

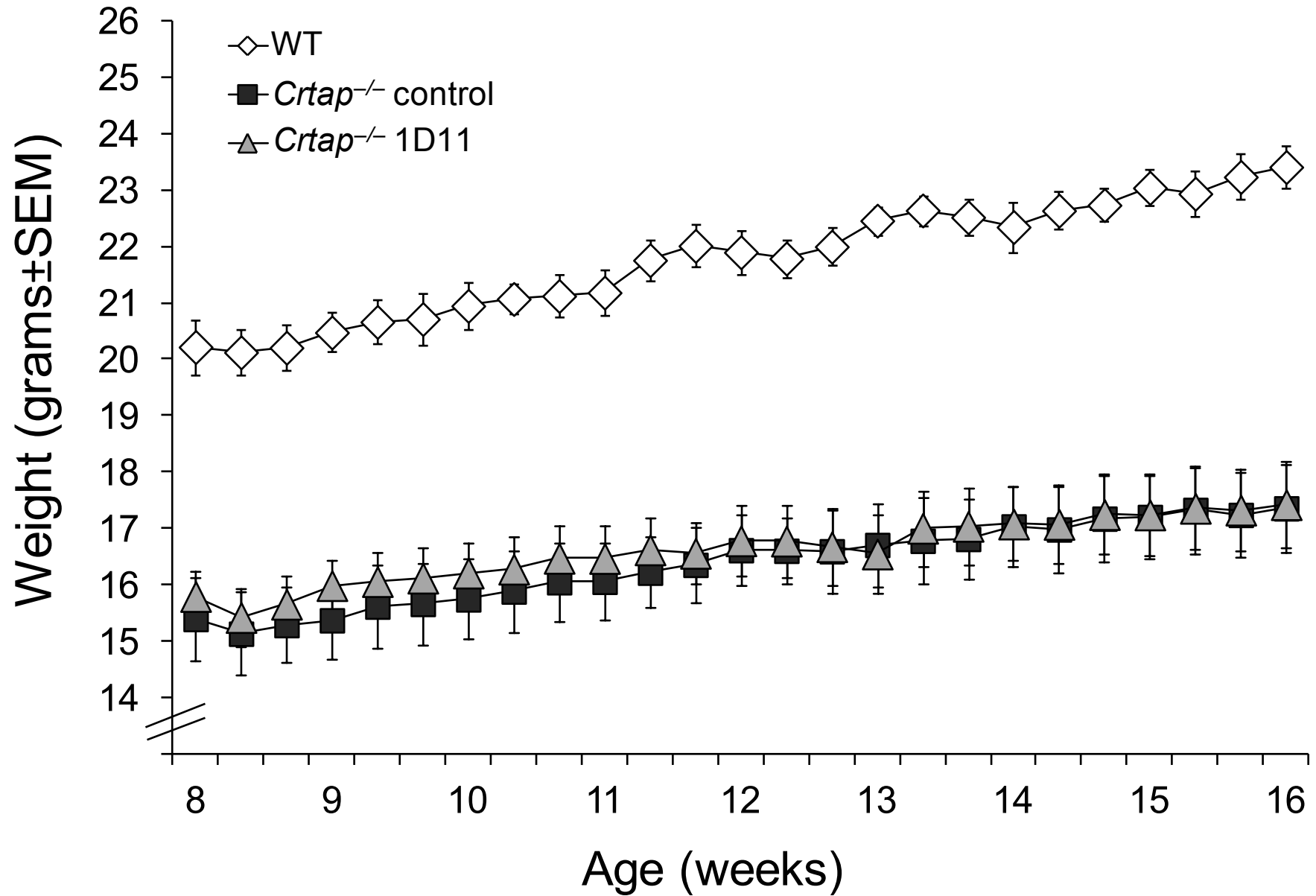
## Supplementary Information

Excessive TGF $\beta$  signaling is a common mechanism in Osteogenesis Imperfecta

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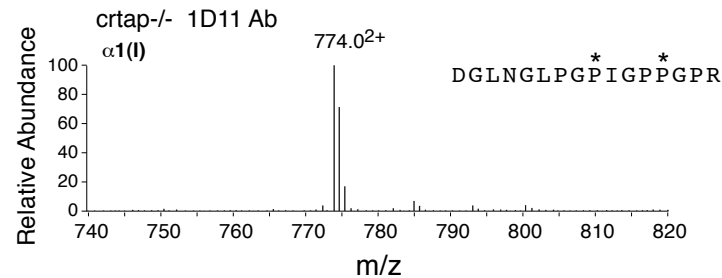
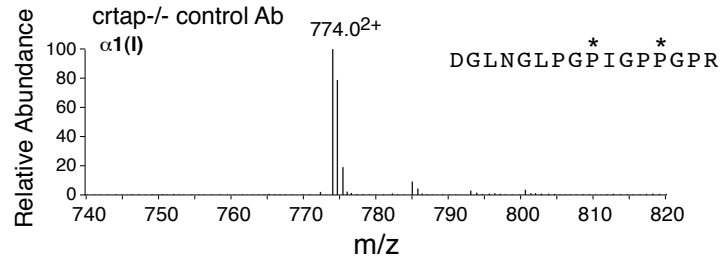
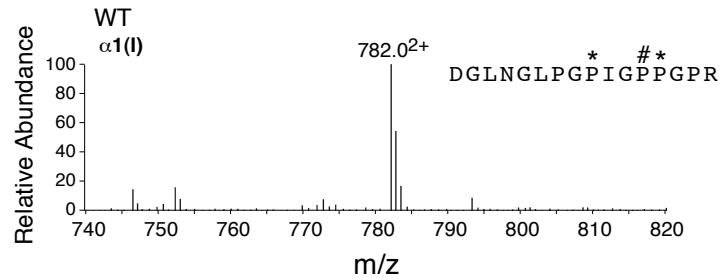
# Supplementary Figure 1



**Supplementary Figure 1.** Weight curves showing weight of WT, control *Crtap*<sup>-/-</sup> mice and 1D11-treated *Crtap*<sup>-/-</sup> mice during the study period. N=8 per group.

