

1 **Altered vertebrae morphology and bone mineralization in a zebrafish model of CHARGE**  
2 **syndrome**

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25 **Conflict of Interest**

26 The authors declare no conflicts of interest

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33 **Abstract**

34 CHARGE syndrome patients present features of idiopathic scoliosis in over 60% of cases,  
35 reduced bone mineral density and in a few cases osteopenia. While several clinical cases and  
36 studies regarding the spinal deformities in CHARGE syndrome bearing *CHD7* mutations are  
37 well-documented, the underlying mechanisms remain elusive. Here, we detect and  
38 quantitatively analyze skeletal abnormalities in adult and young *chd7<sup>-/-</sup>* larvae.

39 We show that young *chd7<sup>-/-</sup>* larvae present with scoliosis and kyphosis already at 9 dpf. Gene  
40 expression analysis confirmed the reduction of osteoblast markers and Ppar $\gamma$  targets. MicroCT  
41 analyses identified abnormal Weberian apparatus structure and vertebral body morphology in  
42 *chd7<sup>-/-</sup>* mutants, with highly mineralized inclusions, along with variances in bone mineral  
43 density and bone volume. Additionally, we detect a specific depletion of *Col2a1a* in the  
44 zebrafish vertebral cartilage, in line with a significantly reduced number of chondrocytes.

45 Our study is the first to elucidate the mechanisms underlying morphological changes in  
46 vertebrae of adult *chd7<sup>-/-</sup>* zebrafish and decreased spinal integrity. The *chd7<sup>-/-</sup>* zebrafish will be  
47 beneficial in future investigations of the underlying pathways of spinal deformities in CHARGE  
48 syndrome.

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## 56 **Introduction**

57 Impaired bone development and spinal dysmorphisms are a major health concern in various  
58 genetic disorders. Among these, Idiopathic scoliosis is a commonly observed phenotype and  
59 the most common skeletal abnormality in children.<sup>(1)</sup> Idiopathic scoliosis is a curvature of the  
60 spine with unidentified genetic cause. However, many risk genes have been connected to the  
61 underlying mechanisms.<sup>(2)</sup> One of these related genetic disorders is the congenital  
62 multisystemic CHARGE syndrome (CS) named after the major characteristics of **C**oloboma,  
63 **H**ear defects, **A**tresia chonae, **R**etarded growth, **G**enital and **E**ar abnormalities.<sup>(3-5)</sup> Though,  
64 considering the wide variety of observed phenotypes, the list of characteristics has been  
65 continuously augmented to include craniofacial abnormalities and spinal deformations.<sup>(6-8)</sup>  
66 Along these lines, CS is closely associated with skeletal deformities such as idiopathic scoliosis,  
67 kyphosis and hemivertebrae.<sup>(9)</sup> In fact, idiopathic scoliosis is observed in a majority of CS  
68 cases, with studies showing over 60% of patients having diagnosed scoliosis at an average age  
69 of just over 6 years.<sup>(10-12)</sup> In some cases, the areal bone mineral density (aBMD) is reduced.<sup>(13)</sup>  
70 CS is most commonly caused by a mutation in the chromodomain ATP-dependant helicase 7  
71 (CHD7). Mutations are distributed evenly throughout the gene and the vast majority of cases  
72 are sporadic, with only very few cases of familial CS.<sup>(7, 14-16)</sup> Analysis of pathways related to  
73 CHD7 show involvement in neural crest differentiation/proliferation/migration and stem cell  
74 quiescence.<sup>(17-19)</sup> Furthermore, studies focusing on CS have identified regulatory mechanisms  
75 of CHD7 in the immune response and more strikingly in brain development.<sup>(20, 21)</sup> Most  
76 recently, CHD7 has been connected to abnormal GABAergic development resulting in an  
77 autistic-like behavior (Jamadagni et al., unpublished data). While the function of CHD7 has  
78 been extensively studied, little attention has been given to the underlying mechanisms in  
79 skeletal development.<sup>(22)</sup> Some studies have directly linked CHD7 to osteogenesis.<sup>(23, 24)</sup>  
80 Investigation in cell cultures have shown a dependency on CHD7 for successful differentiation,

81 more specifically to a differentiation complex with SETD1B, NLK and SMAD1 resulting in  
82 depleted osteogenesis upon depletion of CHD7 in favour of adipogenesis by regulating PPAR $\gamma$   
83 target genes.<sup>(23, 24)</sup> Various models to investigate this role in vivo have been proposed. A mouse  
84 model for Chd7 deficiency, termed “looper” presents with ear ossicle malformations, but no  
85 spinal deformities.<sup>(25)</sup> However, mouse models for Chd7 deficiency have limitations, as null  
86 mutants are embryonically lethal by 10 days.<sup>(26)</sup> Yet, a recent zebrafish model has proven  
87 valuable in the modelling of *chd7* dependent CS revealing a reduction in vertebrae  
88 mineralization of young larvae.<sup>(27)</sup>

89 Zebrafish have become increasingly relevant in the study of fundamental bone development  
90 and bone related disorders<sup>(28-30)</sup>, including scoliosis, osteoporosis, age related osteoarthritis.<sup>(31,</sup>  
91 <sup>32)</sup> Notably, the spinal structure in zebrafish has been characterized in detail and comprises three  
92 major regions: the Weberian apparatus, a specialized structure consisting of the first four  
93 vertebrae connecting the auditory system and the swim bladder to amplify sound vibrations, the  
94 precaudal vertebrae which are connected to neural arch and spines to form the chest and, the  
95 caudal vertebrae of the tail region with neural and hemal arch.<sup>(33)</sup> Simplicity of analysis in  
96 zebrafish to investigate spinal structures has been used to understand the effects of mechanical  
97 loading on bone mineralization.<sup>(30, 34)</sup> Zebrafish have specific advantages in the analysis of  
98 skeletal development such as the closely related structure to humans, rapid bone development  
99 with first mineralization of vertebrate occurring after only 5 days post fertilization (dpf), as well  
100 as *ex utero* development and transparency of young larvae simplifying the use of *in vivo* staining  
101 and transgenic lines to allow for the analysis of early calcification in zebrafish bony  
102 structures.<sup>(35-38)</sup> Furthermore, the simplicity of high-throughput drug screening in this teleost  
103 model makes the model highly useful for investigating new drug targets and effect on skeletal  
104 development.<sup>(39, 40)</sup> These advantages have raised interest due to their high remodelling  
105 efficiency of skeletal structures including their possibility to regenerate fins, as well as the

106 effectiveness to reproduce spinal deformity phenotypes.<sup>(41, 42)</sup> Some of these studies in zebrafish  
107 show a close link to the Collagen Family.<sup>(43, 44)</sup> These highly conserved pathways between  
108 zebrafish and humans allow us to investigate the underlying pathomechanisms linking CHD7  
109 to idiopathic scoliosis and other skeletal abnormalities.

110 We recently generated a stable *chd7*<sup>-/-</sup> zebrafish mutant using CRISPR/Cas9 that replicates  
111 hallmarks of CS (Jamadagni et al., unpublished data), consistent with our previous findings  
112 using a transient *chd7* morpholino knockdown model.<sup>(27)</sup> In this study we describe for the first  
113 time a detailed analysis of skeletal development in a zebrafish *chd7*<sup>-/-</sup> mutant model for CS.  
114 Early larvae screens reveal a delay of mineralization of vertebrae bodies linked to deficient  
115 osteoblast differentiation. Gene expression and IHC analysis reveals striking reduction in  
116 osteoblast, chondrocyte, and collagen matrix markers, particularly displaying diminished levels  
117 of Col2a1. Finally, extensive MicroCT analysis in adults show abnormal mineralization and  
118 morphological structures in precaudal and caudal vertebrae.

