

## **ERK1 and ERK2 play essential roles in osteoblast differentiation and in supporting osteoclastogenesis**

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Abbreviations:

BSP, bone sialoprotein; E, embryonic day; ERK, extracellular signal-regulated kinase;

Ihh, Indian hedgehog; MAPK, mitogen activated protein kinase; MEK, MAPK/ERK; OCN, osteocalcin;

OPG, osteoprotegerin; P, postnatal day; RANKL, Receptor Activator of Nuclear Factor-Kappa B ligand

1 **Abstract**

2 Osteoblasts and chondrocytes arise from common osteo-chondroprogenitor cells. We show here  
3 that inactivation of *ERK1* and *ERK2* in osteo-chondroprogenitor cells causes a block in  
4 osteoblast differentiation and leads to ectopic chondrogenic differentiation in the bone forming  
5 region in the perichondrium. Furthermore, increased MAPK signaling in mesenchymal cells  
6 enhances osteoblast differentiation and inhibits chondrocyte differentiation. These observations  
7 indicate that ERK1 and ERK2 play essential roles in the lineage specification of mesenchymal  
8 cells. The inactivation of ERK1 and ERK2 resulted in reduced beta-catenin expression,  
9 suggesting a role for canonical Wnt signaling in ERK1 and ERK2 regulation of skeletal lineage  
10 specification. Furthermore, inactivation of ERK1 and ERK2 significantly reduced *RANKL*  
11 expression, accounting for a delay in osteoclast formation. Thus, our results indicate that ERK1  
12 and ERK2 not only play essential roles in the lineage specification of osteo-chondroprogenitor  
13 cells, they also support osteoclast formation in vivo.

## 1 **Introduction**

2 The extracellular signal-related kinase/ mitogen-activated protein kinase (ERK MAPK) pathway  
3 is activated by various stimuli including a number of growth factors and cytokines. The  
4 activation of the Raf members of MAPK Kinase Kinase leads to the activation of the MAPK  
5 Kinase, MEK1 and MEK2. MEK1 and MEK2 then phosphorylate and activate MAPK, ERK1  
6 and ERK2. ERK1 and ERK2 phosphorylate various cytoplasmic and nuclear target proteins,  
7 ranging from cytoplasmic adaptor proteins and transcription factors to kinases including RSK  
8 (4,17,18,23,36,46). In this pathway, multiple mutations have recently been identified that cause  
9 syndromes with various skeletal manifestations. Missense activating mutations in KRAS, BRAF,  
10 MEK1, and MEK2 have been identified in Costello, Noonan, LEOPARD, and  
11 Cardio-facio-cutaneous syndromes, while loss-of-function mutations in RSK2, a downstream  
12 kinase of ERK1 and ERK2, cause Coffin-Lowry syndrome (2,42). These observations highlight  
13 the importance of the ERK MAPK pathway in human skeletal development.

14 Both chondrocytes and osteoblasts arise from common osteo-chondro progenitor cells.  
15 Bone growth is achieved through two major ossification processes, endochondral ossification  
16 and intramembranous ossification in which chondrocytes and osteoblasts are involved  
17 (5,13,30,31,38). In normal endochondral ossification, the skeletal element is formed as a

1 cartilaginous template that is subsequently replaced by bone. Condensed mesenchymal cells  
2 differentiate into chondrocytes. Chondrocytes first proliferate in columnar stacks to form the  
3 growth plate and then exit the cell cycle and differentiate into hypertrophic chondrocytes.  
4 Hypertrophic chondrocytes are removed by apoptotic cell death, and the cartilaginous matrix is  
5 resorbed by chondroclasts/osteoclasts and replaced by trabecular bone. Chondroclast/osteoclast  
6 formation is supported by Receptor Activator of Nuclear Factor-Kappa B ligand (RANKL)  
7 secreted from osteoblasts and bone marrow stromal cells (16,45). In intramembranous  
8 ossification, mesenchymal cells directly differentiate into bone-forming osteoblasts; cortical  
9 bone is formed by osteoblasts that arise from the osteo-chondroprogenitor cells in the  
10 perichondrium. Previous studies in mice have indicated that the ERK MAPK signaling cascade  
11 participates in chondrocyte and osteoblast differentiation. Expression of a constitutively active  
12 mutant of MEK1 in chondrocytes caused a dwarf phenotype and inhibited hypertrophic  
13 chondrocyte differentiation (28). In addition, expression of a constitutively active mutant of  
14 MEK1 in mature osteoblasts accelerated bone development, whereas dominant-negative MEK1  
15 was inhibitory at the embryonic stage (11). The roles of ERK/MAPK signaling in mesenchymal  
16 cell differentiation require further investigation. In the current study using the *Cre-loxP* system,  
17 we examined the roles of ERK1 and ERK2 in early steps of chondrocyte and osteoblast

1 differentiation and two major ossification processes, endochondral ossification and  
2 intramembranous ossification by generating loss of function models. In addition to the loss of  
3 function models, we also generated a gain of function model: mice that overexpress a  
4 constitutively active mutant of MEK1 in undifferentiated mesenchymal cells and examined the  
5 roles of ERK/MAPK signaling in osteo-chondro progenitor cells.

6  
7 We show here that ERK1 and ERK2 are essential for osteoblast differentiation, and that  
8 ERK1 and ERK2 inhibit ectopic cartilage formation in the perichondrium. We also show that  
9 increased MAPK signaling in osteo-chondroprogenitor cells inhibits cartilage formation, and  
10 increased MAPK signaling in the perichondrium stimulates bone formation. Furthermore,  
11 inactivation of ERK1 and ERK2 significantly reduced *RANKL* expression, accounting for a delay  
12 in osteoclast formation. Thus, our results indicate that ERK1 and ERK2 not only play essential  
13 roles in the lineage specification of osteo-chondroprogenitor cells, they also support osteoclast  
14 formation *in vivo*.

1 **Materials and methods**

2 **Mutant animals**

3 The institutional animal care and use committee of Case Western Reserve University approved  
4 all animal procedures. All animal care and use were performed in accordance with the  
5 institutional animal use methods and policies. *ERK1*-deficient mice, *Prx1-Cre*, and *Col2a1-Cre*  
6 transgenic mice were described previously (21,32,37). The floxed *ERK2* allele was created by  
7 inserting *loxP* sites flanking exon2 (35). To generate *Prx1-MEK1* transgenic mice, the cDNA for  
8 FLAG-tagged MEK1(S218/222E, Δ32-51) and an *IRES-LacZ* cassette were cloned downstream  
9 of a 2.4 kb *Prx1* promoter (24,28). The construct was microinjected into the pronuclei of  
10 fertilized C57BL/6 x DBA/2 hybrid eggs. *ROSA-LacZ* reporter mice were purchased from  
11 Jackson Laboratories (41).

