *Ctsk-Cre* cells (red) in the periosteum expressed osteoblast marker COL1A1. Scale bar, 1000  $\mu$ m (up), 50  $\mu$ m (middle), 10  $\mu$ m (below). (E) Immunostaining of TRAP in trabecular bone showing that most of *Ctsk<sup>+</sup>* cells in trabecular bone are TRAP<sup>+</sup> osteoclasts. Scale bar, 10  $\mu$ m. (F and G) Flow cytometry analysis showing that *Ctsk-Ai9* cells isolated from the cortical bone of *Ctsk-CKO; Rosa26-Ai9* tibiae displayed increased CD44 expression and re-expression of LKB1 using a lentivirus reduced number of CD44 positive cells in *Ctsk-Ai9* cells ( $n = 4$ ). Data represent means  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , and \*\*\* $P < 0.001$ , by two-way ANOVA (G).

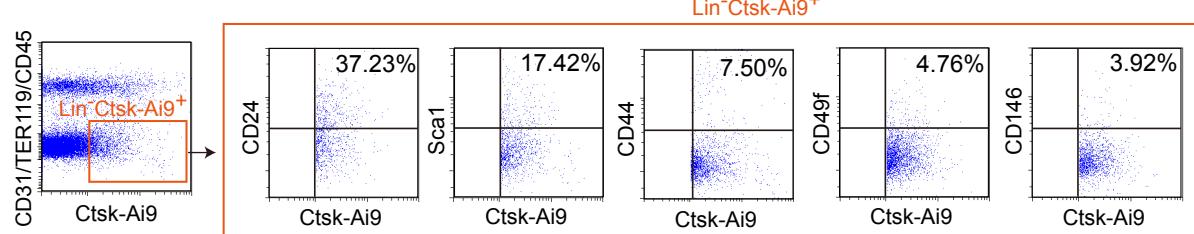
A



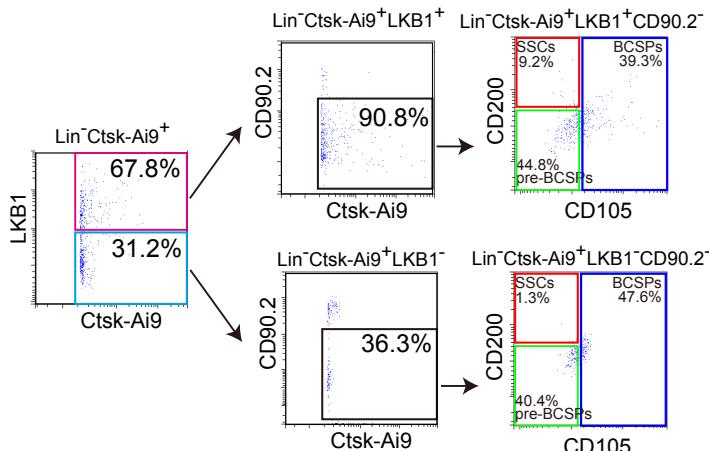
B



C