## 119 **Results**

### 120 *Zebrafish chd7*<sup>-/-</sup> larvae show spinal deformities

121 To investigate if spinal deformities are present at an early developmental stage, we screened  
122 young *chd7*<sup>-/-</sup> larvae at 9 dpf (n=110) for morphological abnormalities. 39% of *chd7*<sup>-/-</sup> larvae  
123 exhibited highly varying, scoliosis-like phenotypes at both precaudal and caudal regions of the  
124 spine (Fig. 1a). Additionally, some of these also presented with a kyphosis-like phenotype  
125 (Supplemental Fig. 1a). Even with the morphological phenotype, fish were reactive to a touch  
126 response.

### 127 *chd7*<sup>-/-</sup> larvae have a spinal mineralization deficit that is nutrient dependent

128 Given the spinal deformities in 9dpf *chd7*<sup>-/-</sup> larvae, we next sought to investigate the efficiency  
129 of mineralization during larval development. Calcein staining for fluorochrome labeling of

130 bones revealed no delay in the onset of mineralization of the first precaudal vertebrae at 6 dpf.

131 Upon further development, at 9dpf, *chd7*<sup>-/-</sup> larvae showed a significant reduction in the number

132 of calcified vertebrae (Fig. 1c). Notably, *chd7*<sup>-/-</sup> larvae were less efficient in complete

133 calcification of vertebrae towards the posterior region of the spine (Fig. 1b, c).

134 Since calcification of the bone matrix is dependent on mineral uptake from food, we tested how

135 the onset and development of vertebral bone structure calcification in 9 dpf *chd7*<sup>-/-</sup> zebrafish

136 was dependent on nutrition accessibility. Notably, wildtype and *chd7*<sup>-/-</sup> larvae showed no

137 difference in the number of mineralized vertebrae in a starvation situation between 5 dpf and 9

138 dpf. Expectedly, we observed significant differences when food was available from 5 dpf until

139 9 dpf, with *chd7*<sup>-/-</sup> zebrafish having a reduced number of mineralized vertebrae compared to

140 controls. The *chd7*<sup>-/-</sup> larvae had the same number of mineralized vertebrae when food was

141 available or absent (Fig. 1b, d). Starvation in zebrafish larvae between 5 dpf and 9 dpf also

142 resulted in smaller yolks, which is to be expected, considering the high dependency on nutrition

143 supply by the yolk. Surprisingly, we found that Weberian vertebrae, which are the first 4

144 vertebrae of the spinal column, were more rapidly calcified in *chd7*<sup>-/-</sup> larvae than in controls

145 (Fig. 1b arrows, Fig. 1e).

146 We investigated if mineralization was affected in later development. Insufficient mineralization

147 was also evident in 4-week-old juvenile zebrafish with spinal deformities. Alizarin red staining

148 revealed decreased mineralization of caudal vertebrae, with striking deficiency toward the

149 vertebral body (Fig. 1f).

#### 150 *Osteoblast differentiation is impaired in chd7*<sup>-/-</sup> larvae

151 CHD7 has been shown to regulate osteogenesis by controlling PPAR $\gamma$  to promote skeletal

152 development and regulate expression of promoting genes. To understand if the Ppar $\gamma$  pathway

153 is directly involved in the observed bone mineralization deficiency and abnormal morphology

154 in *chd7*<sup>-/-</sup> larvae, we tested for Ppar $\gamma$  target gene expression and osteoblast differentiation

155 marker genes at the onset of the defect at 9 dpf (Fig 1g). The key target for Pparg, *runx2a* and  
156 *runx2b*, were significantly downregulated in *chd7<sup>-/-</sup>* larvae compared to controls. Further  
157 markers for osteogenesis were affected, including a significant downregulation of osteoblast  
158 markers *sp7/osterix*, *bglap* and *postna*. The *chd7<sup>-/-</sup>* larvae also had significantly lower levels of  
159 osteocyte markers and other genes regulating mineralization such as *acp5a* and *sost*, while *mgp*  
160 remained unchanged compared to control larvae. Additionally, we detected a significant  
161 upregulation of osteoclast marker *ctsk* in *chd7<sup>-/-</sup>* larvae.

### 162 *Abnormal development of the Weberian apparatus in adult chd7<sup>-/-</sup> zebrafish*

163 To assess the effects of *chd7* deficiency on vertebral growth and maturation, we analyzed  
164 vertebrae morphology and mass using microCT in adult zebrafish. Morphologically, adult *chd7<sup>-/-</sup>*  
165 zebrafish were unchanged in body length, however, appeared strikingly leaner in body stature  
166 compared to adult control zebrafish.

167 Since mineralization in the young larvae varied from anterior to posterior, we decided to  
168 analyze the three major structures of the zebrafish spinal column: The Weberian apparatus,  
169 precaudal and caudal vertebrae. The Weberian apparatus and its specialized structures, the  
170 supraneurals, intercalarium, tripus and parapophysis, were thinner and smaller in morphology  
171 in *chd7<sup>-/-</sup>* adults in comparison to controls (Fig. 2 b). We further performed a detailed analysis  
172 of volumetric bone mineral density (vBMD), bone volume (BV), total volume (TV), and bone  
173 volume fraction (BV/TV) of 1-year-old *chd7<sup>-/-</sup>* larvae and control zebrafish. We did not detect  
174 any significant differences in vBMD, BV or TV (Fig. 2 c-h). However, we found a significantly  
175 greater variance in TV of both intercalarium and parapophysis, as well as in BV of the  
176 parapophysis in 1-year-old *chd7<sup>-/-</sup>* zebrafish, as determined by F-test analysis (Supplemental  
177 Table 1). The same results were observed in 2-year-old *chd7<sup>-/-</sup>* compared to control zebrafish  
178 (Supplemental Table 2).

179 *Abnormal mineralization and malformations in the body end plates of precaudal vertebrae in*  
180 *adult zebrafish*

181 Vertebrae structures were analyzed by assessing key angles of the structure (Fig 3a) and  
182 mineralization features. MicroCT analysis of precaudal vertebrae identified abnormal patterns  
183 of mineralization and abnormal structures, that were especially pronounced at the arches and  
184 transverse processes in 1-year old *chd7<sup>-/-</sup>* and control zebrafish (Fig. 3b, c). Precaudal vertebrae  
185 showed significantly higher overall vBMD, while the distribution of mineralization between  
186 the body and arch remained unaffected (Fig. 3d, e). However, structurally, precaudal vertebrae  
187 of *chd7<sup>-/-</sup>* mutants show an increase of bone volume towards the body end plates of vertebrae,  
188 which in most cases show major malformations manifesting as highly mineralized inclusions  
189 (Fig. 3b, arrows). Additionally, vertebrae in 1-year-old *chd7<sup>-/-</sup>* zebrafish presented with a larger  
190 BV/TV compared to controls (Fig. 3g-i). Analysis of the vertebrae neural arch revealed  
191 distortion with a significantly larger rising angle, as well as a larger variance of the angles (Fig.  
192 3f). Vertebrae showed a significantly changed body angle and significant variance of body  
193 angle and vertebrae length (Supplemental Table 3). While a depletion of vBMD in the arches  
194 and abnormal vertebrae body structure was observed in 2-year-old fish, the effects seen on total  
195 vBMD and BV were not apparent in the few samples that survived until this stage  
196 (Supplemental Fig. 2 and Supplemental Table 4). Notably, one of our *chd7<sup>-/-</sup>* mutants showed  
197 vertebral fusion of the analyzed precaudal vertebra and following vertebra, which was not  
198 observed in any of the controls (Supplemental Fig. 1). This further demonstrates the wide  
199 variation in severity of the bone phenotype in *chd7<sup>-/-</sup>* zebrafish.