12

13 **In situ hybridization, immunohistochemistry, TRAP staining, and BrdU incorporation**

14 **assay**

15 In situ hybridization was performed using <sup>35</sup>S-labeled riboprobes. Nuclei were visualized with  
16 Hoechst 33258. Immunohistochemical staining was performed using the PicTure Kit  
17 (Invitrogen) or MM biotinylation kit (Biocare). The following antibodies were used: ERK1/2

1 (SantaCruz), type X collagen (Quartett), FLAG M5 (Sigma), and  $\beta$ -catenin (BD). For  
2 immunofluorescent detection of ERK1/2, Alexa Fluor 594-conjugated secondary antibody  
3 (Invitrogen) was used. Fluorescence-based TRAP staining and alkaline phosphatase staining  
4 were performed using ELF97 (Invitrogen) as a phosphatase substrate (6,10). To reduce  
5 background, TRAP staining using ELF97 was performed in the presence of 50 mM tartrate.  
6 Naphthol AS-BI Phosphoric Acid-based TRAP staining was performed using Acid Phosphatase  
7 Leukocyte kit (Sigma). For examining cell proliferation, mice were injected intraperitoneally  
8 with BrdU. Embryos were harvested 4 h after injection. BrdU incorporation was detected using a  
9 BrdU staining kit (Zymed). All images were acquired with Leica DM 6000B and DM IRB  
10 microscopes. Photographs were taken with a digital camera (DC500; Leica) using Leica  
11 Application Suite 1.3 software. The signal of in situ hybridization in dark field images was  
12 colored using Photoshop software (Adobe), and brightness and contrast were adjusted by  
13 applying brightness/contrast adjustments to the whole image, with the strict intent of not  
14 obscuring, eliminating, or misrepresenting any information present in the original, including  
15 background. For in situ hybridization, the signal images were overlaid on images of Hoechst  
16 33258.  
17

1 **RNA analysis and semi-quantitative and real time PCR**

2 Total RNA was isolated using RNeasy kit (Qiagen). RNA was reverse-transcribed to cDNA with  
 3 High Capacity cDNA Archive Kit (Applied Biosystems). Real-time PCR was performed on the  
 4 Applied Biosystems 7500 Real-time PCR detection system. TaqMan probe sets were designed  
 5 and synthesized by Applied Biosystems (*Erk2*; Mm00442479\_m1, *Rankl*; Mm00441908\_m1,  
 6 *Opg*; Mm00435452\_m1, *Dkk1*; Mm00438422\_m1, *Tcf1*; Mm03053891\_s1, *Cbfb*;  
 7 Mm00491551\_m1, *c-Fos*; Mm00487425\_m1, *Fral*; Mm00487429\_m1, *Fra2*;  
 8 Mm00484442\_m1, *JunB*; Mm00492781\_s1, *Krox20*; Mm00456650\_m1, *Col2a1*;  
 9 Mm01309562\_g1, *Col10a1*; Mm00487041\_m1, *Gapdh*; 4352932E). To compare gene  
 10 expression levels, the comparative cycle threshold (Ct) method was used. *Gapdh* was used as an  
 11 endogenous control to correct for potential variation in RNA loading or in efficiency of  
 12 amplification. Semi-quantitative PCR was performed on the Applied Biosystems GeneAmp PCR  
 13 system 9700 using the following primer sets; *Runx2*, (forward)  
 14 5'-GAACCAAGAAGGCACAGACA-3' and (reverse) 5'-AACTGCCTGGGGTCTGAAAA-3',  
 15 PCR product: 452bp; *Osteocalcin*, (forward) 5'-CTCTGTCTCTCTGACCTCACAG-3' and  
 16 (reverse) 5'-CAGGTCCTAAATAGTGATACCG-3', PCR product: 252bp; *BSP*, (forward)  
 17 5'-GAAACGGTTTCCAGTCCAG-3' and (reverse) 5'-CTGAAACCCGTTTCAGAAGG-3', PCR

1 product: 568bp; *Coll1a2*, (forward) 5'-TGAAGTGGGTCTTCCAGGTC-3' and (reverse)  
 2 5'-GACCAGGCTACCAACAAGT-3', PCR product: 200 bp; *Alkaline phosphatase*, (forward)  
 3 5'-AATGCCCTGAAACTCCAAAAGC-3' and (reverse)  
 4 5'-CCTCTGGTGGCATCTCGTTATC-3', PCR product: 472 bp; *Gapdh*, (forward)  
 5 5'-ACCACAGTCCATGCCATCAC-3' and (reverse) 5'-TCCACCACCCTGTTGCTGTA-3',  
 6 PCR product: 452 bp . The band intensity was compared while the band intensity and cycle  
 7 numbers are linear.

8

9 **Cell culture**

10 Primary rib chondrocytes and calvaria osteoblasts were isolated as described previously (12,28).  
 11 Cells were infected with Adenovirus expressing Cre, GFP (Gene Transfer Vector Core,  
 12 University of Iowa), or constitutively active MEK1 (Vector Biolabs) at 150-200 MOI.  
 13 Osteoblasts were grown in differentiation medium (alpha MEM, 10% FCS, 5 mM  
 14  $\beta$ -glycerophosphate, 100  $\mu$ g/ml ascorbic acid). Alizarin red and von Kossa staining were done  
 15 following standard protocols. For in vitro osteoclast differentiation, spleen cells were isolated  
 16 and cultured in alpha MEM supplemented with 10% FCS, 10 ng/ml RANKL (R&D), and L929  
 17 cell-conditioned medium containing M-CSF.

1

2 **Western blot analysis**

3 Total cellular protein was prepared by lysing cells in 62.5mM TrisHCl pH 6.8, 2% SDS, 10%  
4 glycerol, 1 mM sodium orthovanadate, 1 mM sodium fluoride, 1 mM  $\beta$ -glycerophosphate,  
5 supplemented with proteinase inhibitor cocktail Complete Mini (Roche). Protein concentrations  
6 were determined by BCA Protein Assay kit (Pierce). Twenty to thirty micrograms of protein was  
7 separated by 10% SDS/PAGE and electrophoretically transferred to PVDF filters (Millipore).  
8 The filters were blocked in 5% nonfat dry milk in Tris-buffered saline (pH 7.5) containing 0.1%  
9 Tween 20 and then incubated with the following antibodies: ERK1 and ERK2 (SantaCruz),  
10 Osterix (SantaCruz), ATF4 (SantaCruz), RSK2 (Cell Signaling Technology), and  $\beta$ -actin (Cell  
11 Signaling Technology). Filters were then incubated with the second antibody (horseradish  
12 peroxidase-conjugated anti-rabbit or anti-goat IgG), and the signal was detected by enhanced  
13 chemiluminescence (Pierce). Images were captured using KODAK Image Station 4000MM.

1 **Results**

2 **Inactivation of *ERK2* in mesenchymal cells of *ERK1*-null mice cause severe limb deformity**  
3 **and bone defects**

4 We first analyzed *ERK1*-null mice and *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice, in which *ERK2* was  
5 inactivated in the limb and head mesenchyme using the *Prx1-Cre* transgene. These mice did not  
6 show obvious skeletal abnormalities (data not shown). To totally inactivate *ERK1* and *ERK2*, we  
7 further inactivated *ERK2* in the *ERK1*-null background. *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice  
8 were born at the expected Mendelian ratio. Skeletal preparation of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;  
9 *Prx1-Cre* mice using alcian blue and alizarin red staining showed severe limb deformity at  
10 postnatal day 0 (P0) (Fig. 1A, Supplemental Fig. 1). In addition, *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre*  
11 mice showed bone defects in the calvaria. The lambdoid suture did not close at least up to P12  
12 (Fig. 1B). Histological examination showed no distinct cortical bone formation in the humerus of  
13 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice at P5 (Fig. 1C). *ERK2* inactivation was confirmed in the  
14 tibia, femur and humerus of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* embryos at embryonic day (E)16.5  
15 by real-time PCR (Fig. 1D, and data not shown). *ERK2* mRNA was decreased about 90%  
16 compared with littermate *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup> embryos. The inactivation of ERK1 and ERK2  
17 was also confirmed by immunohistochemistry using an antibody that recognizes both ERK1 and

1 ERK2 (Fig. 1E). Immunoreactivity of chondrocytes and cells in the bone forming area was  
2 significantly reduced in *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* mice. These observations indicate that  
3 ERK1 and ERK2 are essential for bone formation.

