

Table 1S Histomorphometric analysis of 9-wk-old *Mx1-cre;Ctsk^{fl/fl}* mice and their control littermates

Parameters	Male		Female	
	<i>Ctsk^{fl/fl}</i> (n=6)	<i>Mx1;Ctsk^{fl/fl}</i> (n=5)	<i>Ctsk^{fl/fl}</i> (n=4)	<i>Mx1;Ctsk^{fl/fl}</i> (n=5)
BV/TV (%)	9.38±2.28	16.68±5.40*	5.01±2.54	16.42±2.39*
Tb.Th (µm)	36.93±8.14	40.27±6.63	26.10±1.95	39.45±2.24*
Tb.N (/mm)	2.55±0.31	4.06±0.76*	1.88±0.83	4.15±0.42*
Tb.Sp (µm)	361±52	213±54*	588±263	204±29*
MS/BS (%)	30.82±4.24	33.09±3.09	27.74±4.57	30.55±4.74
MAR (µm/day)	2.15±0.29	2.92±0.43*	2.80±0.23	3.72±0.25*
BFR/BS (µm ³ /µm ² /year)	241±42	352±60*	286±64	418±86*
BFR/BV (%/year)	1324±326	1731±288	2095±397	1975±382
BFR/TV (%/year)	139±20	292±87*	116±58	334±66*
Ob.S/BS (%)	12.19±7.64	19.01±7.12	27.52±10.62	26.16±6.77
N.Ob/T.Ar (/mm ²)	49.02±22.90	126.04±45.70*	81.27±33.41	197.55±45.70*
N.Ob/B.Pm (/mm)	10.14±5.79	15.42±5.50	23.48±9.64	24.11±6.14
OV/TV (%)	0.06±0.09	0.25±0.22	0.09±0.10	0.55±0.36*
OS/BS (%)	4.78±6.10	8.71±7.02	6.49±4.45	15.91±8.57
O.Th (µm)	2.71±0.85	3.33±0.47	3.13±0.76	4.08±0.48
Oc.S/BS (%)	1.70±0.93	5.43±2.98*	1.59±0.42	3.69±0.51*
N.Oc/T.Ar (/mm ²)	3.04±1.32	15.05±8.85*	2.06±1.02	10.42±1.05*
N.Oc/B.Pm (/mm)	0.62±0.30	1.80±1.02*	0.55±0.13	1.26±0.12*
ES/BS (%)	0.78±0.42	1.38±0.95	0.46±0.08	0.82±0.34

Results are mean ± SD. **p*<0.05 versus *Ctsk^{fl/fl}* mice, unpaired *t* test

Table 2S Histomorphometric analysis of 9-wk-old male *CD11b;Ctsk^{fl/fl}* mice and their control littermates

Parameters	<i>CD11b;Ctsk^{+/+}</i> (n=7)	<i>CD11b;Ctsk^{fl/fl}</i> (n=7)
BV/TV (%)	12.23±1.85	15.79±3.73*
Tb.Th (µm)	33.29±2.64	39.36±5.21*
Tb.N (/mm)	3.66±0.38	3.99±0.63
Tb.Sp (µm)	242±30	217±45
MS/BS (%)	30.98±4.38	36.86±16.16
MAR (µm/day)	1.46±0.30	2.55±0.74*
BFR/BS (µm ³ /µm ² /year)	169±59	317±104*
BFR/BV (%/year)	924±299	1690±927
BFR/TV (%/year)	132±48	269±90*
Ob.S/BS (%)	5.24±2.45	10.92±6.11*
N.Ob/T.Ar (/mm ²)	31.19±13.81	68.88±42.53*
N.Ob/B.Pm (/mm)	4.30±1.85	8.34±4.39*
OV/TV (%)	0.04±0.04	0.10±0.09
OS/BS (%)	2.23±1.66	4.00±3.46
O.Th (µm)	2.36±0.35	3.07±0.56
Oc.S/BS (%)	0.91±0.65	2.31±1.23*
N.Oc/T.Ar (/mm ²)	2.63±1.62	6.67±3.96*
N.Oc/B.Pm (/mm)	0.36±0.23	0.79±0.41*
ES/BS (%)	0.47±0.34	0.81±0.36