# Supplementary Figure 2

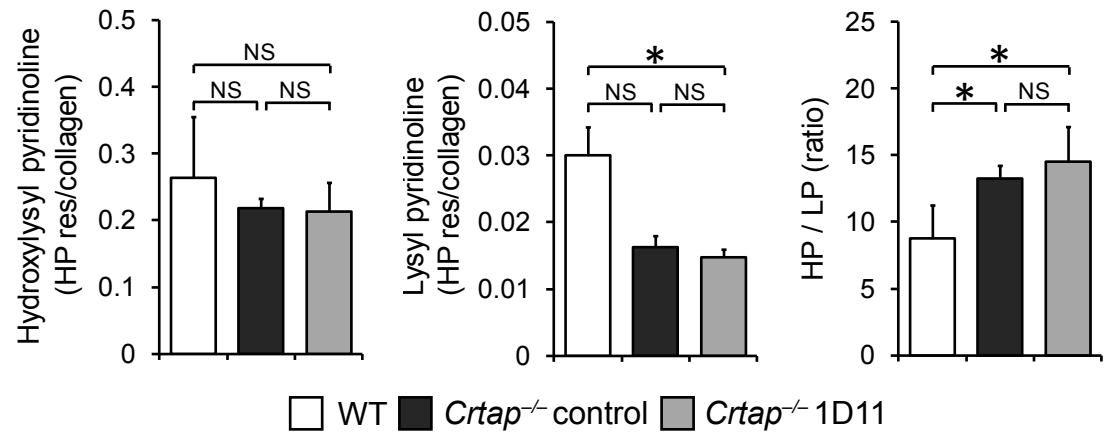
**a**



**b**

3Hyp	WT	<i>Crtap</i> <sup>-/-</sup> control	<i>Crtap</i> <sup>-/-</sup> 1D11
P986 alpha 1(I)	96.6% (2.1)	5.8% (3.3)	2.0% (1.2)

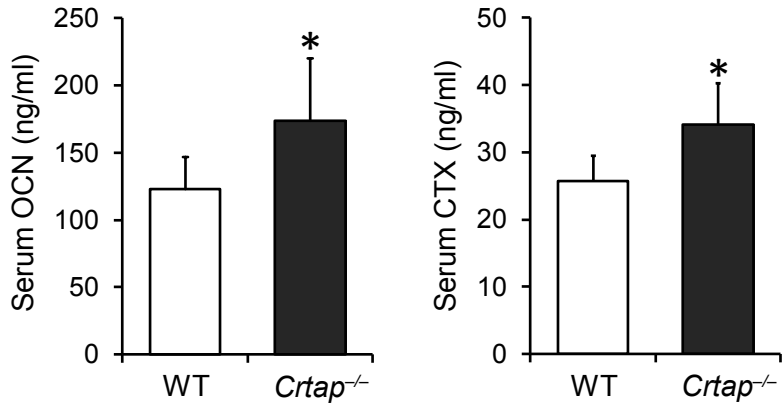
**c**



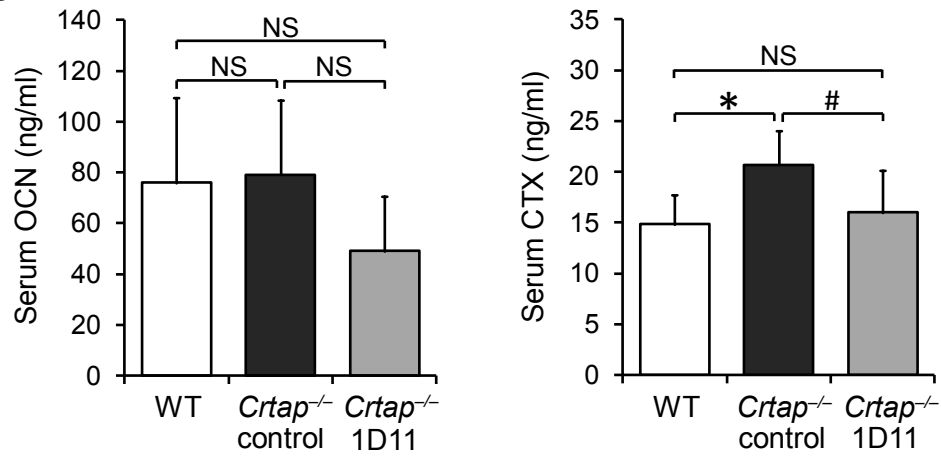
**Supplementary Figure 2.** No effect of TGF $\beta$  inhibition on the abnormal type I collagen post-translational modification in *Crtap*<sup>-/-</sup> mice. (a) Tandem mass spectra of extracted type I collagen from tibia of WT, control *Crtap*<sup>-/-</sup>, and 1D11- treated *Crtap*<sup>-/-</sup> mice (16 week old mice, after treatment for 8 weeks). (b) Status of collagen residue Pro986 alpha 1(I) 3-hydroxylation in bone samples of bone samples of WT, control *Crtap*<sup>-/-</sup> and 1D11- treated *Crtap*<sup>-/-</sup> mice assessed by tandem mass spectra analyses. Mean of percentage of 3-hydroxylated residues ( $\pm$ SD) is shown, n=5 per group. (c). Hydroxylysyl pyridinoline (HP), lysyl pyridinoline crosslinks (LP) levels and HP/LP ratio of bone type I collagen of WT, control *Crtap*<sup>-/-</sup> and 1D11- treated *Crtap*<sup>-/-</sup> mice. Results are given as means $\pm$ SDs, n=4 mice per group, \*P<0.05 for *Crtap*<sup>-/-</sup> vs. WT. NS, not significant.