4

#### 5 **ERK1 and ERK2 are essential for osteoblast differentiation**

6 To examine osteoblast differentiation, we performed in situ hybridization analyses. Early  
7 osteoblast marker *Coll1a1* and master transcription factors for osteoblast differentiation, *Runx2*,  
8 *Osterix*, and *Atf4* were normally expressed in the long bones and calvaria of *ERK1*<sup>-/-</sup>;  
9 *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* mice, indicating ERK1 and ERK2 are not required for their expression  
10 (Fig. 2A,C, and data not shown). However, the expression of *Osteocalcin*, a marker of mature  
11 osteoblasts, was undetectable in the long bones and calvaria of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre*  
12 mice (Fig. 2B,C). Because *Osterix* regulates *Osteocalcin* expression downstream of *Runx2*, these  
13 observations indicate that osteoblast differentiation was arrested after *Osterix* expression and  
14 before differentiation into fully mature osteoblasts.

15

16 To examine whether the block in osteoblast differentiation is due to the cell autonomous  
17 effects of *ERK1* and *ERK2* inactivation, we isolated calvaria osteoblasts from *ERK1*<sup>-/-</sup>; *ERK2*

1 <sup>flox/flox</sup> embryos. The cells were infected with adenovirus expressing Cre recombinase (Ad-Cre) or  
2 a control virus expressing GFP (Ad-GFP). RT-PCR showed marked reduction of *Osteocalcin* and  
3 *Bsp* expression in cells infected with Ad-Cre, while *Alkaline phosphatase* expression was slightly  
4 reduced and *Colla2* expression remained unaltered (Fig. 3A). These observations indicate that  
5 the block in osteoblast differentiation is cell autonomous, and ERK1 and ERK2 are required for  
6 osteoblast differentiation into mature osteoblasts. Consistent with in vivo observations, the  
7 inactivation of *ERK1* and *ERK2* did not abolish *Runx2* mRNA. In addition, *ERK1* and *ERK2*  
8 inactivation did not affect Osterix, ATF4, and RSK2 protein expression (Fig. 3B). The block in  
9 osteoblast differentiation was also confirmed by von Kossa staining (Fig. 3C).

10

11 We also isolated the tibia, femur, and humerus of E16.5 embryos and examined  
12 expression of transcription factors implicated in osteoblast differentiation by real time PCR. We  
13 found that a number of transcription factors were downregulated in the skeletal elements of  
14 *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Prx1-Cre* embryos. *Krox20* was strongly inhibited in bones lacking ERK1  
15 and ERK2 (Fig. 3D). In addition, AP-1 family members *Fra1*, *Fra2*, and *c-Fos* were all  
16 downregulated in bones lacking ERK1 and ERK2. In contrast, *JunB* expression remained  
17 unchanged, indicating that the downregulation of *Krox20*, *Fra1*, *Fra2*, and *c-Fos* is not a

1 consequence of general suppression of transcription. Furthermore, *Cbfb*, which encodes a  
2 co-regulator of runt-domain transcriptional factors, was also downregulated in bones lacking  
3 ERK1 and ERK2 (27,47). Therefore, severe bone phenotype of *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Prx1-Cre*  
4 mice likely involves multiple regulatory factors of osteoblast differentiation. Consistent with  
5 these observations, at least *Krox20* and *c-Fos* were strongly inhibited in calvaria cells infected  
6 with Ad-Cre (Fig. 3E).

7

### 8 **Inactivation of ERK1 and ERK2 in mesenchymal cells cause ectopic cartilage formation in** 9 **the perichondrium**

10 Interestingly, we observed ectopic cartilage formation in the perichondrium of *ERK1*<sup>-/-</sup>;  
11 *ERK2*<sup>lox/lox</sup>; *Prx1-Cre* mice (Fig. 4A). The ectopic cartilage formation was observed as early as  
12 E13.5 in the mutant humerus (Supplemental Fig. 2B). Cells in the ectopic cartilage expressed a  
13 master transcription factor for chondrocyte differentiation, *Sox9*, and a cartilage-specific marker,  
14 *Col2a1*, indicating chondrogenic differentiation. These observations suggest that  
15 osteo-chondroprogenitor cells in the perichondrium were blocked in their differentiation into  
16 osteoblasts, and instead differentiated into chondrocytes. Interestingly, the cells in the ectopic  
17 cartilage also expressed markers for prehypertrophic chondrocytes, *Indian hedgehog* (*Ihh*), and

1 *Parathyroid hormone (PTH)/PTH-related peptide receptor* as well as markers for hypertrophic  
2 chondrocytes, *Col10a1* and *Mmp13* (Fig. 4A, Supplemental Fig. 2G,H, and data not shown). It is  
3 possible that chondrocytes in the ectopic cartilage accelerated differentiation into hypertrophic  
4 chondrocytes in the absence of ERK1 and ERK2.

5

6         Since the inactivation of  $\beta$ -catenin in mesenchymal cells causes similar ectopic cartilage  
7 formation in the perichondrium (7,12,34), we examined  $\beta$ -catenin expression by  
8 immunohistochemistry. Strong  $\beta$ -catenin expression was observed in the bone-forming region of  
9 the periosteum and perichondrium of control embryos at E15.5 (Fig. 4B). In contrast,  $\beta$ -catenin  
10 staining was decreased in the periosteum and perichondrium of *ERK1<sup>-/-</sup>; ERK2<sup>fl/fl</sup>; Prx1-Cre*  
11 embryos. We further examined *Tcf1* and *Dkk1* expression by real time PCR, because both *Tcf1*  
12 and *Dkk1* expression depends on intact  $\beta$ -catenin signaling (12,34). We observed strong  
13 inhibition of *Tcf1* and *Dkk1* expression, supporting the notion that  $\beta$ -catenin signaling is reduced  
14 in the skeletal elements (Fig. 4C).

15

16         Ectopic chondrocyte differentiation in the perichondrium in association with a block in  
17 osteoblast differentiation has been also reported in *Osterix*-null mice and in mice in which *Ihh*

1 signaling is disrupted (22,29). *Osterix*, *Ihh* and its downstream target *Patched* were normally  
2 expressed in the skeletal elements of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fl/fl</sup>; *Prx1-Cre* embryos (Fig. 2A, 4A,  
3 Supplemental Fig. 2G). In addition, coexpression of a constitutively active mutant of MEK1 did  
4 not affect the transcriptional activity of *Osterix* in transient transfection experiments *in vitro*  
5 (data not shown). These observations suggest that *Osterix* and *Ihh* signaling are not affected by  
6 MAPK signaling.

7

8 **Inactivation of ERK1 and ERK2 caused an expansion of terminally differentiated**  
9 **chondrocytes in the growth plates and decreased osteoclasts**

10 The inactivation of ERK1 and ERK2 resulted in a remarkable widening of the zone of  
11 hypertrophic chondrocytes in the growth plate (Fig. 5A). Some of the epiphyseal chondrocytes  
12 closer to the articular surface stained positive for type X collagen, suggesting accelerated  
13 hypertrophy. This notion is further supported by an increase in *Col10a1* expression in *ERK1*<sup>-/-</sup>;  
14 *ERK2*<sup>fl/fl</sup> primary chondrocytes that were infected with Ad-Cre (Fig 7C). In addition, *in situ*  
15 hybridization indicated that the widening of the zone of hypertrophic chondrocytes is associated  
16 with a remarkable expansion of terminally differentiated hypertrophic chondrocytes (Fig. 5B).  
17 Although the matrix showed intense staining with anti-type X collagen antibody, these cells

1 expressed *Col10a1* at a reduced level and instead expressed markers of terminally differentiated  
2 hypertrophic chondrocytes, *Vegf*, *Mmp13*, and *Osteopontin*. These observations suggest that the  
3 expansion of the hypertrophic zone is caused by impaired removal of terminally differentiated  
4 hypertrophic chondrocytes. TUNEL staining did not show significant difference between control  
5 and mutant mice, suggesting chondrocyte apoptosis is not affected (data not shown).