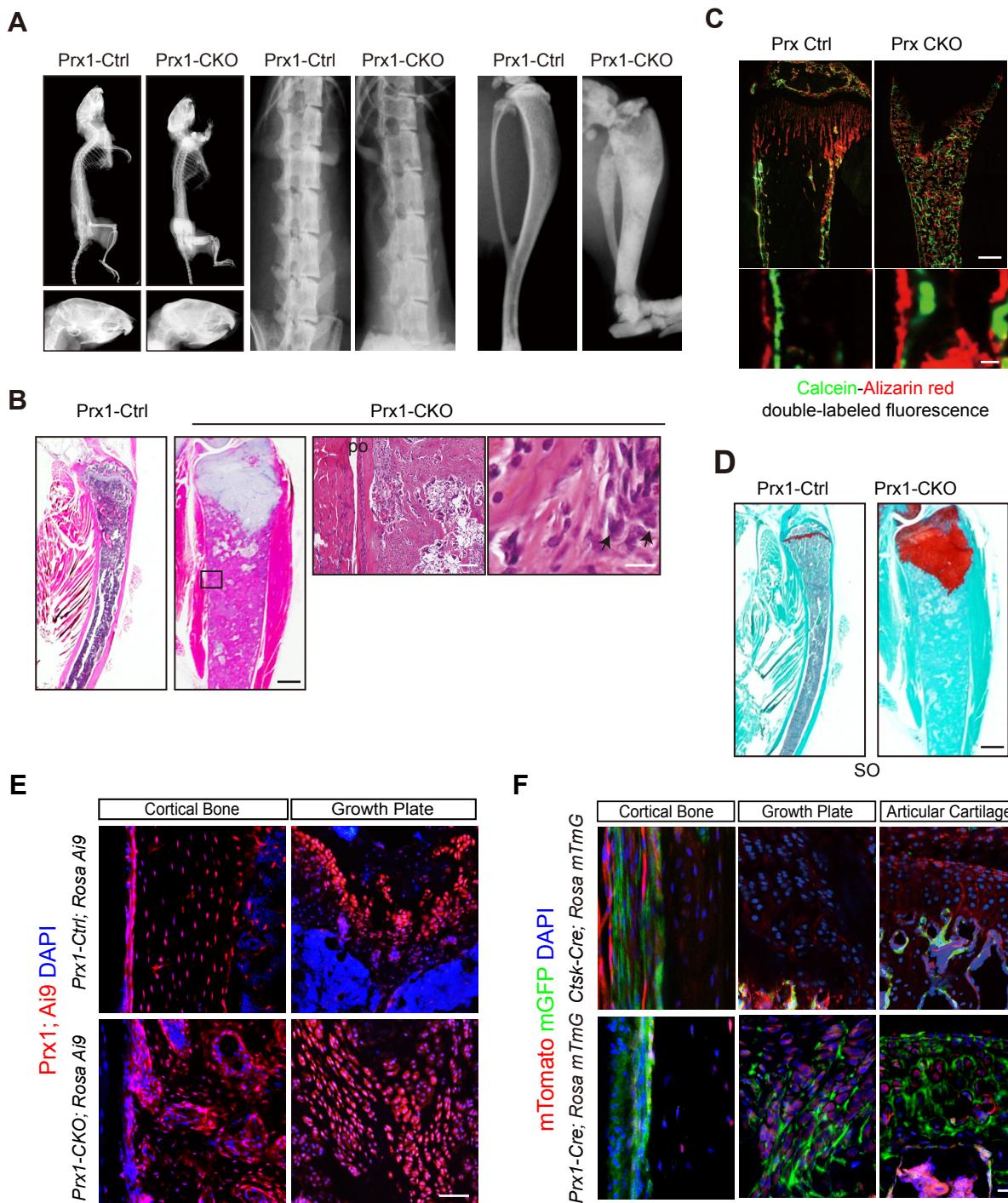


D



**Supplemental Figure 4. Ctsk-Cre expressing cells display mesenchymal stem cell markers and differentiation potential.**

(A) Ctsk-Ai9 cells were able to differentiate into chondrocyte (Alcian Blue staining and Sox9 immunostaining), osteoblast (ALP staining and OSX immunostaining) and adipocyte (Oil Red O and Bodipy staining). Scale bar, 10  $\mu$ m. (B and C) Flow cytometry analysis of Ctsk-Ai9 cells from the periosteum of cortical bones of 4-week-old Ctsk-Ctrl; Rosa26-Ai9 mice. (D) Flow cytometry analysis of periosteum-derived Ctsk<sup>+</sup>Lkb1<sup>+</sup> and Ctsk<sup>+</sup>Lkb1<sup>-</sup> cells.

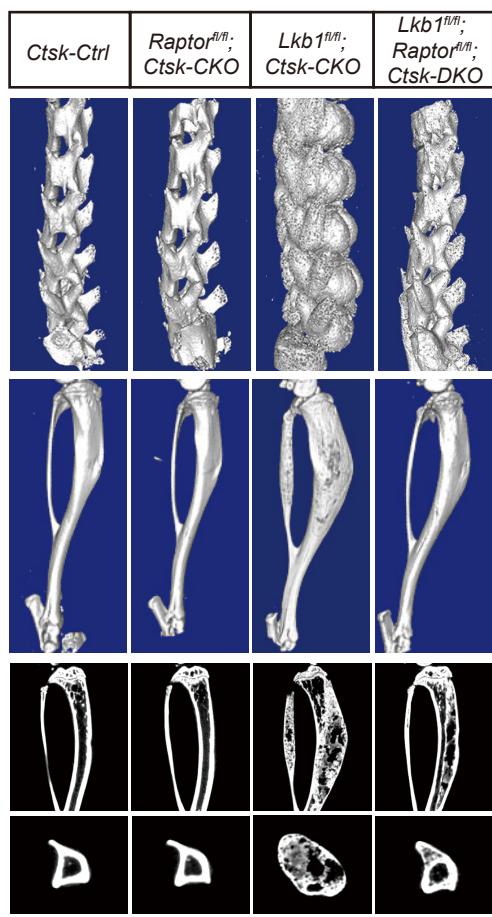


**Supplemental Figure 5. Loss of *Lkb1* in *Prx1-Cre* expressing cells causes both osteogenic tumor and cartilage tumor.**