# Supplementary Figure 3

**a**



**b**



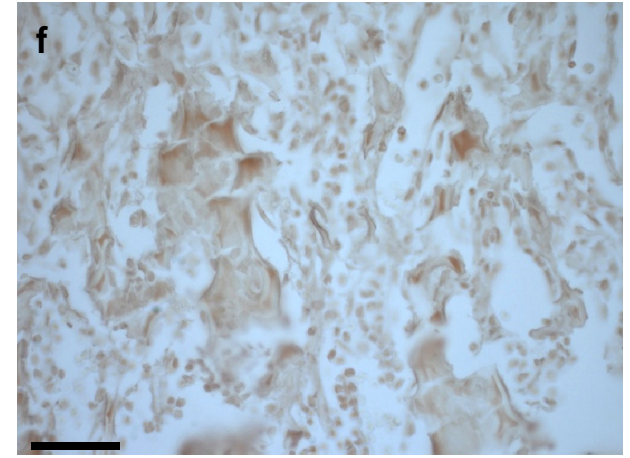
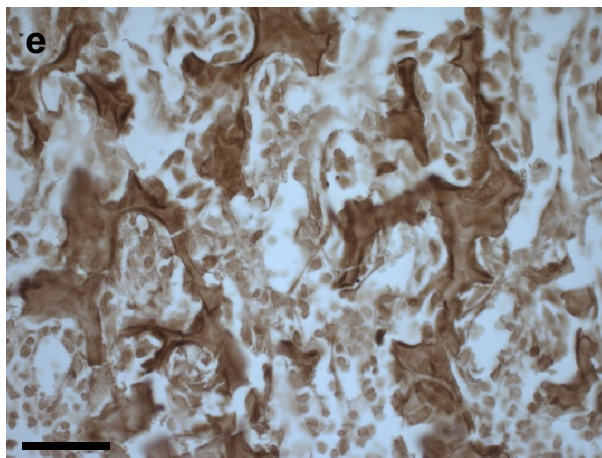
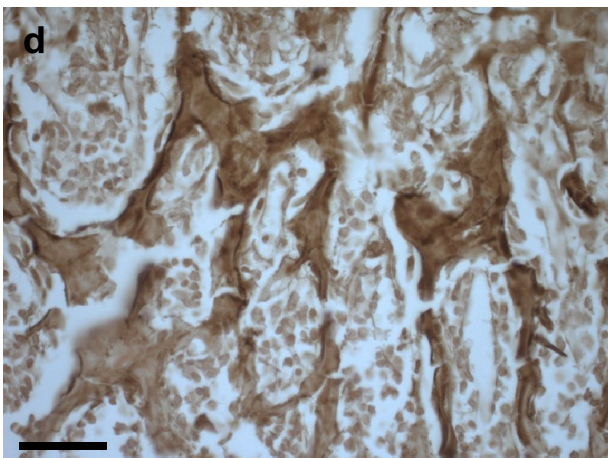
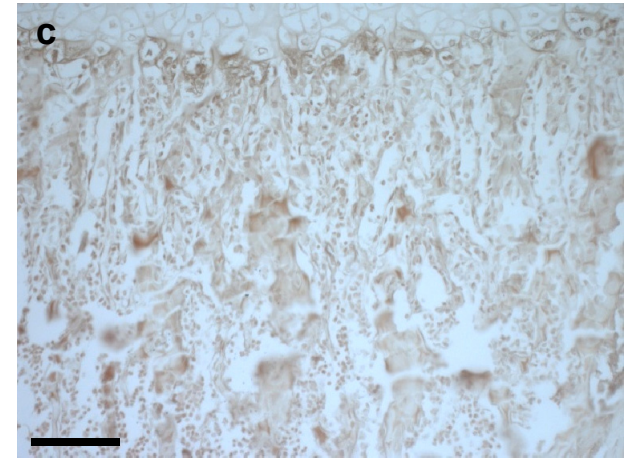
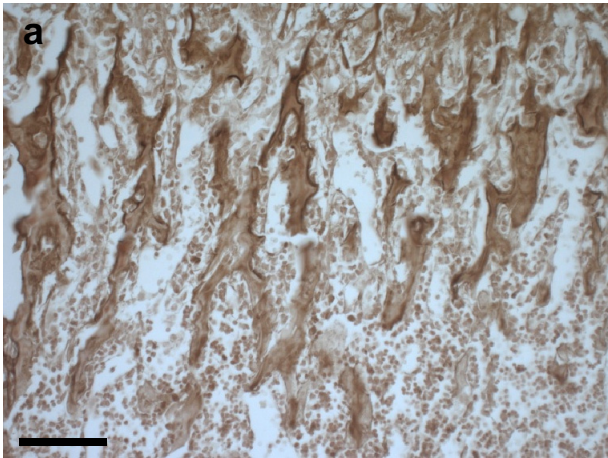
**Supplementary Figure 3.** Serum bone-turnover markers osteocalcin (OCN) and C-terminal cross-linked telopeptide of bone collagen (CTX) at start (a. 8 weeks of age) and end of the treatment study (b. 16 weeks of age). (a) Results are given as means±SDs, n=8 for WT, n=14 for *Crtap*<sup>-/-</sup> mice. (b) Results are given as means±SDs, n=8 for WT, n=7 per *Crtap*<sup>-/-</sup> group. \*P<0.05 for *Crtap*<sup>-/-</sup> vs. WT, #P<0.05 for *Crtap*<sup>-/-</sup> 1D11 vs. *Crtap*<sup>-/-</sup> control. NS, not significant.