6

#### 7 **Inactivation of ERK1 and ERK2 caused a decrease in osteoclast formation and RANKL** 8 **expression**

9 We also examined osteoclasts by TRAP staining. TRAP-positive cells were decreased in the long  
10 bones of *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* embryos at E16.5 (Fig. 5C), suggesting that the  
11 decreased osteoclasts accounts at least in part for the delayed cartilage removal in *ERK1<sup>-/-</sup>;*  
12 *ERK2<sup>fllox/fllox</sup>; Prx1-Cre* embryos. Consistent with the reduced number of osteoclasts,  
13 immunohistochemistry for MMP9 showed reduced staining in *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre*  
14 embryos (Supplemental Fig. 3A).

15

16 Since osteoclastogenesis is regulated by Receptor Activator of Nuclear Factor-Kappa B ligand  
17 (RANKL) and osteoprotegerin (OPG) produced by mesenchymal cells, we examined *RANKL*

1 and *OPG* expression in the long bones of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* embryos. *RANKL*  
2 expression was strongly inhibited in the long bones of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* embryos,  
3 while *OPG* expression remained unaltered (Fig. 5D, and data not shown). Because the *Prx1-Cre*  
4 transgene inactivates *ERK2* both in chondrocytes and osteoblasts, we examined *RANKL* and  
5 *OPG* regulation in each cell lineage in vitro. We harvested calvarial osteoblasts and rib  
6 chondrocytes from *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> mice and inactivated *ERK2* by infecting adenovirus  
7 expressing Cre recombinase. In both cell types, *ERK2* inactivation strongly inhibited *RANKL*  
8 expression, while *OPG* expression was not affected (Fig. 5E,F). Consistent with these  
9 observations, treatment with MEK inhibitor U0126 strongly inhibited *RANKL* expression in both  
10 cell types (Supplemental Fig. 3B,C). These observations indicate that *RANKL* expression  
11 depends on MAPK signaling both in the osteoblast and chondrocyte lineages.  
12  
13 Since reduced osteoclastogenesis could be due to unspecific deletion of *ERK1* and *ERK2* in the  
14 osteoclast lineage, we examined ERK expression in osteoclasts of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>;  
15 *Prx1-Cre* mice by double staining for TRAP activity and ERK protein. Fluorescence-based  
16 TRAP staining using ELF97 showed green fluorescence in osteoclasts of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>;  
17 *Prx1-Cre* at P0 (Fig. 5G). The staining intensity for ERK is similar to that of osteoclasts of

1 littermate *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> mice (data not shown). To check the activity of the *Prx1-Cre*  
2 transgene in the monocyte/macrophage lineage, we further isolated spleen cells from *ROSA-LacZ*  
3 reporter mice harboring the *Prx1-Cre* transgene and control *ROSA-LacZ* reporter mice without  
4 the *Prx1-Cre* transgene. These cells were induced to differentiate into osteoclast-like cells in the  
5 presence of M-CSF and RANKL. X-gal staining followed by TRAP staining showed no X-gal  
6 staining in TRAP-positive osteoclast-like cells, indicating that the *Prx1-Cre* transgene is not  
7 active during in vitro osteoclast differentiation (Fig. 5H).

8  
9 To further exclude the possibility that the reduced osteoclast formation in *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>;  
10 *Prx1-Cre* mice is due to defects in osteoclast precursor cells, we isolated spleen cells from  
11 *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* and littermate *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> mice at P0. Spleen cells from  
12 *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* formed osteoclast-like multinucleated giant cells in the presence  
13 of M-CSF and RANKL similar to *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> cells (Fig. 5I and data not shown). These  
14 giant cells displayed TRAP activity and expressed ERK similar to *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> cells.  
15 These observations further support the notion that reduced osteoclastogenesis in *ERK1*<sup>-/-</sup>;  
16 *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* embryos is due to reduced support from mesenchymal cells.

17

1 **ERK1 and ERK2 inactivation in chondrocytes causes severe chondrodysplasia**

2 To further examine the roles of ERK1 and ERK2 in differentiated chondrocytes, we inactivated  
3 *ERK2* in chondrocytes of *ERK1*-null mice using the *Col2a1-Cre* transgene. *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;  
4 *Col2a1-Cre* mice died immediately after birth, presumably due to respiratory insufficiency  
5 caused by the defective rib cage. We obtained *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* mice at a  
6 Mendelian ratio at E18.5. The inactivation of ERK1 and ERK2 in chondrocytes was confirmed  
7 by immunofluorescence using the antibody that recognizes both ERK1 and ERK2 (Fig. 6A). We  
8 further confirmed ERK expression in osteoblasts and osteoclasts of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;  
9 *Col2a1-Cre* embryos by performing ELF97-based fluorescent alkaline phosphatase staining and  
10 TRAP staining along with immunofluorescence for ERK (Fig. 6B,C). The fluorescent signal for  
11 ERK was indistinguishable from that of control *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup> embryos (data not shown).  
12 Consistent with the normal expression of ERK in osteoblasts, calvaria osteoblasts isolated from  
13 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos showed mineralization similar to *ERK1*<sup>+/-</sup>;  
14 *ERK2*<sup>flox/flox</sup> cells in vitro (Supplemental Fig 4).  
15  
16 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* mice showed a strong cartilage phenotype both in the axial and  
17 appendicular skeletons. Skeletal preparation with alcian blue and alizarin red staining showed

1 kyphotic deformity in the spine of *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Col2a1-Cre* embryos (Fig. 6D).  
2 Histological examinations showed an absence of primary ossification centers in the spine and  
3 cranial base at E18.5 (Fig. 6E,F). In long bones, there was a gene dosage-dependent widening of  
4 the zone of hypertrophic chondrocytes (Fig. 7A). *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Col2a1-Cre* embryos  
5 showed an expansion of *Col10a1* expression domain (Fig. 7B). Similar to *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>*;  
6 *Prx1-Cre* embryos, *Col10a1* expression was also observed in chondrocytes that were closer to  
7 the articular surface in *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Col2a1-Cre* embryos, suggesting premature  
8 hypertrophy. To further confirm the role of ERK1 and ERK2 in hypertrophic differentiation of  
9 chondrocytes, we performed *in vitro* experiments. *ERK2* inactivation in primary *ERK1<sup>-/-</sup>*;  
10 *ERK2<sup>flox/flox</sup>* chondrocytes by adenovirus-mediated expression of Cre recombinase resulted in  
11 increased *Col10a1* expression, further supporting the notion that ERK1 and ERK2 inhibit  
12 hypertrophic differentiation of chondrocytes (Fig. 7C). Loss of *ERK1* and *ERK2* also resulted in  
13 severe disorganization of the epiphyseal cartilage. Chondrocytes failed to form columnar  
14 structures in the growth plates (Fig. 7D). BrdU incorporation experiments indicated significant  
15 reduction in chondrocyte proliferation in *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Col2a1-Cre* embryos at E18.5  
16 (Fig. 7E). These observations indicate that ERK1 and ERK2 are required for proper formation of  
17 columnar structures and chondrocyte proliferation.