(A) X-ray images of the whole-body skeletons, skulls, spines and long bones of 20-week-old Prx1-Ctrl and Prx1-CKO mice. (B) Representative images of H&E staining of the tibiae from 20-week-old Prx1-Ctrl and Prx1-CKO mice showing both osteogenic tumor and cartilage tumor formation. Scale bar, 1 mm (left), 100 $\mu$ m (middle), 20  $\mu$ m (right). (C) Calcein-alizarin red double-labeled fluorescence showing increased bone formation in 5-week-old Prx1-CKO mice. Scale bars, 1 mm (up) 100 $\mu$ m (below). (D) SO staining of the tibiae from 20-week-old Prx1-Ctrl and Prx1-CKO mice. Scale bars, 1 mm. (E) Confocal imaging of Prx1-Ai9 cells showing howing Prx1<sup>+</sup> cells were expanded from the cortex to the marrow cavity in the periosteum and also expanded to form a mass of cartilage in the growth plate of Prx1-CKO mice. Scale bar, 50  $\mu$ m. (F) Confocal images of the tibiae of 4-week-old *Ctsk-Cre; Rosa26-mT/mG* reporter mice and *Prx1-Ctrl; Rosa26-mT/mG* reporter mice including cortical bone, growth plate and articular cartilage. Scale bar, 20  $\mu$ m.

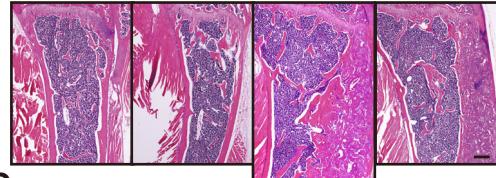
## Supplemental Figure 6

**A**



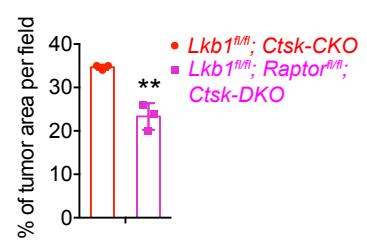
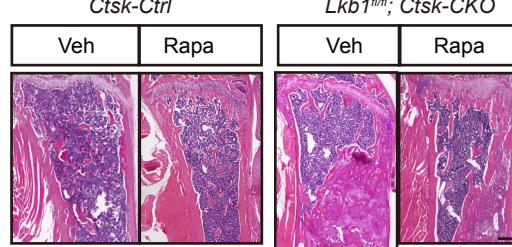
**B**

Ctsk-Ctrl	<i>Raptor</i> <sup>f/f</sup> ; Ctsk-CKO	<i>Lkb1</i> <sup>f/f</sup> ; Ctsk-CKO	<i>Lkb1</i> <sup>f/f</sup> ; <i>Raptor</i> <sup>f/f</sup> ; Ctsk-DKO
-----------	---	---------------------------------------	--



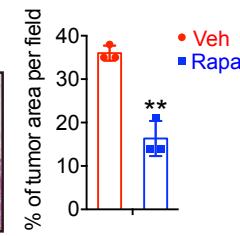
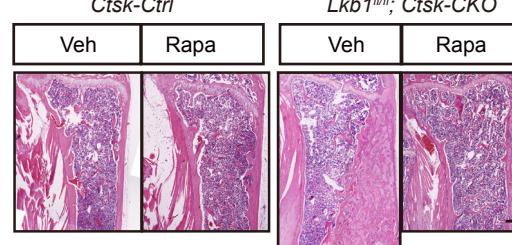
**C** Treatment initiation (twice per week) ↓ Analysis ↓

