Post-embryonic development and aging of the appendicular skeleton in *Ambystoma mexicanum*

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Abstract

Background

The axolotl is a key model organism for studying appendicular regeneration in vertebrates. The axolotl limb complexity resembles that of humans in structure and tissue components; however, axolotl limbs develop post-embryonically. In this work, we evaluated the post-embryonic development of the appendicular skeleton and its changes with aging.

Results

Juvenile skeletal elements are formed mostly by $SOX9^+$/$COL1A2^+$ cartilage cells. Ossification of the appendicular skeleton starts when animals reach a total length of 10 cm, and cartilage cells are replaced by a primary ossification center, consisting of cortical bone and an adipocyte-filled marrow cavity. Vascularization is associated to the ossification center and to the formation of the marrow cavity. We identified the contribution of $Col1a2^+$ cells to bone and adipocytes. Moreover, ossification progresses with age towards the epiphyses of long bones. Even though axolotls are neotenic salamanders, ossification remains responsive to L-thyroxine, increasing the rate of bone formation.

Conclusions

Bone maturation is a continuous process in axolotls that extends throughout the life span of the axolotls. Ossification of the appendicular bones is slow and continues until the complete element is ossified. The cellular components of the appendicular skeleton change accordingly during ossification, creating a heterogenous landscape in each element.

Introduction

The axolotl (*Ambystoma mexicanum*) is a gold standard for studying limb regeneration

in vertebrates. The progenitor cells in the blastema, a key structure for epimorphic

regeneration, are a heterogeneous population that remains restricted to their embryonic

origin (Kragl et al., 2009). Given the cellular heterogeneity of the blastema, the contribution

of specific cell populations during regeneration has been the focus of significant research in

the last decade. An interesting exception is the skeleton: cells embedded in the skeletal matrix

do not participate in regeneration, instead, the skeleton is restored by periskeletal cells and

dermal fibroblasts (Currie et al., 2016; Gerber et al., 2018; McCusker et al., 2016).

While mammals are born with ossified bones, ossification in most amphibians is

coupled with metamorphosis (Miura et al., 2008; Quilhac et al., 2014; Rozenblut and Ogielska,

2005). However, axolotls remain neotenic and, hence, they maintain juvenile features

throughout life and only rarely undergoing metamorphosis. Their adulthood is marked by

reaching sexual maturity, slowing growth rate and by ossification of the appendicular skeleton

(Malacinski, 1978; Vieira et al., 2019). The physiological context in which appendicular

ossification occurs in axolotl as well as the cellular transitions within the skeleton are unclear.

In the anuran amphibium, *Xenopus laevis*, the extent of ossification negatively correlates with

the regenerative potential of the limb. During metamorphosis and as bone ossifies,

regeneration declines in the ossifying areas, while amputation at the cartilaginous joints still

regenerates (Wolfe et al., 2000). Limb regeneration in salamanders remains a feature of adult

animals; however, regeneration also declines with age and after metamorphosis (Monaghan

et al., 2014). It remains unclear if the correlation between appendicular ossification and

decreased regeneration holds true for the axolotl, but more importantly, if the regeneration

mechanisms studied so far are replicated when the cellular and matrix landscape changes in

the skeleton. In this study, we analyzed larvae, adult and aged axolotls to document the

transition of the appendicular skeleton from cartilaginous to ossified skeleton.