Results are mean ± SD. **p*<0.05 versus *CD11b;Ctsk^{+/+}*, unpaired *t* test

Table 3S Cortical bone μ CT analysis of 9-wk-old male *Mx1;Ctsk^{fl/fl}* mice and their control littermates

Parameters	<i>Ctsk^{fl/fl}</i> (n=5)	<i>Mx1;Ctsk^{fl/fl}</i> (n=5)
Total cross-sectional volume (mm ³)	1.40±0.08	1.55±0.19
Cortical volume (mm ³)	0.56±0.06	0.65±0.10
Marrow volume (mm ³)	0.83±0.07	0.90±0.11
Cortical thickness (mm)	0.17±0.02	0.18±0.01

Results are mean ± SD.

Table 4S Cortical bone μ CT analysis of 9-wk-old male *CD11b;Ctsk^{fl/fl}* mice and their control littermates

Parameters	<i>CD11b;Ctsk^{+/+}</i> (n=6)	<i>CD11b;Ctsk^{fl/fl}</i> (n=6)
Total cross-sectional volume (mm ³)	1.20±0.13	1.40±0.14*
Cortical volume (mm ³)	0.50±0.07	0.65±0.11*
Marrow volume (mm ³)	0.70±0.08	0.75±0.05
Cortical thickness (mm)	0.18±0.02	0.20±0.01

Results are mean ± SD. * $p < 0.05$ versus *CD11b;Ctsk^{+/+}* mice, unpaired *t* test

Table 5S Cortical bone μ CT analysis of 13-week-old male *Osx;Ctsk^{fl/fl}* mice and their control littermates

Parameters	<i>Osx;Ctsk^{+/+}</i> (n=5)	<i>Osx;Ctsk^{fl/fl}</i> (n=5)
Total cross-sectional volume (mm ³)	1.37±0.04	1.27±0.13
Cortical volume (mm ³)	0.61±0.04	0.60±0.05
Marrow volume (mm ³)	0.76±0.03	0.67±0.10
Cortical thickness (mm)	0.19±0.01	0.20±0.01

Results are mean ± SD.

Table 6S Histomorphometric analysis of 13-week-old *Osx;Ctsk^{fl/fl}* mice and their control littermates

Parameters	Male		Female	
	<i>Osx;Ctsk^{+/+}</i> (n=6)	<i>Osx;Ctsk^{fl/fl}</i> (n=7)	<i>Osx;Ctsk^{+/+}</i> (n=4)	<i>Osx;Ctsk^{fl/fl}</i> (n=7)
BV/TV (%)	12.72±5.39	8.85±3.38	4.51±2.30	4.79±2.23
Tb.Th (µm)	33.28±7.01	30.95±5.16	27.84±4.96	30.57±3.88
Tb.N (/mm)	3.73±0.98	2.78±0.67	1.55±0.55	1.55±0.62
Tb.Sp (µm)	252±86	348±99	687±276	687±210
MS/BS (%)	19.86±8.28	27.32±5.97	36.68±4.20	35.77±4.79
MAR (µm/day)	1.38±0.41	1.48±0.22	1.95±0.13	1.97±0.26
BFR/BS (µm ³ /µm ² /year)	107±75	146±29	261±29	257±44
BFR/BV (%/year)	581±395	861±212	1971±400	1676±258
BFR/TV (%/year)	94±80	88±30	83±30	85±44
Ob.S/BS (%)	8.93±5.55	7.60±3.87	19.37±9.13	19.36±8.41
N.Ob/T.Ar (/mm ²)	59.22±44.72	37.56±21.21	51.42±24.58	53.70±35.75
N.Ob/B.Pm (/mm)	7.46±4.25	6.76±3.39	17.76±9.80	17.05±6.89
OV/TV (%)	0.07±0.06	0.05±0.04	0.06±0.06	0.05±0.03
OS/BS (%)	2.70±1.84	2.86±1.84	7.32±5.82	6.55±4.50
O.Th (µm)	3.41±0.41	2.86±0.51	2.90±0.36	2.80±0.60
Oc.S/BS (%)	1.58±1.16	2.04±0.97	1.47±0.65	2.20±1.36
N.Oc/T.Ar (/mm ²)	4.88±4.41	3.63±0.97	1.75±1.17	2.70±2.35
N.Oc/B.Pm (/mm)	0.61±0.43	0.66±0.14	0.54±0.26	0.82±0.45
ES/BS (%)	0.66±0.42	0.71±0.63	0.91±0.64	0.81±0.53