1  
 2 Similar to *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* embryos, TRAP-positive osteoclasts were decreased in  
 3 *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Col2a1-Cre* embryos at E16.5, suggesting chondrocytes support  
 4 osteoclastogenesis through ERK1 and ERK2 at this stage (Fig. 8A). The reduced  
 5 osteoclastogenesis is unlikely to be due to unspecific inactivation of ERK in the osteoclast  
 6 lineage, since TRAP-positive osteoclasts of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Col2a1-Cre* embryos express  
 7 ERK (Fig. 6B). In addition, when osteoclast-like cells were generated in vitro from spleen cells  
 8 of *Col2a1-Cre* transgenic mice harboring the *ROSA-LacZ* reporter allele, TRAP -positive  
 9 osteoclast-like cells did not stain positive for X-gal (data not shown). Furthermore, spleen cells  
 10 isolated from E18.5 *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Col2a1-Cre* embryos formed TRAP-positive  
 11 osteoclast-like cells in the presence of M-CSF and RANKL, and these cells express ERK similar  
 12 to *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> cells (Fig. 8B).

13  
 14 **Increased bone formation in mice that express constitutively active mutant of**  
 15 ***MEK1(S218/222E, Δ32-51)* under the control of a 2.4 kb *prx1* promoter**

16 To further examine the roles of MAPK in mesenchymal cells, we generated *Prx1-MEK1*  
 17 transgenic mice that express a constitutively active mutant of *MEK1(S218/222E, Δ32-51)* and

1 *LacZ* under the control of a 2.4 kb *prx1* promoter (Fig. 9A). X-gal staining showed the transgene  
2 expression in the developing limb bud, the perichondrium and periosteum of the long bones,  
3 some of the periarticular chondrocytes, and the lambdoid suture in the cranium (Fig. 9B,C,F).  
4 Protein expression of the FLAG-tagged MEK1 mutant was also confirmed by  
5 immunohistochemistry using anti-FLAG antibody (Fig. 9D). In remarkable contrast to the  
6 defective bone formation in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice, *Prx1-MEK1* transgenic mice  
7 showed a dramatic increase in cortical bone formation, fusion of long bones (Fig.10A) and carpal  
8 and tarsal bones (Fig. 9E), and accelerated closure of the lambdoid suture (Fig. 9F). We observed  
9 relatively normal expression of *Gdf5* and *Wnt 14* at the presumptive joint region (data not  
10 shown), suggesting that the bone fusions are not due to the altered joint formation, but due to  
11 increased bone formation. The increase in cortical bone thickness was preceded by a thickening  
12 of *Runx2*-, *Osterix*-, and *Bsp*-expressing perichondrium, suggesting that MAPK signaling  
13 recruits and directs osteo-chondral progenitor cells toward the osteoblastic lineage (Fig. 10B).  
14 We also observed accelerated osteocalcin expression in the perichondrium of *Prx1-MEK1*  
15 transgenic mice compared with wild type mice (Supplemental Fig 5A), suggesting an accelerated  
16 osteoblast differentiation by the increased MAPK signaling.  
17

1 **Inhibition of cartilage formation in Prx1-MEK1 transgenic mice**

2 In contrast to the increased bone formation, we observed inhibition of cartilage formation in the  
3 transgenic mice. Skeletal preparation of E14.5 embryos showed a delay in the formation of  
4 cartilage anlagen, and alcian blue staining of histological sections showed smaller cartilage  
5 anlagen in the transgenic mice at E15.5 (Fig. 10C and Supplemental Fig. 5C).

6 Immunohistochemistry for the constitutively active MEK1 indicated that cartilage develops  
7 within the transgene-expressing mesenchymal condensation (Supplemental Fig. 5B). In addition,  
8 alcian blue staining was reduced in the periarticular cartilage corresponding to the expression  
9 domain of the constitutively active MEK1 (Fig. 9D, arrows). Furthermore, *Col2a1* expression  
10 was decreased in the developing cartilage primordia (Fig. 10D). Consistent with these  
11 observations *in vivo*, adenovirus-mediated expression of a constitutively active mutant of MEK1  
12 in primary chondrocytes strongly inhibited *Col2a1* expression (Fig. 10E). In addition, FGF18, a  
13 potent activator of the MAPK pathway, also downregulated *Col2a1* expression, and the  
14 downregulation was inhibited by U0126 (Fig. 10F). Collectively, these observations indicate that  
15 increased MAPK signaling inhibits cartilage formation.

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## 1 Discussion

2 In this study, we showed *ERK1* and *ERK2* are essential for osteoblast differentiation and bone  
3 formation. *Osteocalcin*-expressing mature osteoblasts did not develop in the absence of ERK1  
4 and ERK2, indicating ERK1 and ERK2 are essential for differentiation into mature osteoblasts.  
5 Despite severe impairment in osteoblast differentiation, master transcription factors for  
6 osteoblast differentiation, *Runx2*, *Osterix* and *Atf4* are normally expressed, indicating ERK1 and  
7 ERK2 are not required for their expression. In addition to transcriptional regulation, ERK1 and  
8 ERK2 can regulate the activity of transcription factors posttranslationally. Indeed, Runx2 has  
9 been shown to be phosphorylated and activated by ERK MAPK (11,43). In addition, Atf4 is  
10 phosphorylated and activated by RSK2, which is a cytoplasmic substrate of ERK1 and ERK2  
11 (44). Phosphorylation by ERK1 and ERK2 is essential for the complete activation of RSK2  
12 (40,50). Therefore, reduced activity of Runx2 and Atf4 may have a role in the severe defects in  
13 osteoblast differentiation. However, downstream targets of Runx2, such as *Osterix*, *Atf4*, *Vegf*,  
14 *Ihh*, and *Col10a1*, are normally expressed in the absence of ERK1 and ERK2, suggesting Runx2  
15 is not totally inactive in the absence of ERK1 and ERK2 (29,44,48,49,51). In addition, the bone  
16 phenotype of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* mice is apparently more severe than that of *Atf4*  
17 and *Rsk2*-null mice, suggesting additional mechanisms for regulating osteoblast differentiation.

1

2 Consistent with this notion, we found that a number of transcription factors implicated  
3 in bone formation were downregulated in *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* embryos. Notably,  
4 *Krox20*, a zinc finger transcription factor expressed in endosteal and periosteal osteoblasts, was  
5 strongly downregulated in the absence of ERK1 and ERK2. *Krox20* enhances the *Osteocalcin*  
6 promoter activity (19), and *Krox20*-null mice show a strong decrease in trabecular bone  
7 formation and Osteocalcin-positive cells (20). In addition, AP-1 family members, *Fra1*, *Fra2*,  
8 *c-Fos*, and *Cbfb*, which encodes a co-regulator of runt-domain transcriptional factors, were all  
9 downregulated in the skeletal elements lacking ERK1 and ERK2. Since these transcriptional  
10 regulators are all implicated in bone formation (1,9,15,26,27,47), these observations strongly  
11 suggest that ERK1 and ERK2 regulate osteoblast differentiation and bone formation through  
12 multiple regulators.