## Supplementary Figure 4

WT

*Crtap*<sup>-/-</sup>

control



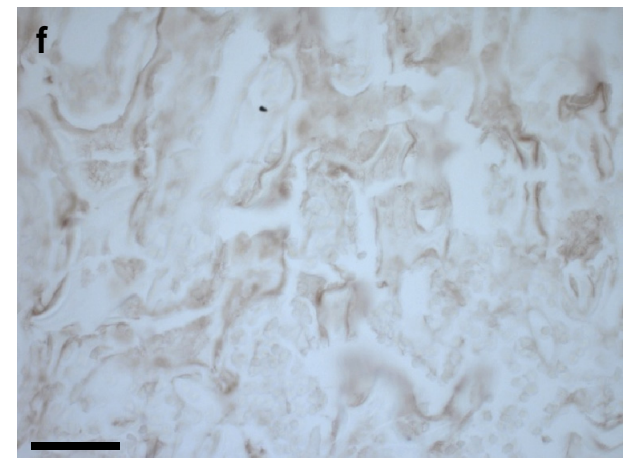
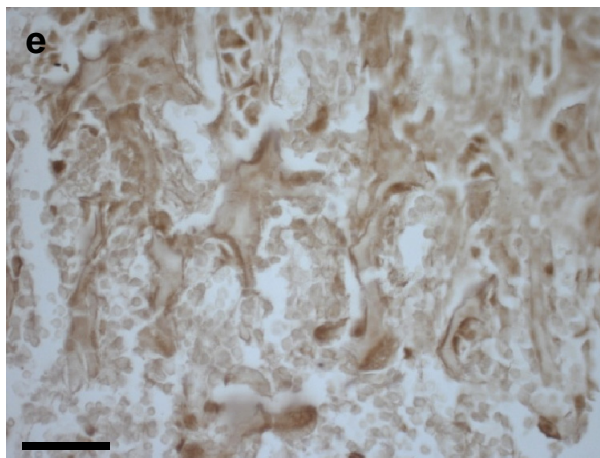
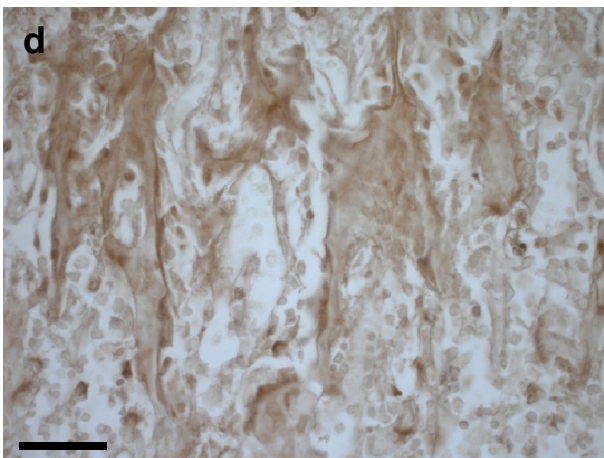
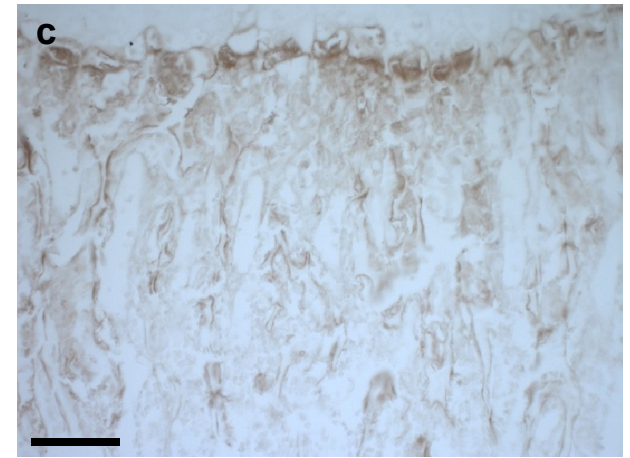
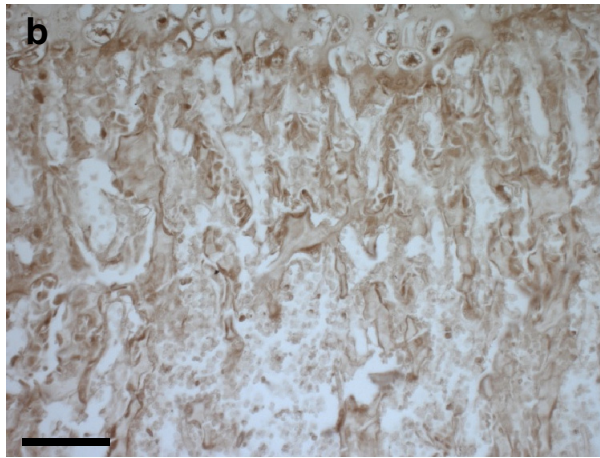
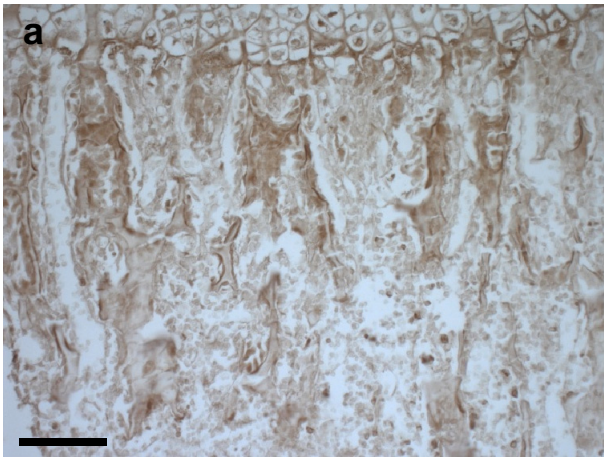
**Supplementary Figure 4.** Immunostaining for decorin in the distal femur metaphysis of WT and *Crtap*<sup>-/-</sup> mice is shown in 20X (a–c) and 40X magnification (d–f). Control femurs were incubated in secondary antibody only. (n=3 per genotype; scale bars=100µm (a–c), 50µm (d–f)).