200 *Abnormal caudal vertebrae in chd7<sup>-/-</sup> zebrafish vary in arch angles and reduced mineralization*

201 We next determined key structural features (Fig. 4a) and mineralization characteristics of  
202 caudal vertebrae. The *chd7<sup>-/-</sup>* zebrafish presented with warped hemal and neural arches and  
203 abnormal body structure compared to controls (Fig. 4b). Similar to precaudal vertebrae, the



204 growth zones of the body end plates were enlarged and showed inclusions of highly mineralized  
205 matrix (Fig. 4b). Unlike control zebrafish, *chd7*<sup>-/-</sup> mutants had abnormal vertebrae body  
206 structure, which usually exhibited an hourglass shape and resulting in greater variance of  
207 measured body angles (Supplemental Table 3).

208 In line with the neural arch screened in the precaudal vertebrae, the hemal arch in the caudal  
209 vertebrae showed significantly wider arch rising angles (Fig. 4f), while the neural arch  
210 displayed significantly greater variance (Fig. 4g) in 1-year-old *chd7*<sup>-/-</sup> zebrafish. In contrast to  
211 precaudal vertebrae data, the caudal vertebrae showed significantly larger variance of Total  
212 vBMD and of both arches and vertebral body vBMD (Supplemental Table 3). A significant  
213 insufficiency in mineralization of the arches compared to the vertebrae body was also measured  
214 in *chd7*<sup>-/-</sup> zebrafish compared to controls (Fig 4 d, e). Further analysis also revealed a significant  
215 variance in BV, TV and BV/TV in *chd7*<sup>-/-</sup> zebrafish compared to controls (Fig h, i).

216 While significantly distorted arches were still observed in 2-year old mutants, the effects on  
217 vBMD and BV were not noticed in the few surviving fish that could be screened (Supplemental  
218 Fig. 3 and Supplemental Table 4).

#### 219 *Col2a1* of the ECM is depleted in *chd7*<sup>-/-</sup> zebrafish

220 To obtain a more complete analysis of vertebrae integrity, we analyzed the extracellular matrix  
221 (ECM) structures of the vertebral column in 9 dpf larvae and 1-year-old *chd7*<sup>-/-</sup> zebrafish and  
222 controls. Hematoxylin-Eosin (HE) staining of precaudal vertebrae sections showed a significant  
223 reduction in the number of nuclei measured in the vertebral cartilage of 1-year-old *chd7*<sup>-/-</sup>  
224 zebrafish and controls (Fig. 5a, b). More detailed analysis using Safranin/Fast Green O staining,  
225 revealed a significant reduction in the number of chondrocytes in the vertebral cartilage in *chd7*<sup>-/-</sup>  
226 zebrafish in line with the HE staining (Fig. 5c, d). Additionally, both stainings indicate a  
227 hypertrophic cell morphology in *chd7* mutants. We further screened for major components of  
228 the cartilage matrix. Gene expression analysis at 9 dpf revealed a significant downregulation of

229 the ECM related gene *col2a1a* (Fig. 5e). Immunofluorescence showed a striking depletion of  
230 Col2a1 in the vertebral cartilage with few cells remaining, which produce a sufficient Col2a1  
231 signal (Fig 5f). This pattern was effectively reproduced in all regions of the spine of *chd7*<sup>-/-</sup>  
232 zebrafish.

233

## 234 **Discussion**

### 235 *Efficiency of model*

236 CS is a multisystemic disorder likely involving a variety of underlying mechanisms that include  
237 CHD7. Therefore, different preclinical models are required to individually identify the various  
238 molecular mechanisms. Even though mouse models for CS exist, none have been reported to  
239 show the bone phenotype observed in CS patients. Our study presents a *chd7* deficient zebrafish  
240 that has a bone phenotype resembling that of CS patients, which we used to investigate skeletal  
241 development in CS. While scoliosis in CS patients has been attributed to poor muscular  
242 development, our study proposes an additional factor involving impaired early mineralization  
243 of vertebrae. In line with observations in patients, we observed spinal deformities in *chd7*<sup>-/-</sup>  
244 zebrafish, indicative of scoliosis.<sup>(10-12)</sup> Interestingly, our analysis reveals vertebral abnormalities  
245 are already present in young larvae of *chd7*<sup>-/-</sup> zebrafish. Additional clinical studies are required  
246 to investigate the onset of spinal deformities in CS patients. However, early analysis of bone  
247 density and spinal development has been suggested by CS guidelines.<sup>(45, 46)</sup> Detailed analysis  
248 may reveal less severe or early onset phenotypes. Similarly, we observed that adult *chd7*<sup>-/-</sup>  
249 zebrafish, which appeared without spinal deformities on visual inspection, showed signs of  
250 spinal abnormalities in microCT analysis, suggesting that mostly mild phenotypes reach this  
251 age. This is supported by the high lethality rate shown by Jamadagni et al. (unpublished), yet  
252 the underlying cause of lethality is still unclear.

253 Progressive skeletal disorders such as osteopenia and osteoporosis are characterized by a  
254 reduction in bone mineralization over time leading to increased fracture risk.<sup>(47)</sup> CS patients  
255 have been shown to present with significantly reduced bone mineral density.<sup>(13)</sup>  
256 Correspondingly, reduced vertebrae mineralization in *chd7*<sup>-/-</sup> zebrafish by 12 dpf was observed  
257 in a previous study involving morpholinos in zebrafish which had observed a reduced  
258 mineralization of vertebral structures.<sup>(27)</sup> Confirming previous data, we observed insufficient  
259 mineralization of vertebra along the posterior spinal column by 4 weeks of age. In earlier  
260 analysis of larvae at 6dpf, we were not able to detect a delay in the onset of mineralization, but  
261 already by 9 dpf we saw a significant decrease in the number of calcified vertebrae.  
262 Furthermore, mineralization appeared to be less efficient in posterior vertebrae, which could  
263 signify an impairment in ossification, since vertebrae in zebrafish undergo ossification from an  
264 anterior to posterior direction.<sup>(48)</sup> An exception from this rule is the Weberian vertebrae C1 and  
265 C2, which are ossified later in wildtypes but earlier in *chd7*<sup>-/-</sup> mutants.