13

14 In addition to the block in osteoblast differentiation, we observed ectopic cartilage  
15 formation in the perichondrium of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* mice (Fig. 4A, Supplemental  
16 Fig. 2). These observations suggest that osteochondral progenitor cells were blocked in their  
17 differentiation into osteoblasts and differentiated into chondrocytes. Similar phenotype has been

1 reported in *β-catenin* conditional knock out mice (7,12). We observed decreased  $\beta$ -catenin  
2 protein levels in the perichondrium of *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>; Prx1-Cre* mice. Consistent with this  
3 observation, expression of *Tcf1* and *Dkk1*, which is dependent on  $\beta$ -catenin signaling, was  
4 strongly inhibited in these mice. These results suggest a role for reduced  $\beta$ -catenin signaling in  
5 the ectopic cartilage formation in the perichondrium. Loss of ERK1 and ERK2 may result in low  
6  $\beta$ -catenin protein levels through intracellular crosstalk between ERK and canonical Wnt  
7 signaling. Such interaction has been reported in hepatocellular carcinoma, in which ERK  
8 inactivates GSK-3 $\beta$  leading to the stabilization of  $\beta$ -catenin (8). Alternatively, the effect of ERK  
9 inactivation on  $\beta$ -catenin expression may be indirect, and loss of ERK1 and ERK2 results in a  
10 block in osteoblast differentiation before differentiating osteoblasts express  $\beta$ -catenin at an  
11 increased level. The inactivation of  $\beta$ -catenin in calvarial mesenchymal cells using the *Prx1-Cre*  
12 or *Derm1-Cre* transgene caused chondrogenic differentiation (7,12). In contrast, loss of ERK1  
13 and ERK2 did not cause ectopic cartilage formation in the calvariae, indicating distinct roles in  
14 the lineage specification of cranial mesenchyme. Further investigation is necessary to elucidate  
15 the interaction between MAPK and  $\beta$ -catenin in the lineage specification of mesenchymal cells.

16

17 In remarkable contrast to the defective bone formation in *ERK1<sup>-/-</sup>; ERK2<sup>flox/flox</sup>*,

1 *Prx1-Cre* mice, *Prx1-MEK1* transgenic mice that express a constitutively active mutant of MEK1  
2 in the perichondrium/periosteum showed a dramatic increase in cortical bone formation. The  
3 increase in cortical bone thickness was preceded by a thickening of *Runx2*-, *Osterix*-, and  
4 *Bsp*-expressing perichondrium, suggesting that MAPK signaling recruits and directs  
5 osteo-chondral progenitor cells toward the osteoblastic lineage. In addition, *Prx1-MEK1*  
6 transgenic mice showed a delay in the formation of cartilage anlage. This is consistent with the  
7 notion that MAPK signaling inhibits chondrogenic differentiation of mesenchymal cells.  
8 Interestingly, the accelerated cranial suture closure and bone fusions in *Prx1-MEK1* transgenic  
9 mice were similar to those of human skeletal syndromes caused by activating mutations in  
10 Fibroblast growth factors receptor 2 (FGFR2) (14). These include Apert, Pfeiffer, Jackson-Weiss,  
11 and Crouzon syndromes. These syndromes show craniosynostosis and variable degrees of limb  
12 abnormalities, including cutaneous and osseous syndactyly and fusion of various bones. Since  
13 the MAPK pathway is one of the major downstream pathways of FGF signaling, these  
14 observations suggest that the MAPK pathway is responsible for some of the clinical features of  
15 FGFR2-related skeletal syndromes. This notion is further supported by the recent observation in  
16 mice that express an Apert syndrome mutation S252W, in which MEK inhibitor U0126 treatment  
17 rescued craniosynostosis (39).

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In contrast to osteoblast differentiation, loss of ERK1 and ERK2 did not affect the formation of cartilage anlage, indicating that ERK1 and ERK2 are dispensable for cartilage formation. However, loss of ERK1 and ERK2 caused severe disorganization of the epiphyseal cartilage and significant reduction in chondrocyte proliferation. These observations indicate that ERK1 and ERK2 are required for the proper organization of the epiphyseal cartilage and chondrocyte proliferation. The cartilage phenotype of *ERK1*, *ERK2* conditional knock out mice was substantially alleviated by one functional allele of either ERK1 or ERK2, indicating one allele of either ERK1 or ERK2 is sufficient for restoring the growth plate architecture and chondrocyte proliferation. We have previously shown that chondrocyte proliferation is not affected in mice that express a constitutively active mutant of MEK1 in chondrocytes (28). Apparently, basal MAPK signaling is sufficient for chondrocyte proliferation, and further increasing MAPK signaling does not stimulate proliferation. Recently, *B-raf* has been conditionally inactivated in chondrocytes in the *A-raf*-null background (33). Surprisingly, these mice did not show obvious cartilage abnormalities. Our results support the notion that intact *C-raf* signaling activates ERK1 and ERK2 to a level sufficient for normal cartilage development.

1           In both *ERK1*<sup>-/-</sup>; *ERK2*<sup>flx/flx</sup>; *Prx1-Cre* mice and *ERK1*<sup>-/-</sup>; *ERK2*<sup>flx/flx</sup>; *Co2a1-Cre*  
2 embryos, we observed premature *Col10a1* expression in chondrocytes in the epiphysis,  
3 suggesting accelerated hypertrophic differentiation. Interestingly, *Col10a1* positive chondrocytes  
4 were observed mainly at the lateral edges of the epiphysis in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flx/flx</sup>; *Prx1-Cre*  
5 embryos, while chondrocytes in the center expressed *Col10a1* at lower levels. This might be  
6 related to the timing of ERK inactivation using the *Prx1-Cre* transgene. The inactivation of  
7 ERK1 and ERK2 upregulated *Col10a1* expression in chondrocytes in vitro, further supporting  
8 the notion that ERK1 and ERK2 inhibits hypertrophic differentiation. These observations are  
9 consistent with our previous observations that increased MEK1 signaling in chondrocytes  
10 inhibits hypertrophic differentiation in transgenic mice (28).

11  
12           Another important function of ERK1 and ERK2 revealed in this study is the role of  
13 ERK1 and ERK2 in supporting osteoclastogenesis. Osteoclast formation is stimulated by  
14 RANKL that is secreted from osteoblasts, osteoblast precursor cells as well as bone marrow  
15 stromal cells, and its action is counterbalanced by its decoy receptor OPG (3,16,45). Our in vitro  
16 analyses indicated that ERK1 and ERK2 are required for *RANKL* expression both in the  
17 osteoblast and chondrocyte lineages. Consistent with this observation, loss of ERK1 and ERK2

1 in the developing skeleton caused a strong decrease in *RANKL* expression and reduced formation  
2 of osteoclasts. Most interestingly, chondrocyte-specific inactivation of ERK1 and ERK2 resulted  
3 in reduced osteoclastogenesis. Similar decrease in *RANKL* expression and osteoclast formation  
4 has been shown in mice in which *VDR* is conditionally inactivated in chondrocytes (25). These  
5 observations indicate that chondrocytes regulate osteoclast formation at the developing primary  
6 ossification center. Our results indicate that ERK1 and ERK2 play essential roles in supporting  
7 osteoclastogenesis through *RANKL* expression.

8  
9 In summary, our results demonstrate that ERK1 and ERK2 are essential for osteoblast  
10 differentiation, and ERK1 and ERK2 inhibit chondrogenic differentiation in the perichondrium  
11 (Fig. 11). Increased MAPK signaling promotes differentiation of mesenchymal cells into  
12 osteoblasts and inhibits chondrogenic differentiation. Based on these observations, we propose  
13 that ERK1 and ERK2 play essential roles in the lineage specification of mesenchymal cells.  
14 Furthermore, ERK1 and ERK2 regulate *RANKL* expression both in the osteoblast and  
15 chondrocyte lineages, which in turn regulates osteoclast formation. Further analyses will provide  
16 novel insights into the roles of MAPK in mesenchymal cells and skeletal development.