## Supplementary Figure 5

WT

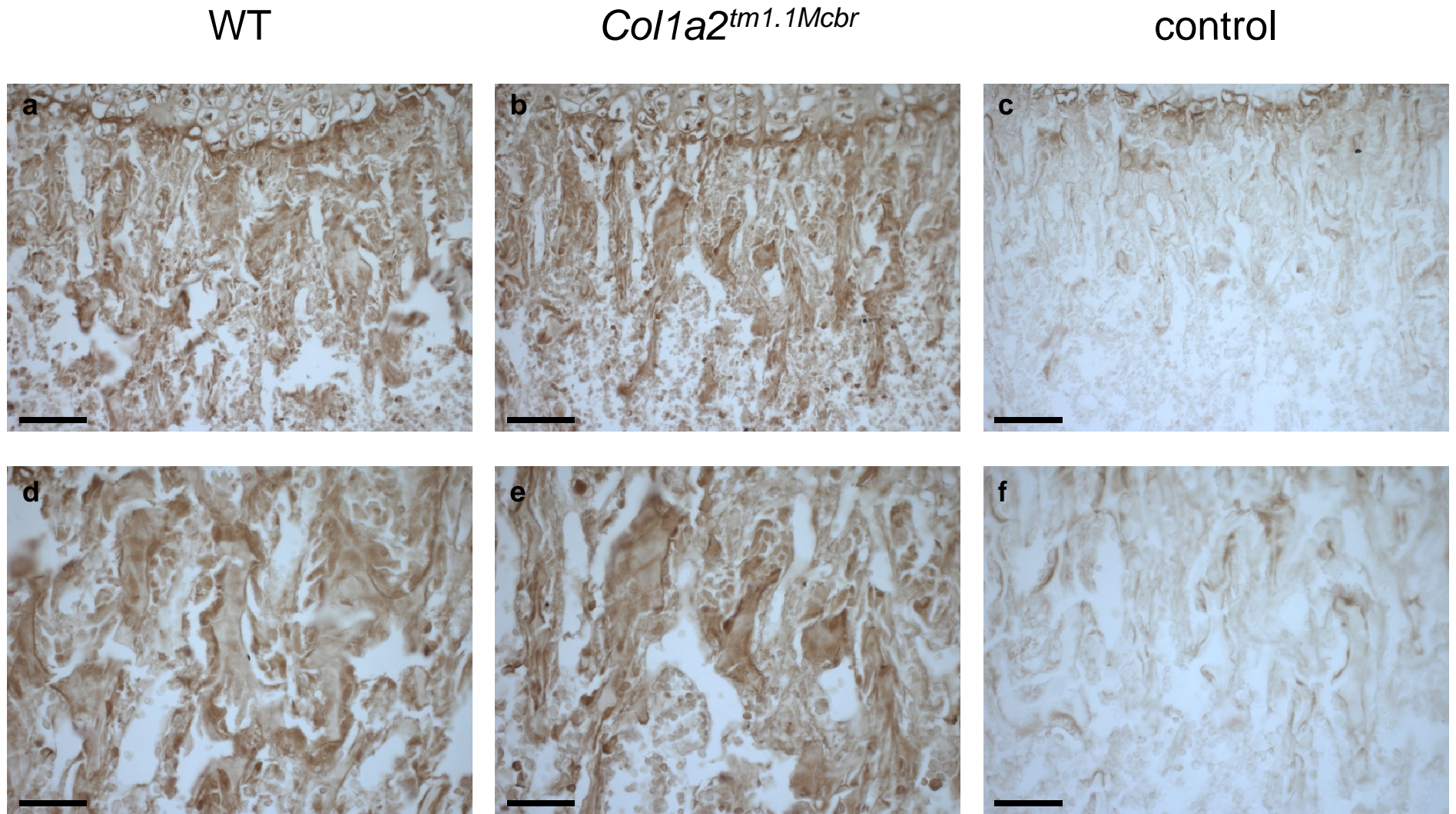
*Crtap*<sup>-/-</sup>

control



**Supplementary Figure 5.** Immunostaining for TGFβ1 in the distal femur metaphysis of WT and *Crtap*<sup>-/-</sup> mice is shown in 20X (a–c) and 40X magnification (d–f). Control femurs were incubated in secondary antibody only (n=3 per genotype; scale bars=100μm (a–c), 50μm (d–f)).