#### 266 *Nutrient dependency*

267 Our study showed that the onset of calcification was not delayed in *chd7* deficient zebrafish.  
268 However, progressing calcification was decreased already 4 days post feeding onset, even  
269 though access to food was identical in control and mutants. In contrast, differences in  
270 calcification of vertebrae in control and mutants without access to food was not significant.  
271 Previous studies have shown that feeding delay until 9 dpf does not affect future fish growth  
272 and viability and was therefore our cut-off point.<sup>(49)</sup> This data suggests that possibly an  
273 underlying mechanism involving either reduced uptake or metabolizing nutrients may play a  
274 part in the observed phenotype and this link remains to be investigated. Bone calcification and  
275 mineralization is dependent on proper nutrition and is often discussed regarding optimal bone  
276 health in humans.<sup>(50)</sup> Insufficient levels of calcium and nutritional uptake are considered an  
277 increased risk for bone injury in CS patients.<sup>(11)</sup> To optimize nutritional uptake, the

278 supplementation of calcium and other nutrients to counteract poor bone health has been  
279 proposed for CS patients.<sup>(51)</sup> In fact, case reports show supplementation with calcium and  
280 vitamin D significantly increased aBMD in CS patients.<sup>(52)</sup> A possible nutrient deficiency in  
281 CS, could be linked to gastrointestinal nutrient uptake difficulties in *chd7*<sup>-/-</sup> zebrafish<sup>(53)</sup>, yet  
282 this needs to be further investigated. Thus, this possible deficiency in nutrient uptake could  
283 further hinder ossification in *chd7*<sup>-/-</sup> mutants. We propose using our zebrafish model of CS  
284 model to test the effect of guidelines such as calcium and nutrient supplementation, nutrient  
285 metabolism and mechanical loading in form of swim training to enhance bone strength in CS.

### 286 *Loss of chd7 reduces osteogenesis in zebrafish*

287 Osteoblast precursors commonly express differentiation markers such as *runx2* and  
288 *Osterix/sp7*.<sup>(54)</sup> Upon maturation of these osteoblasts, expression of mineralization related  
289 genes such as *bglap* (osteocalcin) are expressed. Analysis of these three genes showed that they  
290 were all significantly downregulated in *chd7*<sup>-/-</sup> larvae. This suggests a deficiency in mature  
291 osteoblasts, required for the mineralization of the spinal column. Misregulation or loss of these  
292 genes, such as *osterix/sp7* has been directly linked to poor bone mineralization in zebrafish and  
293 humans.<sup>(55-57)</sup> Notably, among all osteogenesis related gene expressions significantly reduced  
294 in *chd7*<sup>-/-</sup> larvae, *sost*, *ctsk*, *acp5* and *col2a1* have been identified as direct regulatory targets of  
295 Chd7 by Chip-Seq.<sup>(58)</sup> Furthermore, the regulation of osteogenesis has been linked to a  
296 regulatory complex with PPAR $\gamma$ .<sup>(23)</sup> Cell culture experiments have proposed the role of CHD7  
297 in osteogenesis by inhibiting PPAR $\gamma$  target genes by a protein complex involving CHD7, NLK  
298 and SETDB1. Upon loss of CHD7 this complex will fail to inhibit PPAR $\gamma$  target genes and  
299 therefore inhibit osteogenesis in preference over adipogenesis.<sup>(23)</sup> Strikingly, our qPCR results  
300 are consistent with downregulation of Ppar $\gamma$  target genes such as *runx2a* and *runx2b*.  
301 Furthermore, gene expression relevant for bone calcification and remodelling are misregulated

302 such as *sost* and *ctsk*. Thus, our study is the first to indicate the conservation of the proposed  
303 pathway shown in cell culture in an *in vivo* model.

#### 304 *Weberian apparatus*

305 The Weberian apparatus is a specialized structure that connects the auditory system and the  
306 swim bladder in a complex with the first 4 vertebrae to amplify sound vibrations.<sup>(33)</sup> Our results  
307 show that key structures in the apparatus fail to efficiently form and/or mineralize. Two of these  
308 structures, the tripus and the parapophysis, are formed to connect to the swim bladder. Both are  
309 malformed in all tested *chd7*<sup>-/-</sup> mutants to varying degrees. This malformation may imply an  
310 inefficiency in its function concerning both the swim bladder and auditory system. Ear  
311 abnormalities are a major symptom in CS patients and functionality is affected. Furthermore,  
312 the *chd7*<sup>-/-</sup> mutant shows swim bladder defects (Breuer et al., unpublished) that could be  
313 additionally affecting the morphology in tripus and parapophysis or vice versa. Even though  
314 the Weberian apparatus is extremely specialized for carp and carp-like fish (Ostariophysi), it is  
315 likely that it underlies the same pathways for osteogenesis as other parts of the spine.

#### 316 *Abnormal vertebrae and vBMD variability*

317 We examined volumetric bone mineral density (vBMD) and other parameters (e.g. BV, TV) in  
318 vertebral bone structures. Most strikingly, we detected large variance in density and all  
319 parameters examined, as well as some significant changes in both precaudal and caudal  
320 vertebrae. The large variance observed in *chd7*<sup>-/-</sup> mutants is in line with CS patients showing  
321 varying severity in spinal deformities.<sup>(12, 59)</sup> Furthermore, previous studies have shown that  
322 even minor changes in the described parameters can impact the risk for spinal deformities and  
323 fractures in disease and under treatment.<sup>(60, 61)</sup>

324 Another interesting feature we observed in most *chd7*<sup>-/-</sup> mutants, were highly mineralized  
325 inclusions, most commonly, localized towards the growth zone of the vertebrae. We did not

326 identify the source of these highly mineralized structures, but they could be caused by highly  
327 localized clusters of osteoblasts during vertebrae mineralization. Another possibility is the  
328 occurrence of calcified cartilage, which is more mineralized in comparison to bone. Although  
329 speculative, the occurrence of mineralized cartilage could be due to incomplete endochondral  
330 ossification, which would be in line with the reduction in *sp7/osterix* and *col2a1* expression<sup>(62,</sup>  
331 <sup>63)</sup> Since these inclusions are highly mineralized, it seems likely that their appearance impairs  
332 the bone structurally as higher mineralized collagen is stiffer, yet less tough.<sup>(64)</sup> It will be of  
333 interest to see if future studies can take a closer look at these previously unidentified features.  
334 While these minor defects could be involved in the structural integrity of the spine in CS  
335 patients, these minor features may be missed during dual-energy x-ray absorptiometry (DEXA)  
336 screens of CS patients when assessing aBMD.

337 Analysis revealed a decreased mineralization in the caudal vertebral arches of the mutant fish.  
338 This likely reduces the mechanical competence of these vertebrae in comparison to healthy  
339 vertebrae. A compromised mechanical competence alongside low BMD also increases fracture  
340 risk and can be an underlying cause of scoliosis.<sup>(65-67)</sup> Unfortunately, natural history BMD data  
341 of CS patients throughout early development is not available, to the best of our knowledge.  
342 However, aBMD analysis of patients with idiopathic scoliosis in CS via DEXA scans has been  
343 studied and shows significant reduction in some cases.<sup>(13, 68)</sup> In general, there is an increased  
344 fracture risk and incidence of scoliosis corresponding to decreased BMD, such as in osteopenia  
345 and osteoporosis.<sup>(47, 66)</sup> In line with this reduced mineralization an increased fracture and injury  
346 risk has been reported for CS patients.<sup>(13)</sup> Accordingly, proper observation and focus on early  
347 DEXA analysis is considered for CS guidelines.<sup>(45, 46)</sup>

348 Our analysis throughout development of *chd7*<sup>-/-</sup> deficient zebrafish suggests that only mild  
349 phenotypes survive onto later stages with severity decreasing as age progresses. However, the  
350 high lethality in both patients and our model is connected to a multisystemic phenotype in which

351 one mechanism is unlikely to be exclusive underlying cause for this characteristic. The link  
352 between *Chd7* and severe spinal and vertebrae abnormalities, including kyphosis, fused  
353 vertebrae and hemivertebrae has been observed in our samples.<sup>(7, 9)</sup>