In general, appendicular bones develop by endochondral ossification, a process where

a cartilage anlage is replaced by bone (reviewed in (Kozhemyakina et al., 2015; Olsen et al.,

2000)). During development, mesenchymal progenitors condensate and form a cartilage

primordium which expands by proliferation of chondrocytes. Cells located in the central

diaphysis differentiate into hypertrophic chondrocytes (HCs), which subsequently induce the

recruitment of blood vessels, osteoclasts and osteoblasts, giving rise to the primary
ossification center. Within the diaphysis, the cartilage matrix is degraded, osteoblasts replace
the tissue with cortical bone and a marrow cavity is formed. Simultaneously, osteoblasts
located in the perichondrium form a bone collar around the diaphysis and the periosteum is
established. Many HCs undergo apoptosis during endochondral ossification; however, several
studies have shown a partial contribution from HCs to bone formation through
transdifferentiation into osteoblasts (Giovannone et al., 2019; Jing et al., 2017; Yang et al.,
2014). Moreover, Giovannone *et al* showed that HCs can also transdifferentiate into
adipocytes in the ceratohyal bone in zebrafish, highlighting the importance and participation
of HCs in the ossification process. In axolotl, however, endochondral ossification is a post-
embryonic process. If it is driven by the hormonal changes of sexual maturity, environmental
cues or body mass, is not yet determined.

In this work, we evaluated the post-embryonic development of the appendicular
skeleton of axolotls, from larvae to aged adults. We analyzed the morphological and cellular
changes in the zeugopodial elements, the radius and ulna, and we identified the timing in
which ossification of those elements occurred. Moreover, we examined the cellular landscape
of radius and the possible contribution of chondrocytes to bone formation in ulnas. Finally,
we observed in aged animals (5- to 20-years) a continuous ossification in all limb skeletal
elements. Hence, our work provides a detailed view of how the appendicular skeleton,
particularly the zeugopod, matures from a cartilage anlage towards a bone. Urodeles are the
only vertebrates with the ability to regenerate a whole limb as adults, therefore they offer the
unique opportunity to gain important insights to advance regenerative medicine.

Results

The appendicular skeleton is progressively ossified with growth

Few studies on aged animals beyond the well-determined larval stages (Schreckenberg and Jacobson, 1975) and the neoteny of the axolotl, raises controversy on the limit of its growth. In this study, we used animals grown in fully standardized conditions (Khattak et al., 2014) and we defined their appendicular development in relation to their maturation stage. We created a growth chart (normogram) and observed a correlation between size and age for both total length (TL) and snout to vent length (SV) (Fig. 1A, B). Both measurements presented a similar trend: a rapid growth phase in the first 24 months of age followed by a stationary phase. We documented the growth in the context of sexual maturity in our axolotl colony. The first signs of secondary sexual features in males appear around 10 months of age, and by 12 months these features are fully developed. However, a successful mating starts only between 11 and 12 months of age. Females develop the first secondary sexual features slightly later than males (between 10-11 months of age) but like males, by 12 months these features are fully developed. Fertility is first successful in females between 12-14 months-old

Indeed, in 1-year-old siblings, TL was significantly longer in males than females (males: 20.06 ± 0.6 cm, females: 17.78 ± 0.4 cm, n = 9), while SV remained constant (males: 9.8 ± 0.3 cm, females: 9.3 ± 0.3 cm, n = 9) (Fig 1C). In 2.5-month-old juvenile larvae siblings, TL did not differ (males: 4.7 ± 0.2 cm, females: 4.7 ± 0.2 cm, n = 12). This indicates that the differences in total length between sexes is due to an increased tail growth in males upon sexual maturation.

We then evaluated the post-embryonic development of the appendicular skeleton, particularly the long bones from the zeugopod. We used animals ranging from TL 4 to 20 cm, which represent axolotls from a juvenile stage to adulthood. Using alcian blue/alizarin red staining, we first evaluated the broad morphological changes in the skeleton. Alizarin red is a calcium-binding dye that is routinely used to detect ossification patterns together with alcian blue, which stains polysaccharides in cartilage. We observed a progressive expansion of the alizarin red staining in the skeletal elements, which starts in axolotls with a TL around 10-12 cm (Fig 1D). Still, the epiphyses of those elements remained cartilaginous in animals of TL 20 cm. Importantly, alizarin red also labels mineralized cartilage, which can be seen *in vivo* in axolotls by fluorescence imaging (Fig 1E), hence it does not provide accurate information on