## Supplementary Figure 6



**Supplementary Figure 6.** Immunostaining for TGFβ1 in the distal femur metaphysis of WT and *Col1a2<sup>tm1.1Mabr</sup>* mice is shown in 20X (a–c) and 40X magnification (d–f). Control femurs were incubated in secondary antibody only. (n=3 per genotype; scale bars=100μm (a–c), 50μm (d–f)).

Supplementary Table 1

	<b>BV/TV</b> (%)	<b>Tb.N</b> (1/mm)	<b>Tb.Th</b> ( $\mu$ m)	<b>Tb.Sp</b> (mm)	<b>BMD BV</b> (mg HA/ccm)
<b>Wild type</b>	<b>31.173</b>	<b>4.845</b>	<b>63.988</b>	<b>0.145</b>	<b>703.189</b>
SD	6.543	0.583	9.482	0.034	19.014
<b><i>Crtap</i><sup>-/-</sup> control</b>	<b>8.354</b>	<b>2.130</b>	<b>38.888</b>	<b>0.450</b>	<b>677.416</b>
SD	2.045	0.409	2.931	0.118	16.138
<b><i>Crtap</i><sup>-/-</sup> 1D11</b>	<b>27.953</b>	<b>4.290</b>	<b>64.650</b>	<b>0.171</b>	<b>746.862</b>
SD	5.645	0.435	7.407	0.030	26.564
<b>ANOVA <i>P</i> value</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001<sup>+</sup></b>	<b>&lt;0.001<sup>+</sup></b>	<b>&lt;0.001</b>
<b>Pairwise <i>P</i> values</b>					
Wild type vs. <i>Crtap</i> <sup>-/-</sup> control	<0.001	<0.001	<0.05	<0.05	0.023
Wild type vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.	0.031	n.s.	n.s.	<0.001
<i>Crtap</i> <sup>-/-</sup> control vs. <i>Crtap</i> <sup>-/-</sup> 1D11	<0.001	<0.001	<0.05	<0.05	<0.001

**Supplementary Table 1.** MicroCT analyses of vertebral body L4 of WT, control *Crtap*<sup>-/-</sup> and 1D11 treated *Crtap*<sup>-/-</sup> mice (16 week old mice, after treatment for 8 weeks). Means $\pm$ SDs are shown for bone volume/tissue volume (BV/TV), trabecular number (Tb.N), trabecular thickness (Tb.Th), trabecular separation (Tb.Sp), and bone mineral density of bone volume (BMD BV); n=8 per group, + indicates Kruskal-Wallis one-way ANOVA on ranks where the equal variance test failed. n.s.=not statistically significant.



Supplementary Table 2

	<b>BV/TV</b> (%)	<b>Tb.N</b> (1/mm)	<b>Tb.Th</b> ( $\mu$ m)	<b>Tb.Sp</b> (mm)	<b>BMD BV</b> (mg HA/ccm)
<b>Wild type</b>	<b>10.698</b>	<b>2.808</b>	<b>37.413</b>	<b>0.341</b>	<b>733.864</b>
SD	3.558	0.684	5.071	0.108	25.997
<b><i>Crtap</i><sup>-/-</sup> control</b>	<b>2.873</b>	<b>0.861</b>	<b>32.863</b>	<b>1.209</b>	<b>727.151</b>
SD	1.082	0.237	3.428	0.345	28.765
<b><i>Crtap</i><sup>-/-</sup> 1D11</b>	<b>12.343</b>	<b>2.953</b>	<b>40.663</b>	<b>0.323</b>	<b>756.539</b>
SD	4.523	0.767	5.842	0.121	36.913
<b>ANOVA <i>P</i> value</b>	<b>&lt;0.001<sup>+</sup></b>	<b>&lt;0.001<sup>+</sup></b>	<b>0.015</b>	<b>&lt;0.001</b>	<b>0.162</b>
<b>Pairwise <i>P</i> values</b>					
Wild type vs. <i>Crtap</i> <sup>-/-</sup> control	<0.05	<0.05	n.s.	<0.001	
Wild type vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.	n.s.	n.s.	n.s.	
<i>Crtap</i> <sup>-/-</sup> control vs. <i>Crtap</i> <sup>-/-</sup> 1D11	<0.05	<0.05	0,004	<0.001	

**Supplementary Table 2.** MicroCT analyses of trabecular bone in proximal femurs for WT, control *Crtap*<sup>-/-</sup>, and 1D11- treated *Crtap*<sup>-/-</sup> mice (16 week old mice, after treatment for 8 weeks). Means $\pm$ SDs are shown for bone volume/tissue volume (BV/TV), trabecular number (Tb.N), trabecular thickness (Tb.Th), trabecular separation (Tb.Sp) and bone mineral density of bone volume (BMD BV); n=8 per group. + indicates Kruskal-Wallis one-way ANOVA on ranks where the equal variance test failed. n.s.=not statistically significant.