#### 354 *Collagen and chondrocyte depletion as additional risk factor*

355 Our data in *chd7* deficient zebrafish indicate that reduced vBMD may not be the only  
356 underlying risk factor for idiopathic scoliosis in CS patients. Spinal integrity may be weakened  
357 by a decrease in chondrocyte marker *col2a1* expression in the vertebral cartilage of *chd7*  
358 deficient zebrafish. *Col2a1* is required in the ECM of various cell types including osteoblasts,  
359 chondrocytes, external ligament connective tissue cells, as well as notochord basal cells and  
360 expressed in the corresponding tissues.<sup>(69, 70)</sup> Each cell type is required for proper integrity and  
361 could likely contribute to spinal injuries and malformations. Furthermore, *col2a1a* expression  
362 is observed throughout ossification of the notochord.<sup>(71)</sup> Reduction in this collagen matrix in  
363 the notochord can affect proper calcification. *Col2a1a* is expressed in osteoblasts and  
364 chondrocytes of teleosts and regulated by *sox9*, which in turn is known to be regulated by  
365 *Chd7*.<sup>(72-74)</sup> The hypertrophic morphology of the chondrocytes seen in our study may indicate  
366 signs of a weakened matrix, as well as altered matrix composition and may contribute to an  
367 increased risk in osteoporosis.<sup>(75, 76)</sup> How this regulatory defect may influence other  
368 cartilage/skeletal regions such as the craniofacial cartilage remains to be investigated.

369 In summary, our study is the first to identify skeletal abnormalities linked to *chd7* deficiency.  
370 These abnormalities include morphological changes in vertebrae, reduced bone mineralization,  
371 impaired osteoblast activity, highly mineralized inclusions and *Col2a1* deficiency in the *chd7*  
372 <sup>-/-</sup> zebrafish model organism. The use of this preclinical model will allow for future studies to  
373 further understand the underlying mechanisms of spinal deformities in CS.

374

## 375 **Materials and Methods**

### 376 *Zebrafish husbandry*

377 Wildtype (Cntrl) and mutant (*chd7<sup>-/-</sup>*) zebrafish were kept at 28°C in a 12h/12h dark/light cycle  
378 and maintained in accordance with Westerfield et al.<sup>(77)</sup>. All zebrafish in this study were fed a  
379 steady diet of Skretting®GemmaMicro starting at 5dpf. Embryos were raised at 28.5°C and  
380 staged as previously described by Kimmel et al.<sup>(78)</sup> All experiments were performed in with the  
381 guidelines of the Canadian Council for Animal Care and the local ethics committee.

### 382 *Skeletal stainings and analysis of larvae*

383 Calcein staining was done as described<sup>(38)</sup>. In short, zebrafish larvae at 9dpf were exposed to  
384 calcein (2g/L) (Sigma-Aldrich) in water for 10 minutes. Larvae were then washed at least 3  
385 times in fish water for 10 minutes to remove excess calcein. Larvae were then anesthetized  
386 using tricaine before being imaged using an AxioZoom V16 (Zeiss). For nutrition dependency  
387 tests, larvae were either deprived of food until 9 dpf or fed a regular Skretting®GemmaMicro  
388 diet starting at 5 dpf.

389 Alizarin Red was performed as previously shown<sup>(79)</sup>. In brief, larvae were fixed for 2 hours with  
390 2% PFA. The PFA was then removed by washes in PBS 3x10 minutes and then placed in 25%  
391 glycerol/0.1%KOH. Finally, larvae were stained with 0,05% Alizarin Red (Sigma-Aldrich) in  
392 H<sub>2</sub>O for 30 minutes and stored in 50% glycerol in 0.1%KOH.

### 393 *MicroCT Imaging*

394 Adult zebrafish were euthanized using tricaine (MS-222, Sigma-Aldrich). Tissue was fixed in  
395 4% PFA prior to microCT analysis. Ex vivo microCT at an isotropic voxel size of 10.6 µm  
396 (SkyScan1276, Bruker, Kontich, Belgium) was performed to assess differences in bone mass,  
397 mineral density, and microstructure. The following scanning parameters were used: peak  
398 voltage of 55 kVp, Aluminum 0.25 mm filter, source current of 200 µA, 0.3° rotational steps



399 for full 360° with no frame averaging. Images were reconstructed using standard reconstruction  
400 algorithms provided with the microCT. Spinal deformities of controls and mutants were  
401 analyzed of 1-year (n=8 for Cntrl and n=8 *chd7*<sup>-/-</sup>) and 2-year-old (n=5 for Cntrl and n=5 *chd7*<sup>-/-</sup>)  
402 zebrafish.

#### 403 *Analysis and bone density/volume/angles*

404 From the reconstructed microCT images of the whole spine, the Weberian apparatus, as well as  
405 vertebrae 7 (abdominal) and 20 (caudal) were segmented. Further, the intercalarium, the tripus  
406 as well as the parapophysis were segmented from the Weberian apparatus. For the vertebrae,  
407 the neural as well as the hemal arch were segmented from the vertebral body. To differentiate  
408 between bone and background, a density-based threshold of 0.41 gHA/cm<sup>3</sup> determined by  
409 Otsu's method<sup>(80)</sup> was used. For each segmented bone in the Weberian apparatus outcomes  
410 included: bone volume BV (mm<sup>3</sup>), tissue volume TV (mm<sup>3</sup>), bone volume fraction BV/TV  
411 (mm<sup>3</sup>/mm<sup>3</sup>) and volumetric bone mineral density vBMD (gHA/cm<sup>3</sup>). For abdominal and caudal  
412 vertebrae the outcomes included: bone volume BV, tissue volume TV, bone volume fraction  
413 BV/TV, total bone mineral density T.vBMD, vertebral body vBMD VB.BMD, vertebral arch  
414 vBMD VA.BMD, as well as arch-body angle, body angle, arch opening angle and rising angles  
415 (see Fig.2a and 3a). For caudal vertebrae, the angle outcomes were calculated for hemal as well  
416 as neural arches.

#### 417 *qRT-PCR*

418 Total RNA was isolated from 9 dpf old zebrafish larvae using TriReagent. 1µg of RNA was  
419 used for cDNA synthesis using cDNA vilo kit. qRT-PCR was performed with SYBR Green  
420 mix (BIORAD) with a Lightcycler96® (Roche). Gene expression was analyzed relative to the  
421 housekeeping gene *elf1α*. Primers used for *runx2a*, *runx2b*, *sost*, *bglap*, *sp7/osterix*, *mgp*,  
422 *postna*, *acp5a*, *ctsk*, *col2a1* and *mgp* are shown in supplementals Table 5.

423 *Spinal sections/Stainings/Immunohistochemistry*

424 The zebrafish were euthanized in tricaine (MS-222, Sigma-Aldrich) and fixed for 3 days in 4%  
425 paraformaldehyde at 4°C, after the abdomen was opened to ensure proper fixation. The spinal  
426 cords were dissected and decalcified with EDTA 10 % for 3 days under agitation at RT. After  
427 fixation and decalcification, the zebrafish were embedded in paraffin. Longitudinal sections  
428 (5µm) were obtained and were deparaffinized in xylene and were rehydrated in a graded series  
429 of ethanol. The slides were stained with hematoxylin (STATLAB Medical Products, LLC) for  
430 4 min, and washed with alcohol-acid, and were rinsed with tap water. In the blueing step, the  
431 slides were soaked in saturated lithium carbonate solution for 10 sec, and then rinsed with tap  
432 water. Finally, staining was performed with eosin Y (STATLAB Medical Products, LLC) for 2  
433 min and mounted with Permount™ mounting medium.