the ossification dynamics. To accurately quantify ossified bone volume and trace the start of the ossification process, we analyzed different limbs by µCT scan. Ossification was not detected in animals of TL 10 cm. We observed the first appearance of an ossified ring around both the radius and ulna in animals with a TL of 12 cm (Fig 1F). In animals bigger than 12 cm, ossification extended longitudinally to both ends of the bone. Interestingly, the ossification did not occur homogenously, but rather in a discontinuous manner. A porous appearance resembles the mammalian bone development. In animals of TL ca 25 cm, this porosity disappeared and the tissue became uniformly ossified (Fig 1H), forming compact bone. Ossification extended to the epiphyses of the bone, as observed in the widening ends. Quantification of the bone volumes (Fig 1G) are not significantly different between bones from animals TL 12 and 14 cm, or TL 14 and 16 cm. However, between TL 16 cm and 20 cm, a 4-fold increase in bone volume was observed for radius, and 7-fold for ulna. Moreover, radius volume was significantly higher than ulna (1.4 times increase) in axolotls of TL 20 cm. Even though ossification started when animals had a TL of 12 cm, there is variability between siblings, specially at TL of 14 cm (Fig 1G). Whether this variability results from sexual dimorphism or simply interindividual variation is yet to be determined.

Axolotl bones are vascularized and the marrow cavity is filled with adipocytes

Changes in the skeletal cellular landscape accompany the transition from a cartilage anlage to bone. We assessed these changes in isolated radii from axolotls TL 6 and 12 cm, which corresponded to radii before and during ossification. We performed H&E staining from paraffin sections and observed that the radius was exclusively composed of chondrocytes in animals TL 6 cm (Fig 2A, left panel). We identified a resting zone (RZ), filled with rounded chondrocytes, a proliferative zone (PZ), characterized by flattened chondrocytes, and a hypertrophic zone (HZ), where chondrocyte differentiation leads to an increase in cellular volume. Interestingly, these zones were not uniformly organized, as opposed to the isotropic cellular distribution of mammalian growth plates. The lack of an organized cellular distribution has been previously observed in another salamander model (Quilhac et al., 2014). Throughout the juvenile stages, a calcification ring surrounds the perichondral region of the HZ (arrowhead Fig 2A', 3A). We have observed that the first calcification ring appears in development occurred at stage 48 (data not shown). At this stage, the humerus is formed and the radius and ulna had just started to form by bifurcation from the humerus distal end.

Radii from animals TL 12 cm (Fig 2A, right panel) have a similar cellular distribution of

the cartilage region as younger animals. But in contrast to young animals, we observed the

formation of a primary ossification center in the diaphysis, which contained a marrow cavity

surrounded by cortical bone. The woven-like bone seen on the µCT scan was also observed in

sections as discontinuous tissue surrounding the cartilage, with defined edges between gaps

(Fig 2A''). The bone extended beyond the marrow cavity, surrounding also the HZ. By TL 16

cm, radii stained with H&E showed a further progression in ossification. An expanding marrow

cavity and compact bone were also observed. However, cartilage lining subjacent to the bone

could still be seen suggesting that the transition to bone was not completed. Embedded

osteocytes in the cortical bone matrix and cells lining in the outer and inner sides of the bone

were observed (periosteum and endosteum respectively) (Fig 2B). In both TL 12 and 16 cm,

the marrow cavity was filled with adipocytes identified by their morphology.