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1 **Figure 1**  
 2 *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* embryos and mice. (A) Skeletal preparation after alizarin red and  
 3 alcian blue staining at P1. Limbs were severely deformed in *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* mice.  
 4 (B) Skeletal preparation of the cranium after alizarin red staining. *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre*  
 5 mice showed bone defects in the calvaria at P12. (C) Hematoxylin, eosin, and alcian blue  
 6 staining of the humerus showing an absence of the primary ossification center and cortical bone  
 7 formation in *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* mice at P5. (D) Real-time PCR showed *ERK2*  
 8 inactivation in the tibia and humerus of *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* embryos at E16.5. (E)  
 9 Immunohistochemistry using anti-ERK1 and ERK2 antibody showed reduced immunoreactivity  
 10 in chondrocytes and cells in the bone forming region of tibia of *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre*  
 11 mice at P5. (a) *ERK1<sup>-/+</sup>; ERK2<sup>fllox/fllox</sup>* mice. The boxed area in the upper panel is magnified in the  
 12 lower panel. (b) *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* mice. Upper left panel shows immunostaining  
 13 for ERK1 and ERK2. The boxed areas 1 and 2 are magnified in the corresponding right panels.  
 14 While chondrocytes showed reduced immunoreactivity (1), endothelial cells show positive  
 15 staining (2). Lower left panel shows alcian blue, hematoxylin and eosin staining of a neighboring  
 16 section. Bars indicate 100  $\mu$ m.

1 **Figure 2**

- 2 (A,B,C) In situ hybridization analyses of the femur (A,B) and calvaria (C) showing normal levels  
 3 of *Colla1*, *Runx2*, and *Osterix (Osx)* expression and markedly decreased *Osteocalcin (OCN)*  
 4 expression in *ERK1<sup>-/-</sup>*; *ERK2<sup>flox/flox</sup>*; *Prx1-Cre* embryos and mice. (A) E15.5, (B) E16.5 and P5,  
 5 (C) P1.

1 **Figure 3**  
 2 (A) Semi-quantitative RT-PCR showing reduced *Osteocalcin (OCN)* and *Bone sialoprotein*  
 3 (*BSP*) expression in primary *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>* calvaria mesenchymal cells that were infected  
 4 with adenovirus expressing Cre recombinase (Ad-Cre). Primary calvaria mesenchymal cells  
 5 were isolated from E15.5 *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>* embryos and infected with Ad-Cre or adenovirus  
 6 expressing GFP (Ad-GFP). RNA was extracted 20 days after infection. (B) Western blot analysis  
 7 showing Osterix (OSX), ATF4, and RSK2 expression in primary *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>* calvaria  
 8 cells that were infected with Ad-Cre or Ad-GFP. ERK2 expression was inhibited 80% by Ad-Cre  
 9 infection, while Osterix, ATF4, and RSK2 expression remained largely unaffected. Total cell  
 10 lysates were prepared 10 days after infection. (C) von Kossa staining of *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>*  
 11 calvaria cell cultures 20 days after infection with Ad-Cre or Ad-GFP. Ad-Cre infection inhibited  
 12 mineralization. (D) Real time PCR analysis showed reduced *Krox20*, *c-Fos*, *Fra1*, *Fra2*, and  
 13 *Cbfb* expression in the humerus of *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>; Prx1-Cre* embryos at E16.5, while *JunB*  
 14 was not affected. Real time PCR analysis of the tibia and femur showed similar results. (E) Real  
 15 time PCR analysis showed reduced *Erk2*, *Krox20* and *c-Fos* expression in *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>*  
 16 calvaria cells infected with Ad-Cre. *ERK1<sup>-/-</sup>; ERK2<sup>fllox/fllox</sup>* calvaria cells were infected with  
 17 Ad-Cre or Ad-GFP, and RNA was extracted 9 days after infection.

1 **Figure 4**

2 Ectopic cartilage formation in the perichondrium of *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Prx1-Cre* embryos.

3 (A) Alcian blue staining and in situ hybridization of the femur at E15.5. The ectopic cartilage

4 (arrowheads) in the perichondrium expresses *Sox9*, *Col2a1*, and *Indian hedgehog (Ihh)*. (B)

5 Alcian blue staining (top panel) and immunohistochemical staining of the radius for β-catenin

6 (middle panel). *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Prx1-Cre* embryos showed reduced β-catenin protein levels

7 in the perichondrium at E16.5 (arrows). The boxed area was magnified in the bottom panel. (C)

8 *Tcf1* and *Dkk1* expression quantitated by real-time PCR. *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Prx1-Cre* embryos

9 showed reduced *Tcf1* and *Dkk1* expression in the tibia and humerus at E16.5.

1 **Figure 5**  
 2 Delayed formation of primary ossification centers in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice. (A)  
 3 Immunohistochemistry for type X collagen showed a widening of the zone of hypertrophic  
 4 chondrocytes in the tibia of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice at P0. (B) In situ hybridization  
 5 for *Coll10a1*, *Vegf*, *Mmp-13* and *Osteopontin* showed an expansion of terminally differentiated  
 6 chondrocytes in the hypertrophic zone of the tibia at P0. (C) TRAP staining showed an absence  
 7 of TRAP-positive cells in the tibia of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* embryos at E16.5. Arrows  
 8 indicate TRAP-positive cells. The upper panels in (A, B, C) show *ERK1*<sup>-/-</sup> and lower panels show  
 9 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* mice. (D) Real-time PCR showed reduced *RANKL* expression in  
 10 the tibia and humerus of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Prx1-Cre* embryos at E16.5. (E, F) *ERK2*, *RANKL*  
 11 and *Osteoprotegerin (OPG)* expression examined by real time PCR. Inactivation of *ERK2*  
 12 strongly inhibited *RANKL* expression in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup> rib chondrocytes (E) and calvaria  
 13 osteoblasts (F) *in vitro*. RNA was extracted from *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup> chondrocytes and  
 14 osteoblasts 5 days and 8 days after infection with adenovirus expressing Cre recombinase or GFP.  
 15 (G) ELF97-based fluorescent TRAP staining (green fluorescence) in combination with  
 16 immunofluorescence for ERK protein (red fluorescence), showing the presence of ERK protein  
 17 in TRAP-positive osteoclasts (arrows) in the femoral metaphysis of a *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;

1 *Prx1-Cre* mouse at P0. The immunofluorescent signal for ERK protein was indistinguishable  
 2 from that of *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup> mice (data not shown). Lower panels show magnification of an  
 3 osteoclast indicated by arrowheads in the upper panels. Nuclei were visualized by DAPI. (H)  
 4 X-gal staining followed by TRAP staining showing no β-galactosidase activity in TRAP-positive  
 5 osteoclast-like cells derived from spleen cells of *Prx1-Cre* mice harboring the *ROSA-LacZ*  
 6 reporter allele (right panel). The staining results were indistinguishable from TRAP-positive  
 7 osteoclast-like cells derived from control *ROSA-LacZ* reporter mice (left panel). (I) Spleen cells  
 8 from *ERK1*<sup>-/-</sup>; *ERK2*<sup>fllox/fllox</sup>; *Prx1-Cre* mice formed TRAP-positive multinucleated osteoclast-like  
 9 cells in the presence of M-CSF and RANKL (left panel). Nuclei were visualized by DAPI. These  
 10 osteoclast-like cells express ERK protein (right panel), and the staining intensity was  
 11 indistinguishable from that of osteoclast-like cells generated from spleen cells of *ERK1*<sup>-/-</sup>;  
 12 *ERK2*<sup>fllox/fllox</sup> mice (data not shown).