Results are mean ± SD.

Table 7S Sequences of oligonucleotides for qRT-PCR analysis

Gene	Forward primer	Reverse primer
<i>Alp</i>	CTTGACTGTGGTACTGCTGATCA	GTATCCACCGAATGTGAAAACGT
Type I collagen	CCCAAGGAAAAGAAGCACGTC	ACATTAGGCGCAGGAAGGTCA
<i>Dmp1</i>	ATGACTGTCAGGACGGCTAC	AGTTATAGTGAACCTCTCTAC
<i>Runx2</i>	AGTCCCAACTTCCTGTGCTCC	CGGTAACCACAGTCCCATCTG
Osteopontin	CTCCAATCGTCCCTACAGTCG	CCAAGCTATCACCTCGGCC
<i>RANKL</i>	CAAGCTCCGAGCTGGTGAAG	CCTGAACTTTGAAAGCCCCA
<i>OPG</i>	AAGAGCAAACCTTCCAGCTGC	CACGCTGCTTTCACAGAGGTC
<i>Ctsk</i>	AGGCATTGACTCTGAAGATGCT	TCCCCACAGGAATCTCTCTG
<i>Sphk1</i>	TGAGGTGGTGAATGGGCTAATGGA	AACAGCAGTGTGCAGTTGATGAGC
<i>Sphk2</i>	TGGGCTGTCCTTCAACCTCATACA	AGTGACAATGCCTTCCCCTCACT
<i>Efnb2</i>	TCTGTGTCATCGGTTGGCTACGTT	ACAGACGCACAGGACACTTCTCAA
<i>Wnt10b</i>	AGGCTTCTCCTTCCGTTCAAGTTGT	ATTCCCACCCTTCTGCTGAAGAA
<i>Bmp6</i>	AGAAGGGCACTCTTTCAGGTTCCA	TCACACCACCGAGAGTCAACACAA
<i>Gapdh</i>	TGCACCACCAACTGCTTAG	GGATGCAGGGATGATGTTC