A critical step for bone formation by endochondral ossification is the vascularization

of the tissue (Filipowska et al., 2017). To assess the vascularization of the skeletal elements

from animals TL 6 cm and 12 cm, we generated a new transgenic line that expresses EGFP

under the control of a *Tie2* promoter and enhancer (*Tie2-Tg*) (Schlaeger et al., 1997), labeling

blood vessels in the entire animal. We collected limbs from animals TL 6 cm and the radii from

animals TL 12 cm and performed whole-mount immunofluorescence (IF) against GFP,

followed by tissue clearing and confocal imaging. We observed that the blood vessels

surrounded the skeletal element of TL 6 cm animals, but the cartilage was not vascularized

(Fig 2C, left panel). However, we observed some blood vessel protrusions in close contact with

the perichondrium in the HZ (arrowhead, Fig 2D), which could point to the position where the

ossification of the skeletal element will start. In the radii of TL 12 cm animals (Fig 2C, right

panel), the mid-diaphysis was found to be vascularized, which correlates with the ossification

of the cartilage anlage and formation of the marrow cavity. We did not observe vascularization

of the epiphyses, and thus no secondary ossification center formation at this stage.

Sox9 labels chondrogenic cells while Col1a2 labels all skeletal cells

We have identified the morphological changes of the zeugopod skeleton during

growth. However, the identity of the cells within the skeletal tissue, and their origin, remains

unclear. A key transcription factor involved in cartilage development is Sox9 (Kozhemyakina

et al., 2015). Thus, using a *Sox9-Tg* knock in line, where a T2a self-cleaving peptide followed

by mCherry protein is fused to the endogenous SOX9 protein, we evaluated the distribution

of SOX9$^+$ cells in ulnas from animals TL 6 and 12 cm (Fig 3A). We confirmed that the expression

of mCherry mirrors the real expression of Sox9 by immunofluorescence (Fig 3C). In tissue

sections from TL 6 cm ulnas, most cells were mCherry$^+$, while perichondral cells were mCherry$^-$

(Fig 3A, left panel). We observed a similar distribution in ulnas from animals TL 12 cm (Fig 3A,

right panel), and we found no mCherry$^+$ cell in the ossified portion of the skeletal element (Fig

3D). Moreover, using a *Col1a2-Tg* line, we evaluated the distribution of COL1A2$^+$ cells in ulnas

from animals TL 6 and 12 cm (Fig 3B). This transgenic line expresses fluorescent protein TFP

under the control of the *Col1a2* enhancer and has been shown to label skeletal cells in the

axolotl (Gerber et al., 2018). We observed TFP$^+$ cells throughout the skeletal elements in both

sizes, labelling both cartilage and bone cells (Fig 3B, E), i.e. chondrocytes, perichondral cells,

osteoblasts, osteocytes and periosteal cells.

Osteocalcin is expressed in bone cells and in some hypertrophic chondrocytes

The most definite confirmation of ossification is the identification of mature

osteoblasts in the bone matrix and bone lining cells. To that end, we used immunodetection

of osteocalcin (anti-OCN). As expected, no labeling was observed in ulnas from TL 6 cm (Fig

4A, left panel), which further demonstrated the absence of bone tissue in juvenile larvae. In

TL 12 cm animals, we observed a broad OCN labeling of the bone ECM (Fig 4A, right panel).

We identified OCN$^+$ cells in the skeletal element both embedded in the bone matrix and in the

periosteum (arrowheads, Fig 4B). Interestingly, OCN$^+$ cells were also identified in the lower HZ

of the chondro-osseous junction of the mid-diaphysis. Those cells were mCherry$^+$, identified

by our *Sox9-Tg* line (solid arrowheads Fig 4B, 4C). Previous studies have correlated the

downregulation of *Sox9* in the HZ with the activation of the osteogenic program and the

persistence of SOX9 with an inhibition of bone-related markers (Dy et al., 2012; Lui et al.,

2019). The expression of OCN in HCs could point towards an early activation of an osteogenic

program in those cells. In that regard, osteocalcin expression has been reported in post-HCs

(Pullig et al., 2000). Some studies have demonstrated the ability of HCs to transdifferentiate

into osteoblasts in mammals and zebrafish (Giovannone et al., 2019; Yang et al., 2014).

However, the co-expression of SOX9 and OCN needs to be further explored.