2 weeks      18 weeks

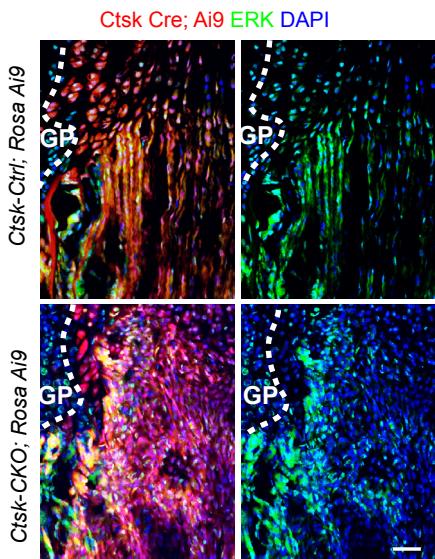
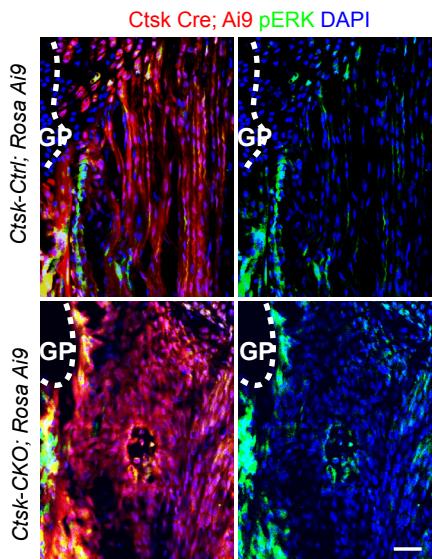
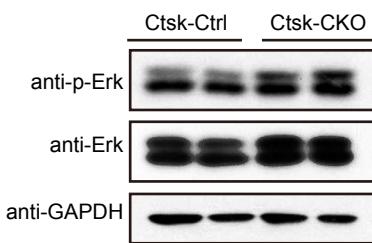
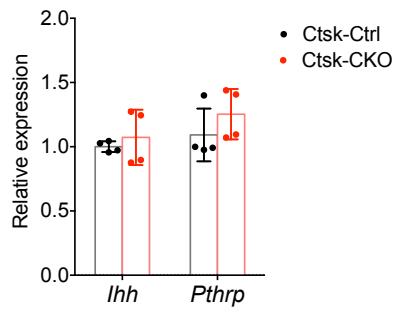
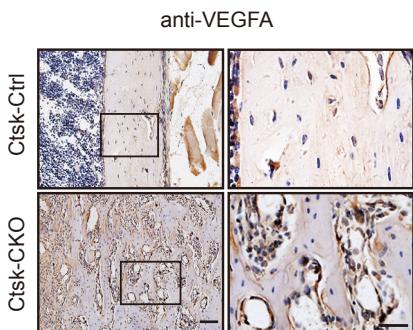
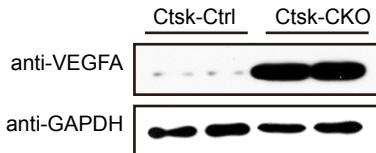
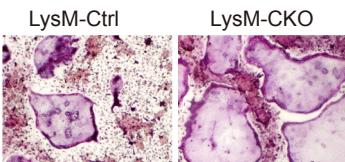
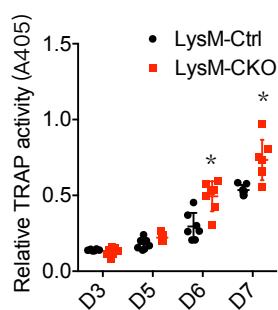


**D** Treatment initiation (daily) ↓ Analysis ↓

16 weeks      4 weeks



**Supplemental Figure 6. Inhibition of mTOR signaling ameliorates the tumor progression of Ctsk-CKO mice.**  
**(A)** MicroCT scans of spines and tibiae from 20-week-old Ctsk-Ctrl, *Raptor*<sup>f/f</sup>; Ctsk-CKO, *Lkb1*<sup>f/f</sup>; Ctsk-CKO and *Lkb1*<sup>f/f</sup>; *Raptor*<sup>f/f</sup>; Ctsk-DKO mice. **(B)** H&E staining of the tibiae from Ctsk-Ctrl, *Raptor*<sup>f/f</sup>; Ctsk-CKO, *Lkb1*<sup>f/f</sup>; Ctsk-CKO and *Lkb1*<sup>f/f</sup>; *Raptor*<sup>f/f</sup>; Ctsk-DKO mice. Scale bar, 300  $\mu$ m. **(C and D)** H&E staining showing the treatment of Ctsk-Ctrl and Ctsk-CKO mice with rapamycin twice per week from 2-week-old to 20-week-old (**C**) or daily from 16-week-old to 20-week-old (**D**) delayed the tumor growth of Ctsk-CKO mice. Scale bars, 300  $\mu$ m. Data represent means  $\pm$  SEM. \*P < 0.05, \*\*P < 0.01, and \*\*\*P < 0.001, by unpaired Student's t test (**B, C and D**).