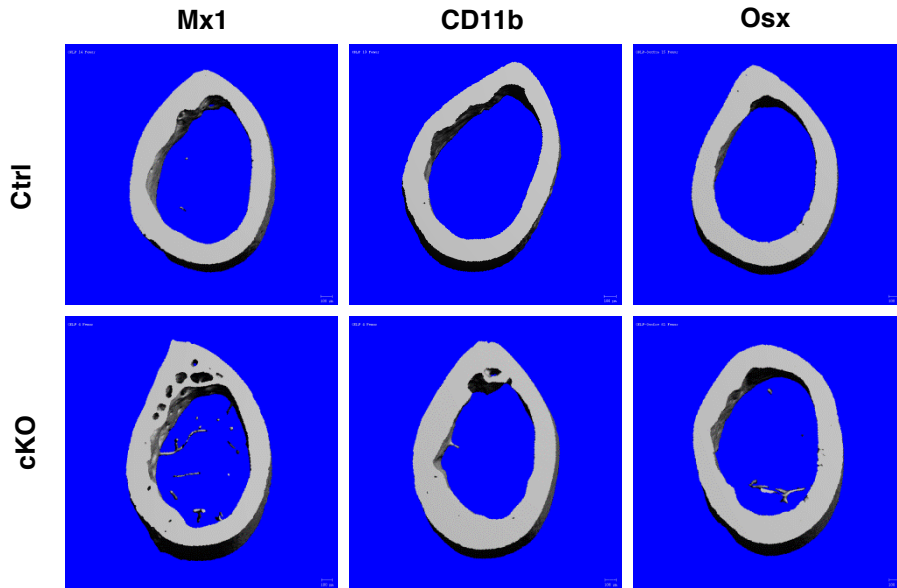


Figure 1S μ CT images of cortical bone in three conditional knockouts (cKO) and their controls (Ctrl). Deletion of *Ctsk* in osteoclasts (CD11b) increases total cross-sectional volume (1.20 ± 0.05 vs 1.40 ± 0.06 , $p < 0.05$) and cortical bone volume (0.50 ± 0.03 vs 0.65 ± 0.05 , $p < 0.05$).

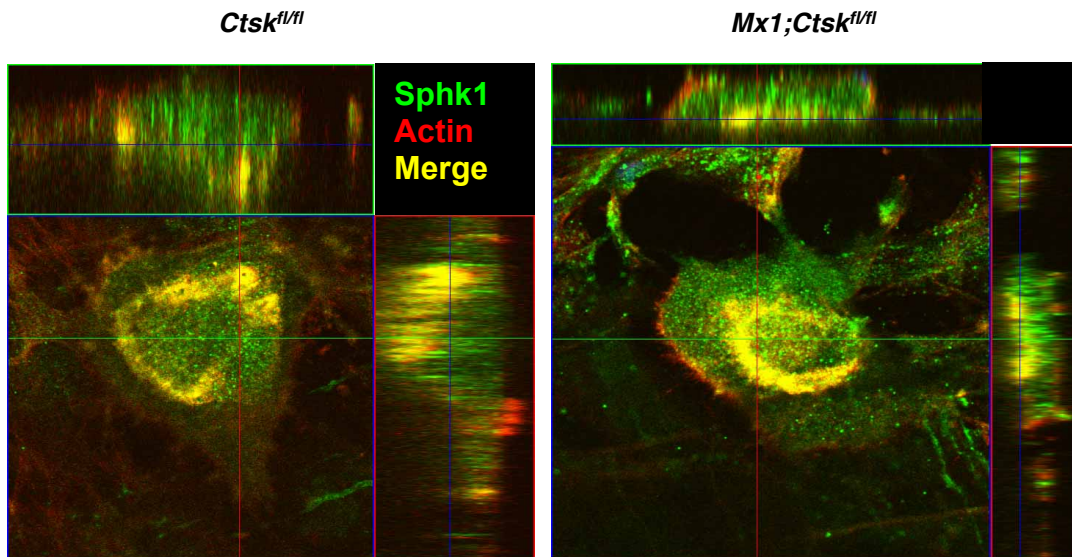


Figure 2S Defective bone resorption in *Mx1;Ctsk^{fl/fl}* osteoclasts. High magnification of Z-axis images of individual podosome showing shallower resorption pit in *Mx1;Ctsk^{fl/fl}* osteoclasts grown on dentin slices. The localization of Sphk1 (green) and actin (red) were analyzed by confocal microscope.