Supplementary Table 3

	<b>Cortical thickness</b> (mm)	<b>BMD BV</b> (mg HA/ccm)	<b>Diameter a.p.</b> (mm)	<b>CSA</b> (mm <sup>2</sup> )	<b>CSMI m.l.</b> (mm <sup>4</sup> )	<b>CSMI a.p.</b> (mm <sup>4</sup> )
<b>Wild type</b>	<b>0.242</b>	<b>1084.726</b>	<b>1.239</b>	<b>0.905</b>	<b>0.134</b>	<b>0.221</b>
SD	0.014	28.375	0.063	0.079	0.023	0.042
<b><i>Crtap</i><sup>-/-</sup> control</b>	<b>0.203</b>	<b>1084.885</b>	<b>1.142</b>	<b>0.731</b>	<b>0.101</b>	<b>0.162</b>
SD	0.020	34.256	0.065	0.082	0.021	0.034
<b><i>Crtap</i><sup>-/-</sup> 1D11</b>	<b>0.221</b>	<b>1096.127</b>	<b>1.186</b>	<b>0.808</b>	<b>0.111</b>	<b>0.186</b>
SD	0.026	39.754	0.080	0.118	0.028	0.038
<b>ANOVA <i>P</i> value</b>	<b>0.003</b>	<b>0.753</b>	<b>0.039</b>	<b>0.005</b>	<b>0.032</b>	<b>0.021</b>
<b>Pairwise <i>P</i> values</b>						
Wild type vs. <i>Crtap</i> <sup>-/-</sup> control	<0.001		0.012	0.001	0.011	0.006
Wild type vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.		n.s.	n.s.	n.s.	n.s.
<i>Crtap</i> <sup>-/-</sup> control vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.		n.s.	n.s.	n.s.	n.s.

**Supplementary Table 3.** MicroCT analysis of cortical bone at the femur midshaft for WT, control *Crtap*<sup>-/-</sup>, and 1D11-treated *Crtap*<sup>-/-</sup> mice (16 week old mice, after treatment for 8 weeks). Means±SDs are shown for cortical thickness, bone mineral density of bone volume (BMD BV), anterior-posterior (a.p.) diameter, cross-sectional area (CSA), and cross-sectional moments of inertia (CSMI) for medio-lateral (m.l.) and anterior-posterior (a.p.) axis; n=8 per group. n.s.=not statistically significant.

Supplementary Table 4

	Maximum load (N)	Stiffness (N/mm)	Energy to failure (N*mm)	Ultimate strength (MPa)	Toughness to failure (MPa)	Elastic modulus (GPa)	Total displacement (mm)	Elastic displacement (mm)	Post-yield displacement (mm)
<b>Wild type</b>	<b>22.831</b>	<b>230.578</b>	<b>7.846</b>	<b>154.704</b>	<b>11.912</b>	<b>6.960</b>	<b>0.444</b>	<b>0.068</b>	<b>0.376</b>
SD	2.860	37.053	3.985	7.478	5.695	0.731	0.207	0.014	0.203
<b><i>Crtap</i><sup>-/-</sup> control</b>	<b>12.943</b>	<b>151.689</b>	<b>1.228</b>	<b>114.496</b>	<b>2.208</b>	<b>6.663</b>	<b>0.127</b>	<b>0.079</b>	<b>0.048</b>
SD	2.402	27.384	0.913	18.240	1.677	1.154	0.057	0.006	0.058
<b><i>Crtap</i><sup>-/-</sup> 1D11</b>	<b>18.818</b>	<b>200.804</b>	<b>1.991</b>	<b>145.633</b>	<b>3.408</b>	<b>7.248</b>	<b>0.156</b>	<b>0.073</b>	<b>0.083</b>
SD	2.337	15.644	0.834	19.074	1.851	0.271	0.055	0.013	0.055
<b>ANOVA <i>P</i> value</b>	<b>&lt;0.001</b>	<b>0.009</b>	<b>0.009</b>	<b>0.004</b>	<b>0.009</b>	<b>0.658</b>	<b>0.015</b>	<b>0.360</b>	<b>0.012</b>
<b>Pairwise <i>P</i> values</b>									
Wild type vs. <i>Crtap</i> <sup>-/-</sup> control	<0.001	0.003	0.005	0.001	0.005		0.009		0.007
Wild type vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.	n.s.	0.017	n.s.	0.017		0.023		0.019
<i>Crtap</i> <sup>-/-</sup> control vs. <i>Crtap</i> <sup>-/-</sup> 1D11	0.015	n.s.	n.s.	0.016	n.s.		n.s.		n.s.

**Supplementary Table 4.** Results of biomechanical testing of femurs from WT, control *Crtap*<sup>-/-</sup> and 1D11 treated *Crtap*<sup>-/-</sup> mice by 3-point bending (16 week old mice, after treatment for 8 weeks). N=6 for WT, n=4 for control *Crtap*<sup>-/-</sup> and n=3 for 1D11 treated *Crtap*<sup>-/-</sup> mice. Results are shown as means±SDs. n.s.=not statistically significant.