1 **Figure 6**

2 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos. (A) Immunofluorescence using anti-ERK1 and

3 ERK2 antibody showed reduced immunoreactivity in chondrocytes in the tibia of *ERK1*<sup>-/-</sup>;

4 *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos at E18.5. (B) ELF97-based fluorescent TRAP staining in

5 combination with immunofluorescence for ERK protein, showing the presence of ERK protein in

6 TRAP positive osteoclasts (arrows) of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos. (C)

7 ELF97-based fluorescent alkaline phosphatase staining in combination with immunofluorescence

8 for ERK protein, showing the presence of ERK protein in alkaline phosphatase positive

9 osteoblasts (arrows) of *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos. (D) Skeletal preparation after

10 alizarin red and alcian blue staining. *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos showed severe

11 kyphotic deformity in the thoracic spine at E18.5. (E, F) Hematoxylin, eosin, and alcian

12 blue-staining of the spine (E) and cranial base (F) showing an absence of ossification centers in

13 *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos at E18.5. (\*) indicates the ossification center in the

14 vertebral body, and arrowheads indicate the corresponding areas in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;

15 *Col2a1-Cre* embryos. (VB) vertebral body, (D) intervertebral disc, (SC) spinal cord, (Pi)

16 pituitary gland.

17

1 **Figure 7**  
 2 (A) Hematoxylin, eosin and alcian blue staining of the femur showing a dosage dependent  
 3 widening of the zone of hypertrophic chondrocytes at E16.5. Bars indicate the width of the zone  
 4 of hypertrophic chondrocytes. (B) In situ hybridization of the tibia showing an expansion of  
 5 *Col10a1*-expressing domains in *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Col2a1-Cre* embryos at E18.5. (C)  
 6 Primary *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup> chondrocytes were infected with adenovirus expressing Cre  
 7 recombinase or GFP at 200 MOI. *Col10a1* expression was examined by real time PCR at 5 days  
 8 after adenovirus infection. *ERK2* inactivation in *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup> chondrocytes increased  
 9 *Col10a1* expression. (D) Disorganization of columnar structures in the tibial growth plates of  
 10 *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Col2a1-Cre* embryos at E18.5. (E) BrdU incorporation of the distal  
 11 femoral growth plate. *ERK1*<sup>-/-</sup>; *ERK2*<sup>lox/lox</sup>; *Col2a1-Cre* embryos showed reduced chondrocyte  
 12 proliferation at E18.5. Data represent mean ± SD. (N.S.); Not significant. Analysis of variance  
 13 was used to detect significant difference. \*P<0.01

1 **Figure 8**

2 (A) TRAP staining of the femur showing decreased TRAP positive cells in *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup>;  
 3 *Col2a1-Cre* embryos at E16.5. (B) ELF97-based fluorescent TRAP staining in combination with  
 4 immunofluorescence for ERK protein. Spleen cells from *ERK1*<sup>-/-</sup>; *ERK2*<sup>flox/flox</sup> (a) and *ERK1*<sup>-/-</sup>;  
 5 *ERK2*<sup>flox/flox</sup>; *Col2a1-Cre* embryos (b,c) formed TRAP-positive multinucleated osteoclast-like  
 6 cells in the presence of M-CSF and RANKL. Nuclei were visualized by DAPI. Lower panels  
 7 show immunofluorescence using anti-ERK antibody (a' and b') or non-immune IgG (c') in  
 8 corresponding cells.

1 **Figure 9**

2 (A) Schematic representation of the construct that drives the expression of a constitutively active  
 3 mutant of *MEK1* and *LacZ* under the control of a 2.4 kb *prx1* promoter. (B) X-gal staining of an  
 4 E15.5 embryo showing transgene expression in the limb and cranium. (C) X-gal staining of the  
 5 distal ulna of an E15.5 embryo showing transgene expression in periarticular chondrocytes,  
 6 periosteum, and perichondrium. (D) Immunostaining of the FLAG-tagged MEK1(S218/222E,  
 7  $\Delta$ 32-51) using anti-M5 FLAG antibody showing transgene expression in periarticular  
 8 chondrocytes (arrows), periosteum (arrowheads), and perichondrium of the distal radius of a  
 9 *Prx1-MEK1* transgenic embryo at E15.5 (middle panel). No immunoreactivity was observed in a  
 10 wild type (Wt) littermate embryo (left panel). The immunostained section was further stained  
 11 with alcian blue and eosin (right panel). The cartilaginous matrix of transgene-expressing  
 12 periarticular chondrocytes (arrows) shows reduced alcian blue staining. (E) Skeletal preparation  
 13 of the forelimbs after alizarin red and alcian blue staining. Transgenic mice showed a thickening  
 14 and shorting of long bones at P8. (Wt) Wild type; (Tg) transgenic. (F) Skeletal preparation of the  
 15 cranium after X-gal and alizarin red staining showing transgene expression in the mesenchyme  
 16 of the lambdoid suture. Transgenic mice showed an accelerated closure of the lambdoid suture at  
 17 E17.5.

1 **Figure 10**  
 2 (A) Cross section of the forelimb stained with hematoxylin and eosin. *Prx1-MEK1* transgenic  
 3 mice showed increased bone formation and fusion of long bones (arrowhead) at P10. (Wt) Wild  
 4 type. (B) Hematoxylin, eosin, and alcian blue staining and in situ hybridization of the tibia at  
 5 E15.5. *Prx1-MEK1* transgenic embryos showed a thickening of the perichondrium, which  
 6 express *Runx2*, *Osterix (Osx)*, and *Bone sialoprotein (BSP)* (arrow heads). (C) Hematoxylin,  
 7 eosin and alcian blue staining of the foot. *Prx1-MEK1* transgenic embryos showed a delay in the  
 8 formation of cartilage anlage at E15.5. Tibia (Ti), Talus (Ta), Calcaneus (Ca) (D) In situ  
 9 hybridization of the carpal bones. *Prx1-MEK1* transgenic embryos showed reduced *Col2a1*  
 10 expression at E15.5. (E) Primary wild type chondrocytes were infected with adenovirus  
 11 expressing a constitutively active mutant of MEK1 (Ad-MEK1) or empty virus (Ad-Null).  
 12 *Col2a1* expression was examined by real time PCR at 48 h after adenovirus infection.  
 13 Expression of a constitutively active mutant of MEK1 strongly inhibited *Col2a1* expression. (F)  
 14 20 ng/ml FGF18 treatment downregulated *Col2a1* expression in wild type primary chondrocytes  
 15 at 24h. The downregulation was inhibited by U0126. (C) control, (F) FGF18, (U) U0126, (F+U)  
 16 FGF18 + U0126.

1 **Figure 11**

- 2 Model for the roles of MAPK in chondrocyte and osteoblast differentiation. While MAPK  
3 inhibits chondrogenic differentiation of osteo-chondroprogenitor cells, MAPK enhances  
4 osteoblast differentiation. MAPK is essential for osteoblast differentiation into  
5 *Osteocalcin*-expressing mature osteoblasts. MAPK also inhibits hypertrophic chondrocyte  
6 differentiation.





