Lineage tracing reveals a contribution of *Col1a2$^+$* cells to bone and adipocyte tissue

Our above observations led us to hypothesize that some HCs could transdifferentiate
into bone cells in the axolotl. To test this idea, we evaluated the contribution of *Col1a2$^+$*
chondrocytes to bone formation.

Using the double transgenic *Col1a2xLPCherry-Tg*, we traced the fate of *Col1a2$^+$* cells
by inducing the permanent labeling of chondrocytes by tamoxifen-induced conversion of the
Cre/Lox reporter cassette in TL 5 cm animals. In these animals the appendicular skeleton is
still exclusively cartilaginous. We allowed the animals to grow until a TL of 14 cm and we
collected the ulnas for sectioning and imaging. We observed a great proportion of the cells to
be mCherry$^+$. They were distributed through the cartilage part, bone matrix, periosteum,
endosteum and in the adipose tissue (Fig 4D, 4E). Moreover, we assessed whether some of
the mCherry$^+$ cells were OCN$^+$. We found mCherry$^+$/OCN$^+$ cells in the endosteum and
periosteum (arrowheads, Fig 4F, 4F', 4F''), showing that *Col1a2$^+$* descendants gave rise, at
least partially, to committed osteoblast in the axolotl zeugopod. However, whether a specific
subpopulation of *Col1a2$^+$* gives rise to all derived cells or if HCs contribute directly to OCN$^+$
cells, cannot be answered with this transgenic approach.

Bone ossification and marrow cavity expand with age
Axolotls have a life span extending to more than 20-years under controlled laboratory
conditions (Vieira et al., 2019). However, little is known about bone morphological features in
aged animals, although it is commonly assumed that axolotls retain cartilaginous epiphyses
and thus there is perennial longitudinal growth of the bones. We collected bones and limbs
from 1-, 5-, 7- year-old axolotls and our only available specimen of 20-years of age. In
histological sections from radii (Fig 5A), we observed an age-related expansion of the marrow
cavity towards the epiphyses. Cortical bone was also expanded longitudinally and radially,
surrounding most of the skeletal element length, even though some parts of the inner surface
of the epiphyses remained cartilaginous in most cases. However, when we sectioned the 20-
year-old radius (n = 1), we observed a partial ossification of the epiphysis (arrowhead, Fig 5A),
which suggests a continuous ossification of the radius with age. Importantly, we did not
identify a secondary ossification center in any of the radii analyzed. Still, we observed the
appearance of pseudo trabeculated bone tissue, which we believe corresponded to the
marrow cavity expansion. In addition, in whole limbs stained with alcian blue/alizarin red, we
observed a calcification in one of the carpal bones in a 5-year-old animal (Fig 5B), as well as

several tarsal bones in a 20-year-old animal (Fig 5C), which suggest that the short bones in the autopod undergo a late ossification in comparison with the long bones in the zeugopod, but also highlights the continuity of this process in adulthood and aging.

In elements from the stylopod and autopod (digits from 2-year old animals and humeri from 5-year-old animals) (Fig 5D, E respectively), we observed the same features described for long bones in the zeugopod, i.e., an ossified ring of cortical bone around a marrow cavity filled with adipocyte tissue which expanded from the diaphysis towards the epiphyses.

This continuous ossification could be the origin of aberrant phenotypes. In 3 out of 6 bones from 7-year-old and in our single 20-year-old animal, we observed that the zeugopodial bones were fused either in their proximal or distal epiphysis. In one case, the fusion occurred with a heterotopic ossification and cartilage formation around the distal end of the radius.

Our results show a continuous ossification of the skeletal elements even in aged animals. Whether the appendicular skeleton would become completely ossified if an animal lived long enough remains unknown.

