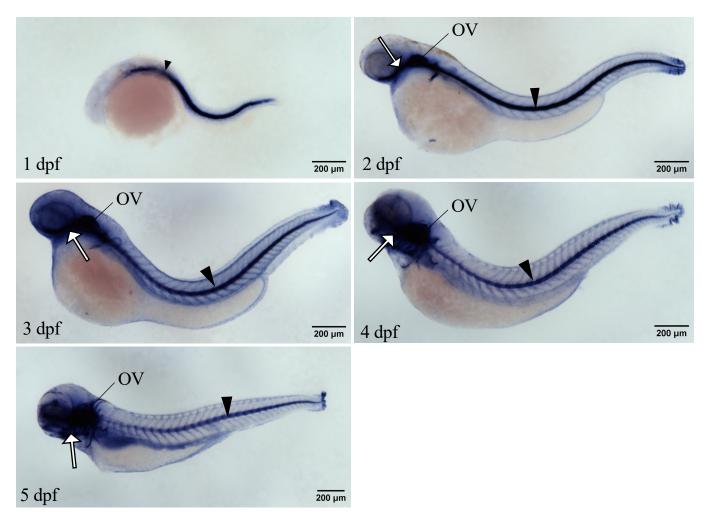
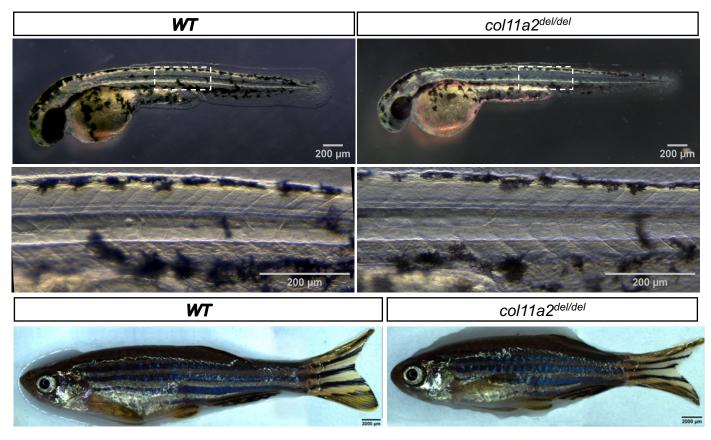


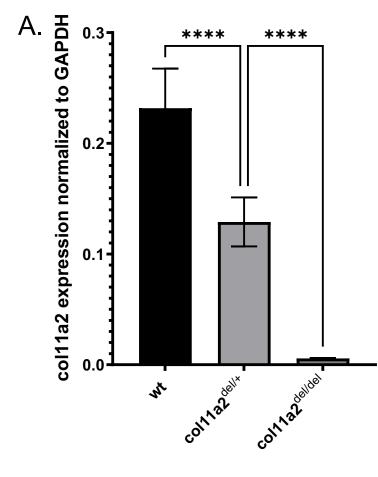
**Supplemental Figure 1. Radiographic illustration of proband vertebral malformations.** (a) Proband 1: x-ray showing fused C4-C5 cervical vertebrae in a 46 year old female (b) Proband 2: CT scan showing fused C3-C5 cervical vertebrae in a 31 year old male. (c) Proband 3: 20-year old male with congenital scoliosis and hemivertebra of T9 demonstrated by x-ray.



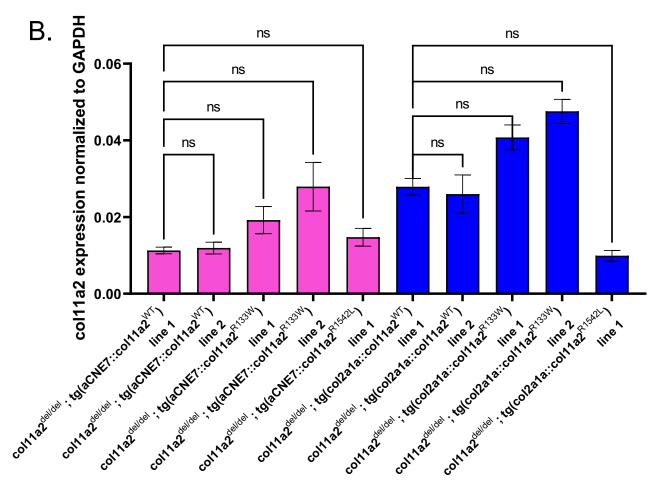
Supplemental Figure 2. Whole mount RNA *in situ* hybridization demonstrating the spatiotemporal expression pattern of *coll1a2* in zebrafish embryos from 1-5dpf. *coll1a2* is expressed in the developing notochord (arrowheads) and craniofacial cartilage (white arrows), consistent with a role in skeletal development.