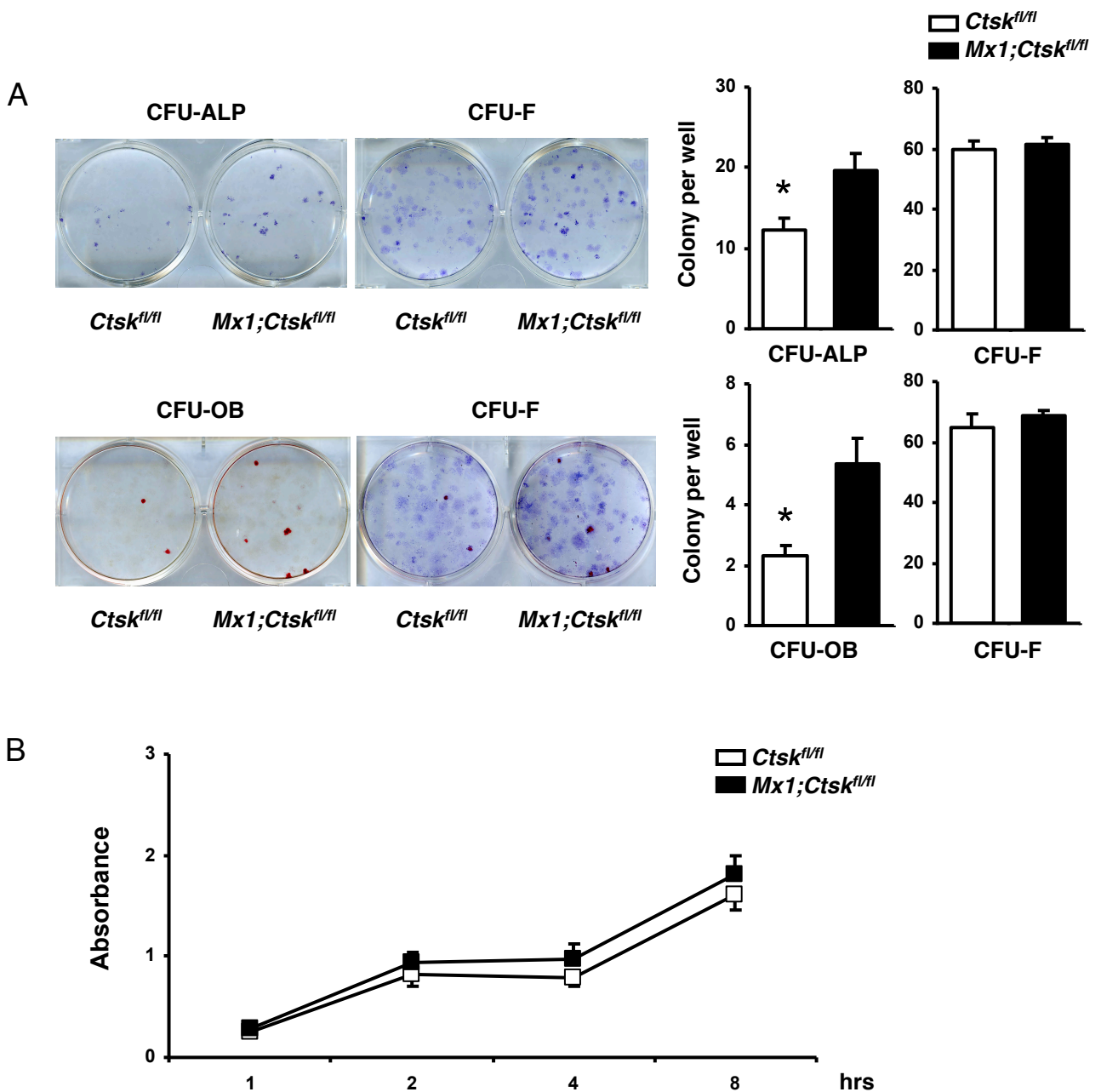


Figure 3S Deletion of *Ctsk* in hematopoietic cells increases osteoblast precursors and differentiation but not proliferation. (A) Cultured osteoblasts derived from long bones of *Mx1;Ctsk*^{fl/fl} mice and control littermates were stained for ALP and alizarin red. *Mx1;Ctsk*^{fl/fl} osteoblasts contain an increased number of CFU-ALP and CFU-OB but not CFU-F. Results are mean \pm SEM. **p* < 0.05 versus controls. (B) Time course of cell proliferation assessed by BrdU labeling showed no changes in proliferation of *Mx1;Ctsk*^{fl/fl} osteoblasts.