Axolotl metamorphosis accelerates bone ossification

Metamorphosis is a critical transformation in the life cycle of many amphibians. This process allows them to access a vaster territory in their search for food and mating partners. In some anurans and urodeles, ossification of the limb skeleton is associated with the metamorphic transformation (Miura et al., 2008; Quilhac et al., 2014). Although axolotls are neotenic and rarely undergo spontaneous metamorphosis, the administration of exogenous thyroxine can induce it. We evaluated whether the ossification in axolotl is still responsive to the hormonal changes of metamorphosis. If this is the case, metamorphosis should accelerate the ossification process in axolotls. We injected thyroxine in axolotls TL 14 cm, where ossification is already covering the mid-diaphysis. Radii were collected at 35 days post-injection and embedded in paraffin for sectioning and staining. Metamorphic axolotls were slightly smaller than their paedomorph siblings, although this difference was not significant (paedomorph TL: 15.4 ± 0.4 cm, SV: 8.6 ± 0.3, n = 5, metamorphic TL: 14.7 ± 0.3, SV: 8.4 ± 0.5, n = 6). By H&E staining, we did not observe differences in the gross morphology (Fig 6A) compared to an un-injected animal. These observations are in agreement with a previous study, in which no significant difference was observed in zeugopodial bones when surface and length were compared between a pre- and post-metamorphic axolotl (Thampi et al., 2018).

However, when compared using µCT scan, we observed an increase in bone volume in
metamorphic animals. Both zeugopodial bones had a bigger ossified area and the ossification
pattern was more uniform than the paedomorph bones analyzed (Fig 6B). Bone volume
quantification showed a significant increase of 4 times in radii volume and over 3 times
increase for ulnas (Fig 6C). These results demonstrate that ossification is accelerated in axolotl
limbs when metamorphosis is induced, and suggest that cells respond to intrinsic programs as
well as hormonal changes.

Discussion

Appendicular skeletogenesis is considered highly conserved among tetrapods, mostly based on studies in mammals and chicks, (Shubin et al., 1997) although the appendicular skeleton has undergone species-specific adaptations due to the environmental influence. Many species display variations in important cellular and temporal parameters of limb development. Mammals are born with ossified bone, while most amphibians develop their limbs post-embryonically. The transition of a cartilaginous appendicular skeleton to a bony skeleton is associated with the hormonal changes during metamorphosis (Kemp and Hoyt, 1969; Rose and Reiss, 1993). Metamorphosis in amphibians is a significant event that precedes sexual maturity, giving the animals a biological advantage. In neotenic salamanders, such as the axolotl, the appendicular cartilage is not only a transitional tissue, and larvae live with a calcified cartilage throughout their juvenile period. Their limbs ossify in the absence of metamorphosis although it was proposed that the epiphyses remain cartilaginous throughout life (Haines, 1942; Thampi et al., 2018) and that this feature would likely contribute to longitudinal bone growth (Cosden-Decker et al., 2012). However, little can be said about the biological clock triggering skeletal changes, since they rarely change to a terrestrial environment.

In this work, we provide a characterization of the maturation of the appendicular skeleton during the life span of the axolotl, considering ossification timing and age-related effects, complementing previous studies in larval stages (Torok et al., 1998) and young adults (Cosden-Decker et al., 2012). In our axolotl colony, animals are grown in fully standardized conditions that allow us to use the total body length to associate it to appendicular changes and biological landmarks such as sexual maturity. Our results show an exponential growth of axolotls in the first 22-24 months of their life, after which growth plateaus. During this exponential growth, animals become sexually mature and ossification of their appendicular bones starts, but for most of the juvenile phase, axolotls present a mineralized ring around the cartilaginous mid-diaphysis. Mineralization of the zeugopodial appendicular skeletal elements was observed as early as developmental stage 48. In contrast, ossification starts around the sexual maturity period, when we can observe the formation of a denser calcified surface and embedded cells with morphological and genetic characteristics of osteoblasts. Moreover, we show that $Col1a2^+$ cells or their progeny participate in the ossification process

by giving rise to osteogenic cells and adipocytes. We report a continuous ossification of

zeugopodial bones and marrow cavity expansion with age. However, this ossification is not

complete, as we observed a cartilage-bone composite in most of diaphyses analyzed. Short

bones, such as tarsals and carpals, undergo a very late ossification when compared to

appendicular long bones. Finally, we see an accelerated zeugopodial ossification in

metamorphic animals, indicating a responsiveness to exogenous thyroxine.