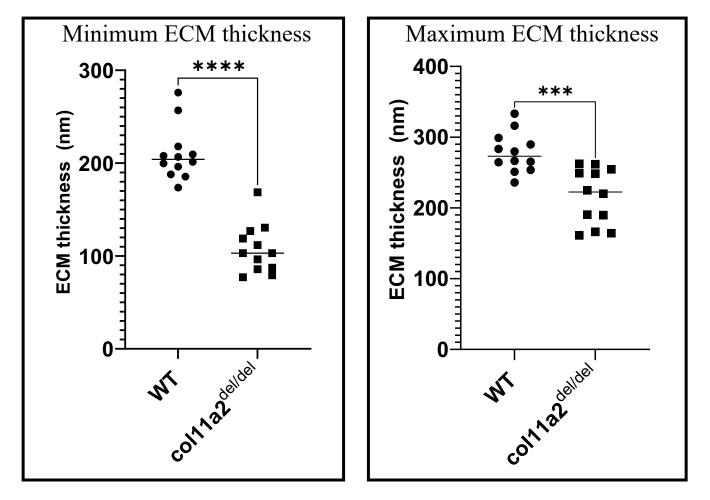


**Supplemental Figure 3.** *coll1a2*<sup>del/del</sup> **mutant animals appear morphologically normal.** Live, whole-mount images of wildtype (left) and *coll1a2*<sup>del/del</sup> mutant (right) fish at 48hpf embryonic (top) and adult (bottom) stages. 2dpf *coll1a2*<sup>del/del</sup> embryos do no exhibit obvious morphological defects. Vertebral fusions in adult *coll1a2*<sup>del/del</sup> animals do not result in axial curvatures, but mutant fish do appear shorter in length than wildtype, and exhibit a craniofacial defect.

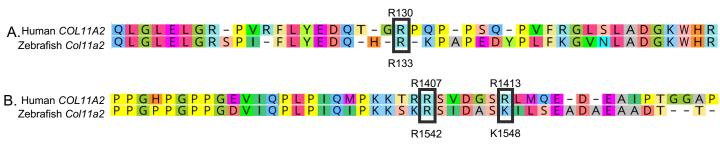


Supplemental Figure 4. RT-qPCR analysis of coll1a2 expression levels in 2dpf embryos. A. collla2 expression in *coll1a2*<sup>*del/+*</sup> embryos is approximately half of of wildtype expression levels and is almost undetectable in *coll1a2*<sup>del/del</sup> mutant embryos. B. No significant differences in coll1a2 expression are observed between wildtype and patient-mutation bearing colla2 transgenic lines. All expression levels were normalized to gapdh expression within the same sample. Each genotype included three biological replicates, with three technical replicates of each. Error bars represent standard deviation and significance was calculated by one-way ANOVA with Dunnett's *post hoc* test. \* p<0.05, \*\* p<0.01, \*\*\* p<0.001, \*\*\*\* p<0.0001.





Supplemental Figure 5. Notochord ECM is significantly thinner in *coll1a2*<sup>del/del</sup> mutant embryos. Quantification of notochord ECM thickness from TEM images of 2 dpf wildtype and *coll1a2*<sup>del/del</sup> mutant embryos. Both the minimum thickness (A) and maximum thickness (B) are significantly reduced in *coll1a2*<sup>del/del</sup> embryos. n=6 for each genotype, thickness measured on both lateral sides of TEM notochord cross-sections. Significance calculated using Student's t-test. \* p<0.05, \*\* p<0.01, \*\*\* p<0.001, \*\*\*\* p<0.0001.



**Supplemental Figure 6. Partial alignment of the human and zebrafish Coll1a2 proteins.** Panel A shows the alignment of the early part of the protein, highlighting conserved human R130 and zebrafish R133 residues. Panel B shows the alignment of the C terminal end of the triple helix domain. Human R1407 is conserved in the fish and aligns with R1542. R1413 in the human protein is not conserved in zebrafish.