**A****B****C****D****E****F****G****H**

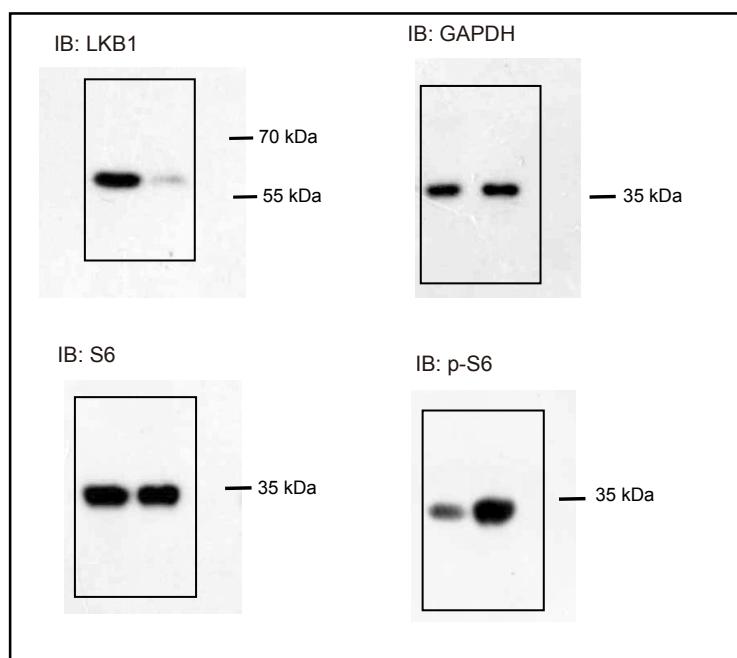
### Supplemental Figure 7. Effect of Lkb1 loss on Ihh, VEGF and osteoclastogenesis.

(A and B) Immunostaining of ERK (green) (A) and pERK (green) (B) in Ctsk<sup>+</sup> (red) cells. (C) Western blot showing expression of ERK and pERK in the tumor region of Ctsk-CKO mice and the cortical bones of control mice. Scale bar, 50 µm. (D) qPCR showing Ihh and PthrP production in Ctsk-CKO and control mice. GP: growth plate. Immunostaining (E) and western blot (F) of VEGFA in the tumor region of Ctsk-CKO mice and the cortical bones of control mice. Scale bar, 100 µm (left), 100 µm (right). (G and H) TRAP staining (G) and quantification of TRAP activity (H) of cells differentiated from LysM-Ctrl and LysM-CKO bone marrow (BM) cells in the presence of M-CSF and RANKL. Data represent means ± SEM. \*P < 0.05, \*\*P < 0.01, and \*\*\*P < 0.001, by unpaired Student's t test (H).