Mineralization *versus* ossification

For most of the initial larval stages, the skeleton is cartilaginous with mineral

deposition. We have observed calcification of the humeral diaphysis as early as Stage 48, when

radius and ulna had just started to form by bifurcation from the humerus distal end. Nye *et al*.

reported mineralization by alizarin red staining at stage 50, when condensations of the

metacarpals are already present (Nye et al., 2003). This discrepancy can be accounted for by

the detection protocol. While Nye *et al.*, used processed samples, we used real-time vital

detection. The calcification of appendicular cartilage remains throughout the juvenile phase

of the axolotl until the anlage starts to ossify.

In contrast to mineralization, ossification occurs after the limbs are fully formed and

around the time animals begin to show secondary sexual characters. Generally, amphibian

metamorphosis precedes sexual maturation, while in neotenic amphibians both sexual

maturation and bone ossification occur in the absence of metamorphosis. In salamanders, as

in other species, metamorphosis provides adaptive features to change habitat and ensure

greater chances of survival. The axolotl and its geographically and phylogenetically related

counterpart, *Ambystoma velasci* (Parra-Olea et al., 2007; Shaffer and McKnight, 1996), are

examples of facultative metamorphs, which means that individuals transform into terrestrial

organisms when the environmental conditions of its habitat change. In *Xenopus laevis*

tadpoles lacking thyroid glands, metamorphosis is halted but growth continues and the

ossification progresses in the hind limb independently of thyroxine (Kerney et al., 2010). This

suggests that appendicular ossification programs could also be triggered in the absence of L-

thyroxine and that probably depends on a critical body mass. Although ossification genetic

programs in axolotl are triggered independently of metamorphosis, inducing metamorphosis

with exogenous thyroxine, accelerates the ossification process as shown in this work. Thus,

cells within the appendicular skeleton remain responsive to changes triggered by thyroxine.

In contrast to the fully ossified diaphysis of post-metamorphic limb newts (*Pleurodeles waltl* total length 14 cm) (Quilhac et al., 2014), the diaphysis of axolotls retains a cartilage anlage subjacent to the cortical bone even in old animals and also in post-metamorphic zeugopodial and autopodial elements. We think this observation had eluded previous reports, because most of them focus on the humerus or humeral-radial joint. We observed in humeri of a 5-, 7- and 20-year-old animal only a modest cartilage layer in the diaphysis. Although in *Xenopus*, the extent of ossification correlates with decreased regenerative potential, axolotls can still regenerate their ossified limbs in paedomorph adults and induced land adults (Young et al., 1985). However, both the potential for regeneration and its fidelity are reduced (Monaghan et al., 2014).

Importantly, our results show that ossification in appendicular skeleton is continuous. Rux *et al* observed some degree of ossification of the metaphysis in the femur of a 10-year-old axolotl (Rux et al., 2018), which suggested that ossification of the appendicular bones continues throughout life. Here, we have shown that this ossification reaches the epiphyses with time.

Contribution of COL1A2^{+} cells during ossification

One crucial difference between axolotl and mammalian bone formation is the differential activation of the *Col1a2* enhancer. In mammals only osteoblasts activate the *Col1a2* enhancer (Bou-Gharios et al., 1996), while in axolotls, chondrocytes show *Col1a2* enhancer activity and *Col2a1* expression (Gerber et al., 2018; Khattak et al., 2013). We have observed *Col1a2* enhancer activity as the first mesenchymal condensates form in the limb bud (stage 47), and it remains active in cells inside the cartilage of long bones and in HCs at the chondro-osseous junction even in 7-year-old animals (personal observation).