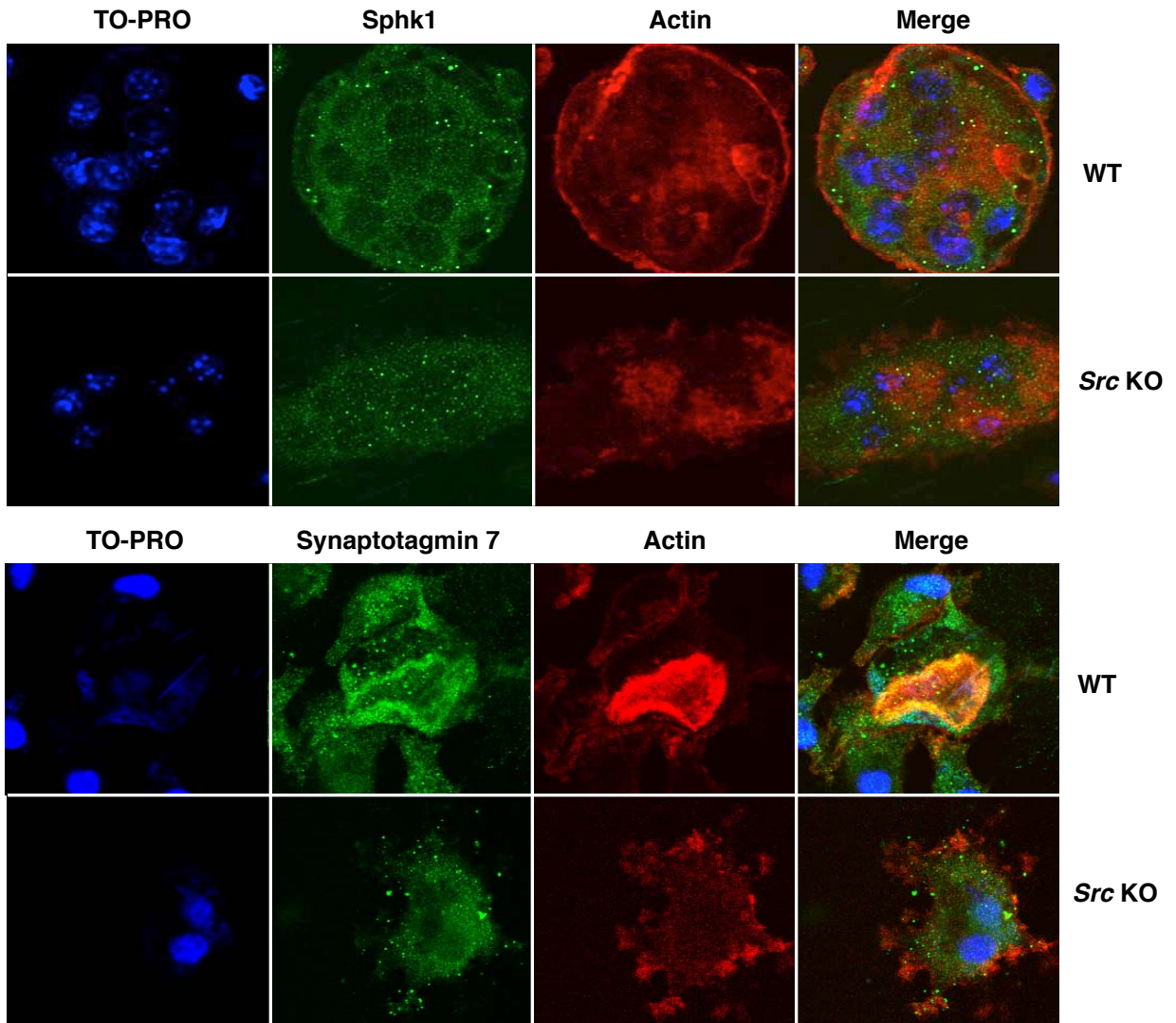


Figure 4S Absence of Src does not alter Sphk1 and Synaptotagmin 7 expression. Osteoclasts were generated by co-culture with osteoblasts on collagen gel in the presence of vitamin D₃ and PGE₂. Cells were replated onto dentin slices and cultured for 48 h and immunolabeled for Sphk1 and synaptotagmin 7 (green) and actin (red). Cells were visualized by scanning confocal immunofluorescence microscopy.