434 For Safranin O/Fast green, after dehydration, the slides were stained with Weigert's  
435 hematoxylin (Sigma-Aldrich) for 4 min, wash under tap water, immersed in alcohol–acid for 5  
436 sec, then washed under water. The slides were stained with Fast green 0.02 % for 2 min, washed  
437 with acetic water 1% for 20 sec, then stained directly in safranin 0.01 % for 5 min and mounted  
438 with Permount™ mounting medium. Images were taken using an AxioZoom V16 (Zeiss) and  
439 area and number of nuclei were determined using ImageJ.

440 Immunofluorescence for Col2a1 was performed as previously shown<sup>69</sup>. In short, Epitope  
441 demasking was performed by digestion with Proteinase K (20µg/ml). Tissue was blocked with  
442 NGS and samples treated with primary antibody for Col2a1 (1 in 10; Developmental studies  
443 hybridoma bank) and secondary antibody Goat-anti-Mouse Alexafluor 488 (1 in 300). Counter  
444 stain was done using DAPI-mounting medium. Images were taken using a LSM780 (Zeiss).

445 *Statistics*

446 Statistical analysis was carried out using Prism-GraphPad® (GraphPad, San Diego, CA, USA).  
447 qPCR data and nuclei quantification were tested with student's t-test. Calcification analysis in  
448 larvae was performed with one-way ANOVA and post-hoc Tukey Test. MiroCT data of Control  
449 vs *chd7<sup>-/-</sup>* mutants was analyzed using student's t-test with Welch's correction and Variance  
450 determined by F-test analysis. Significance was determined at \* $<0,05$ ; \*\* $<0,01$  and \*\*\* $<0,001$

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#### 459 **Author roles**

460 MB, MR, BMW and SP designed the study. MB, MR and CZ performed the experiments. MB  
461 and MR drafted the manuscript. MB, MR, CZ, BMW, SP edited the manuscript.

462

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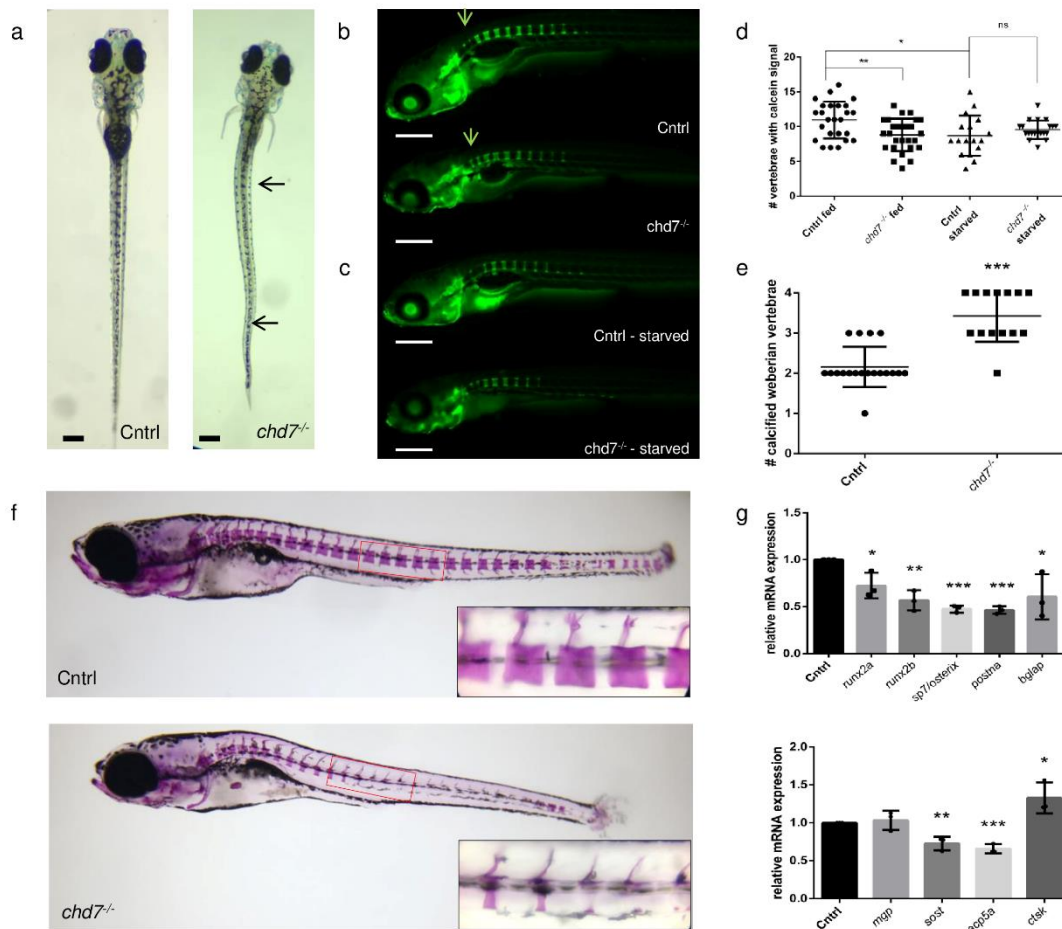
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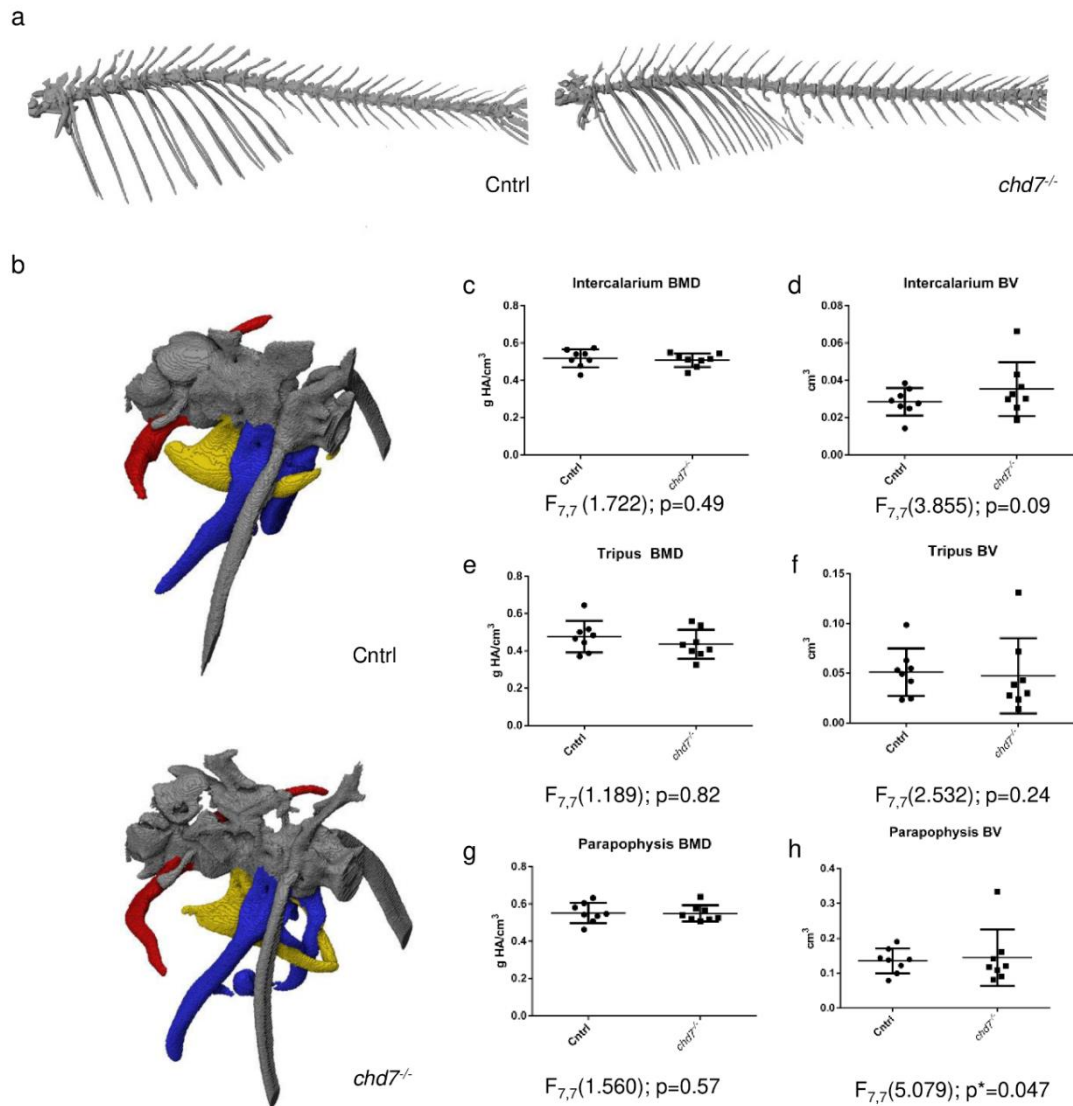