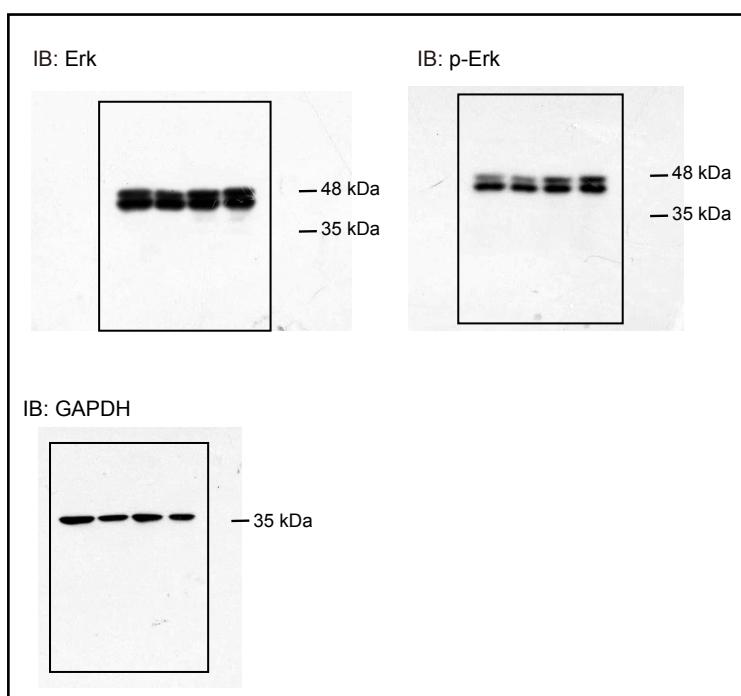
**TableS1- qPCR Primer Sequence**

<b>NO.</b>	<b>PrimerName</b>	<b>Sequence(5'-3')</b>
1	mTrp53-qF	CTCTCCCCGCAAAAGAAAAA
	mTrp53-qR	CGGAACATCTCGAACGCGTTA
2	mRb1-qF	TGCATCTTATCGCAGCAGTT
	mRb1-qR	GTCACACGTCCGTTCTAATTG
3	mMyc-qF	ATGCCCTCAACGTGAACCTC
	mMyc-qR	CGAACATAGGATGGAGAGCA
4	mMdm2-qR	TGTCTGTCTACCGAGGGTG
	mMdm2-qF	TCCAACGGACTTTAACAACTTCA
5	mCdkn2a/p16-qF	AACTCTTCGGTCGTACCCC
	mCdkn2a/p16-qR	GCGTCTTGAGCTGAAGCTA
6	mCdkn2a/p19-qF	CGCAGGTTCTGGTCACTGT
	mCdkn2a/p19-qR	TGTTCACGAAAGCCAGAGCG
7	mCdkn1a/p21-qR	CCTGGTATGTCCGACCTG
	mCdkn1a/p21-qF	CCATGAGCGCATCGCAATC
8	mCdkn2b/p15-qR	CCCTGCCACCCTTACCAAGA
	mCdkn2b/p15-qF	CAGATACTCGCAATGTCACG
9	mCcnd1-qF	GCGTACCCCTGACACCAATCTC
	mCcnd1-qR	CTCCTCTCGCACCTCTGCTC
10	mBub3-qF	GATGGCATCTCCTCGGTTAAG
	mBub3-qR	AATTGGCGGGCACATCGTAG
11	mMapk7-qF	GTGGGGGACGAGTACGAGAT
	mMapk7-qR	TGGTCACCAACATCAAAGCAT
12	mHey1-qF	GCGCGGACGAGAATGGAAA
	mHey1-qR	TCAGGTGATCCACAGTCATCTG
13	mWif1-qF	TCTGGAGCATCCTACCTTGC
	mWif1-qR	ATGAGCACTCTAGCCTGATGG
14	mVegfa-qF	TTACTGCTGTACCTCCACC
	mVegfa-qR	ACAGGACGGCTTGAAGATG
15	mFgfr2-qR	GCCTCTCGAACAGTATTCTCCT
	mFgfr2-qF	ACAGGGTTCATAGGCATGGG
16	mHprt-qF	GTAAAGCAGTACAGCCCCAAA
	mHprt-qR	AGGGCATATCCAACAACAAACTT
17	mRunx2-qF	CCAACCGAGTCATTAAGGCT
	mRunx2-qR	GCTCACGTCGCTCATCTTG
18	mOsx-qF	ATGGCGCCTCTCTGCTTGA
	mOsx-qR	GAAGGGTGGTAGTCATTG
19	mAlp-qF	CGGGACTGGTACTCGGATAA
	mAlp-qR	ATTCCACGTCGGTTCTGTTTC
20	mCol1a1-qF	GCTCCTCTAGGGGCCACT
	mCol1a1-qR	CCACGTCTCACCATGGGG
21	mBsp-qF	GACTTTGAGTTAGCGGCACT
	mBsp-qR	CCGCCAGCTCGTTTCATC
22	mOprn-qF	TCAAGTCAGCTGGATGAACC
	mOprn-qR	CTTGTCTTGTGGCTGTGAA
23	mOcn-qF	GCAGCACAGGTCTAAATAG
	mOcn-qR	GGGCAATAAGGTAGTGAACAG
24	mLkb1-qF	GTGTGCCAAGCTCATGGGT
	mLkb1-qR	ATGTCCTTGTGAACAATGCC
25	mlhh-qF	CTCAGACCGTGACCGAAATAAG
	mlhh-qR	CCTTGGACTCGTAATACACCCAG
26	mPthrp-qF	CATCAGCTACTGCATGACAAGG
	mPthrp-qR	GGTGGTTTGGTGTGGAG

Full unedited gel for Figure 7A



Full unedited gel for Supplemental Figure 7C



Full unedited gel for Supplemental Figure 7F