Supplementary Table 5

	<b>BV/TV</b> (%)	<b>Tb.N</b> (1/mm)	<b>Tb.Th</b> ( $\mu$ m)	<b>Tb.Sp</b> ( $\mu$ m)	<b>N.Oc/BS</b> (1/mm)	<b>Oc.S/BS</b> (%)	<b>N.Ob/BS</b> (1/mm)	<b>Ob.S/BS</b> (%)	<b>N.Ot/B.Ar</b> (1/mm <sup>2</sup> )
<b>Wild type</b>	<b>11.956</b>	<b>4.003</b>	<b>29.554</b>	<b>226.736</b>	<b>3.226</b>	<b>15.051</b>	<b>18.142</b>	<b>20.827</b>	<b>549.002</b>
SD	3.067	0.656	3.536	50.077	0.812	3.089	2.368	2.754	83.665
<b><i>Crtap</i><sup>-/-</sup> control</b>	<b>3.817</b>	<b>1.909</b>	<b>19.502</b>	<b>587.008</b>	<b>4.768</b>	<b>19.885</b>	<b>24.487</b>	<b>26.888</b>	<b>735.561</b>
SD	1.656	0.692	2.560	289.835	0.914	3.407	5.421	6.983	72.617
<b><i>Crtap</i><sup>-/-</sup> 1D11</b>	<b>10.203</b>	<b>3.411</b>	<b>30.182</b>	<b>277.886</b>	<b>1.772</b>	<b>6.829</b>	<b>8.638</b>	<b>9.748</b>	<b>565.044</b>
SD	3.097	0.878	6.982	69.118	0.673	2.235	2.855	4.718	51.014
<b>ANOVA <i>P</i> value</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>0.002</b>	<b>0.003<sup>+</sup></b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>
<b>Pairwise <i>P</i> values</b>									
Wild type vs. <i>Crtap</i> <sup>-/-</sup> control	<0.001	<0.001	0.002	<0.05	0.005	0.012	0.011	n.s.	<0.001
Wild type vs. <i>Crtap</i> <sup>-/-</sup> 1D11	n.s.	n.s.	n.s.	n.s.	0.007	<0.001	<0.001	0.002	n.s.
<i>Crtap</i> <sup>-/-</sup> control vs. <i>Crtap</i> <sup>-/-</sup> 1D11	<0.001	0.003	0.001	<0.05	<0.001	<0.001	<0.001	<0.001	<0.001

**Supplementary Table 5.** Histomorphometry analyses of L4 vertebral bodies of WT, control *Crtap*<sup>-/-</sup>, and 1D11- treated *Crtap*<sup>-/-</sup> mice (16 week old mice, after treatment for 8 weeks). Means $\pm$ SDs are shown for bone volume/tissue volume (BV/TV), trabecular number (Tb.N), trabecular thickness (Tb.Th), trabecular separation (Tb.Sp), number of osteoclasts/bone surface (N.Oc/BS), osteoclast surface/bone surface (Oc.S/BS), number of osteoblasts/bone surface (N.Ob/BS), osteoblast surface/bone surface (Oc.S/BS), and number of osteocytes/bone area (N.Ot/B.Ar); n=6 per group. + indicates Kruskal-Wallis one way ANOVA on ranks where equal variance test failed. n.s.=not statistically significant.

Supplementary Table 6

	<b>BV/TV</b> (%)	<b>Tb.N</b> (1/mm)	<b>Tb.Th</b> ( $\mu$ m)	<b>Tb.Sp</b> (mm)	<b>BMD BV</b> (mg HA/ccm)
<b>Wild type</b>	<b>24.708</b>	<b>3.939</b>	<b>62.533</b>	<b>0.193</b>	<b>716.669</b>
SD	3.282	0.330	3.497	0.025	7.626
<b><i>Col1a2</i><sup>tm1.1Mcb</sup> control</b>	<b>10.608</b>	<b>2.071</b>	<b>51.067</b>	<b>0.435</b>	<b>731.436</b>
SD	1.487	0.195	2.999	0.047	11.792
<b><i>Col1a2</i><sup>tm1.1Mcb</sup> 1D11</b>	<b>29.715</b>	<b>4.244</b>	<b>69.983</b>	<b>0.166</b>	<b>754.115</b>
SD	1.984	0.199	2.329	0.013	7.844
<b>ANOVA P value</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>	<b>&lt;0.001</b>
<b>Pairwise P values</b>					
Wild type vs. <i>Col1a2</i> <sup>tm1.1Mcb</sup> control	<0.001	<0.001	<0.001	<0.001	0.015
Wild type vs. <i>Col1a2</i> <sup>tm1.1Mcb</sup> 1D11	0.002	n.s.	<0.001	n.s.	<0.001
<i>Col1a2</i> <sup>tm1.1Mcb</sup> control vs. <i>Col1a2</i> <sup>tm1.1Mcb</sup> 1D11	<0.001	<0.001	<0.001	<0.001	<0.001

**Supplementary Table 6.** MicroCT analyses of vertebral body L4 of WT, control *Col1a2*<sup>tm1.1Mcb</sup> and 1D11 treated *Col1a2*<sup>tm1.1Mcb</sup> mice (16 week old mice, after treatment for 8 weeks). Means $\pm$ SDs are shown for bone volume/tissue volume (BV/TV), trabecular number (Tb.N), trabecular thickness (Tb.Th), trabecular separation (Tb.Sp), and bone mineral density of bone volume (BMD BV); n=6 per group, n.s.=not statistically significant.