666

667 Fig 1. Mineralization in larval and juvenile *chd7<sup>-/-</sup>* mutants

668 a. Dorsal view of control and *chd7<sup>-/-</sup>* larvae at 9dpf showing severe spinal deformities in 43 out of 110  
 669 larvae screened. Deformities were present at both precaudal and caudal region (arrows). b. Lateral view  
 670 of calcein staining at 9dpf showing mineralization of vertebrae with access to food starting at 5dpf and  
 671 c. calcein staining at 9dpf during starvation. Arrow indicates mineralization at the first 4 (Weberian)  
 672 vertebrae d. Number of vertebrae in wildtype and *chd7<sup>-/-</sup>* (fed and starved) larvae showing at least partial  
 673 calcification signal as tested by calcein and e. Number of weberian vertebrae (vertebrae 1-4) showing  
 674 complete calcification (see arrows in b) f. Lateral view of 4 week old zebrafish stained with alizarin red  
 675 to screen mineralized vertebrae, showing decreased calcification towards posterior region. Bottom right  
 676 corners show magnification of selected caudal vertebrae. g. RT-qPCR of osteogenesis related genes  
 677 screened at 9dpf, showing relative fold change. (Significance: \* $p < 0.05$ ; \*\* $p < 0.01$ )





678

679 Fig 2. Weberian apparatus

680 a. Representative microCT overview image of the spinal cord from a 1-year old control and a *chd7<sup>-/-</sup>*

681 adult zebrafish b. MicroCT Image of the Weberian apparatus from a 1-year old control and a *chd7<sup>-/-</sup>*

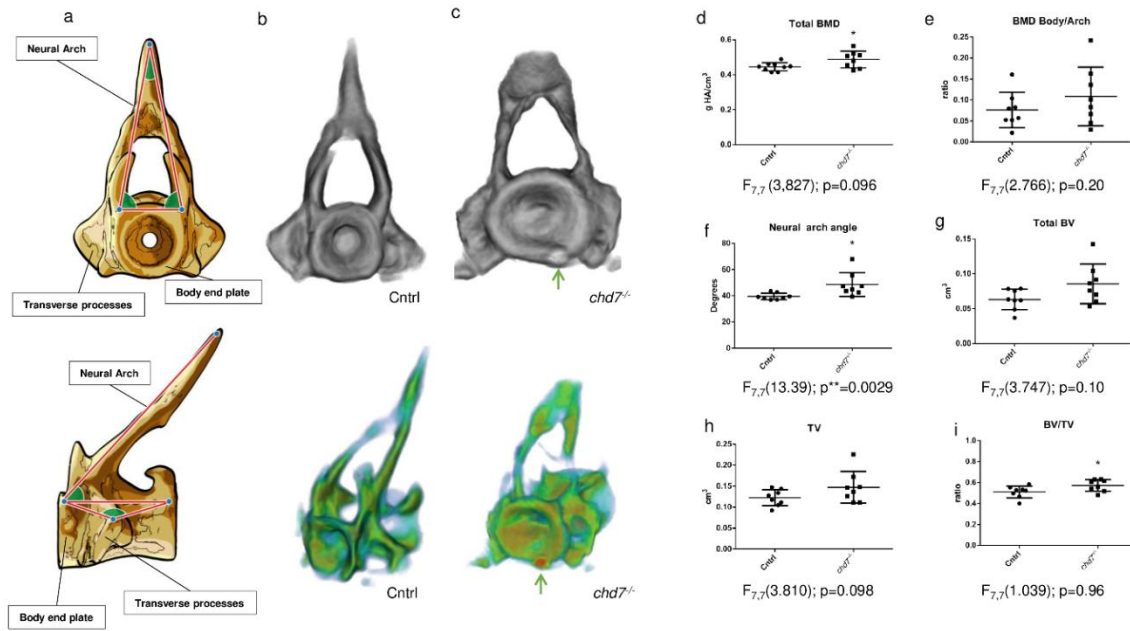
682 mutants, indicating structures of intercalarium (red), tripus (yellow) and parapophysis (blue). c-h.

683 vBMD and BV/TV of intercalarium (c, d) tripus (e, f) and parapophysis (g, h) (n=8/genotype were

684 analyzed). Significance of student t-test with Welch's correction is included in the graphs with: \*p<0.05;

685 \*\*p<0.01. F-test results are indicated underneath as  $F_{\text{degree of freedom numerator, degree of freedom denominator}}$  (F-value);

686 p-value: \*p<0.05; \*\*p<0.01; \*\*\*p<0.001



687

688 Fig 3. Precaudal vertebrae show inefficiency for proper mineralization

689 a. Sketch diagram of precaudal vertebrae (frontal and lateral) indicating structure and measured angles.

690 b-c. (top) Individual rendering of precaudal vertebrae, frontal view of tissue over 0.41 g HA/cm<sup>3</sup>

691 threshold and (bottom) vBMD intensity map (Range blue-red; 0.41 – 1.00 g HA/cm<sup>3</sup>) showing

692 morphological abnormalities and growth zone malformations with highly mineralized inclusions

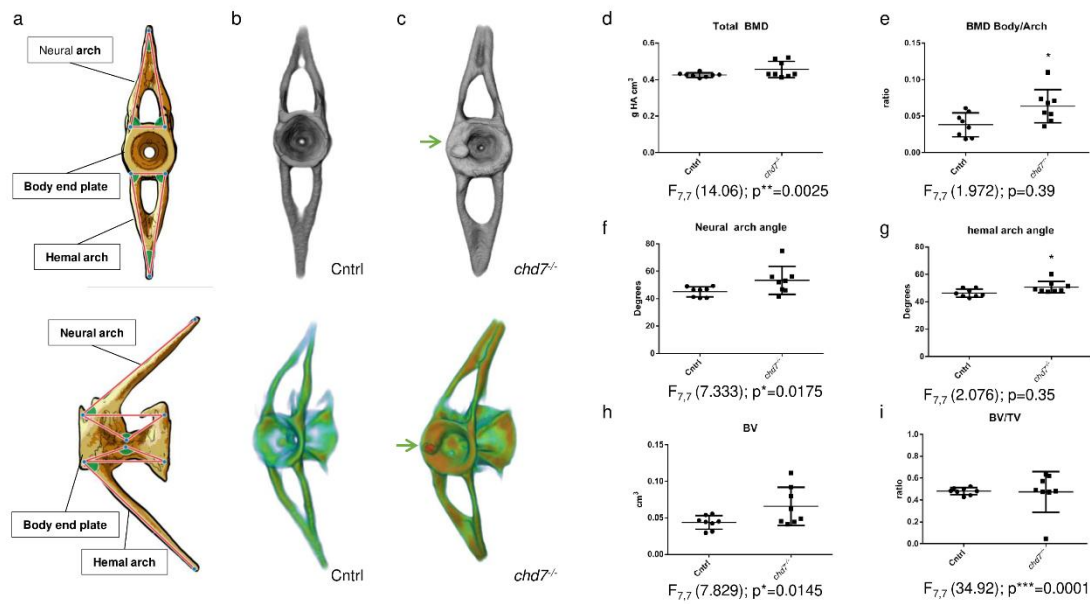
693 (arrow) d. vBMD of whole vertebrae showing increasing density with increasing age and e. increasing

694 ratio of vBMD arch/vertebrae body. f-h neural arch angle, BV, TV and ratio BV/TV. (n=8/genotype

695 were analyzed). Significance of student t-test with Welch's correction is included in the graphs with:

696 \*p<0.05; \*\*p<0.01. F-test results are indicated underneath as F<sub>degree of freedom numerator, degree of freedom denominator</sub>

697 (F-value); p-value: \*p<0.05; \*\*p<0.01; \*\*\*p<0.001



698

699 Fig 4. Caudal vertebrae show abnormal structure

700 a. Sketch diagram of precaudal vertebrae indicating structure and measured angles (frontal and lateral).

701 b-c. (top) Individual rendering of caudal vertebrae, frontal view of tissue over 0.41 g HA/cm<sup>3</sup> threshold

702 and (bottom) vBMD intensity map (Range blue-red; 0.41 – 1.00 g HA/cm<sup>3</sup>) showing morphological

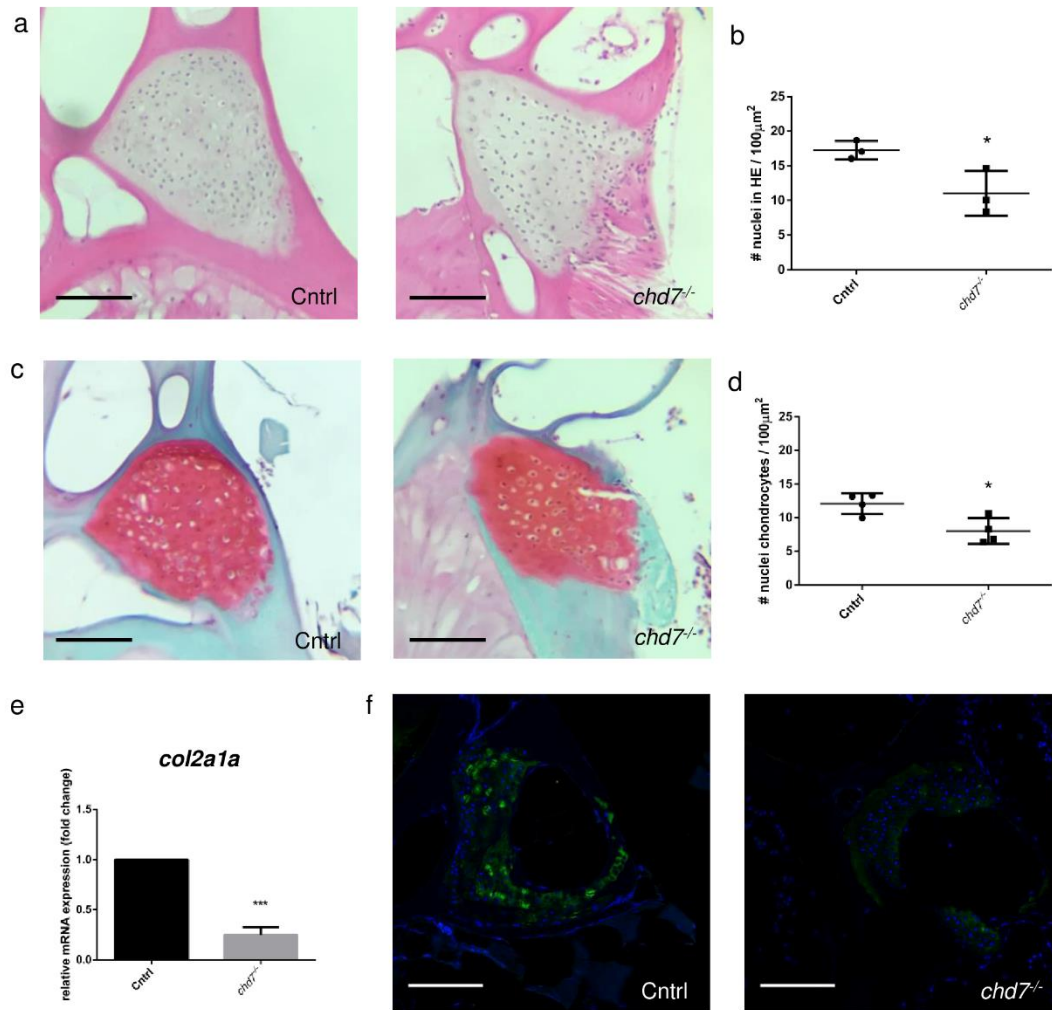
703 abnormalities and growth zone malformations with highly mineralized inclusions (arrow) d. vBMD of

704 whole vertebrae and E. increasing ratio of vBMD arch/vertebrae body. f-h neural arch angle, BV, TV

705 and ratio BV/TV. (n=8/genotype were analyzed). Significance of student t-test with Welch's correction

706 is included in the graphs with: \*p<0.05; \*\*p<0.01. F-test results are indicated underneath as F<sub>degree of</sub>

707 freedom numerator, degree of freedom denominator (F-value); p-value: \*p<0.05; \*\*p<0.01; \*\*\*p<0.001



708

709 Fig 5. ECM Collagen deficiency in 1-year-old *chd7*<sup>-/-</sup> adults

710 a. H&E staining of precaudal vertebrae section of Cntrl and *chd7*<sup>-/-</sup> mutants b. Total nuclei count in  
711 precaudal vertebral cartilage detected in H&E staining per 100µm<sup>2</sup> c. Safranin O/fast green staining of  
712 precaudal vertebrae section Cntrl and *chd7*<sup>-/-</sup> mutants showing cartilage in red. d. Chondrocyte nuclei  
713 count in precaudal vertebral cartilage detected in Safranin O/fast green staining per 100µm<sup>2</sup> e. qPCR of  
714 *col2a1a* at 9dpf. f. Immunofluorescence of Col2a1 in precaudal vertebrae sections of Cntrl and *chd7*<sup>-/-</sup>  
715 mutants with Col2a1 in green and DAPI in blue. Scale bar presents 100µm. Significance of student t-  
716 test with: \*p<0.05; \*\*p<0.01 \*\*\*p<0.001