We consider that understanding the development of the appendicular bones is essential to comprehend how limb regeneration is orchestrated in stages where different cell types and different maturation stages are found at the amputation plane. The common use of larval stage to study regeneration is normally preferred because it concludes faster and it is easier to image.

One cell type largely ignored in regeneration is the adipocyte. We saw the contribution of COL1A2[+] descendants to adipocytes in the marrow cavity. Although our *Col1a2-Tg* line also labels periosteum, their contribution to homeostasis is typically restricted to skeletal cells.

Therefore, we speculate that the contribution of COL1A2$^+$ descendants to adipocytes is most

likely from chondrocytes. This result agrees with work done in zebrafish, in which lineage

tracing of chondrocytes has shown the contribution of these cells to the marrow adipocytes

(Giovannone et al., 2019), similar to what we show in this study. Lopez et al. have shown that

the axolotl bone marrow is nonhematopoietic in young adults (1-year-old) (Lopez et al., 2014).

Our study detects an adipocyte-filled marrow cavity in long bones of the zeugopod in young

and aged animals (Fig 2A, 2B, 5A).

SOX9$^+$/OCN$^+$ cells

During endochondral ossification, the fate of HCs was long considered to be death and

subsequent substitution for bone cells. Unexpectedly, we found that HCs at the chondro-

osseous junction are SOX9$^+$/OCN$^+$. OCN is a non-collagenous protein in the bone matrix

expressed and secreted by committed osteoblast. Although murine HCs can become

osteoblasts and osteocytes during endochondral bone formation (Giovannone et al., 2019;

Jing et al., 2017; Yang et al., 2014), it is surprising to find cells expressing both SOX9 and OCN.

SOX9 is known to block osteoblast differentiation and to maintain columnar chondrocyte

proliferation by delaying pre-hypertrophy, but also directing hypertrophic maturation (Dy et

al., 2012; Hattori et al., 2010; Lui et al., 2019). The roles of SOX9 are various, and its expression

in committed osteoblast has not been shown before, which suggests the presence of chondro-

osteogenic hybrid cells during the normal transition to the ossified skeleton in axolotl.

Chondro-osteogenic hybrids are found generally in reparative instances such as jaw

regeneration in zebrafish (Paul et al., 2016) and murine rib (Kuwahara et al., 2019) and may

hold an unexplored potential for bone regeneration. In future studies, with transgenic lines

labeling specifically HCs, we will be able to conclusively show their contribution to different

cell types in the axolotl skeleton.

Perspectives

The origin of adipocytes during ossification and their main role inside the marrow

cavity needs to be further explored in axolotls. The known role of bone marrow adipocytes is

to promote hematopoietic stem cell renewal. However, marrow cavity of axolotls remains

devoid of a hematopoietic niche (Lopez et al., 2014). We assume that the adipose tissue in the

marrow is white fat based on morphology and the fact that brown adipose tissue has not been

reported in amphibians (Gesta et al., 2007). As such, it is possible that these cells have instead,
an important role in osteogenesis and energy metabolism in the axolotl. Another cell type of
interest is the periskeletal cell. McCusker *et al*. had shown that the periskeletal tissue of older
animals of TL 25 cm is multilayered and more complex than that of larvae of TL 6.5 cm
(McCusker et al., 2016). Importantly, periskeletal tissue is mainly responsible for the
regeneration of the skeleton (Currie et al., 2016; McCusker et al., 2016). The high potential of
these cells has neither been tested in non-regenerative conditions, nor their origin and
contribution to the formation of appendicular bone has been investigated.

Concluding remarks

Taken together, our data provide evidence that bone remodeling is a continuous
process in axolotls, starting from a mineralized cartilaginous skeleton during the larval stage
to a slowly transitioning bone that continues to ossify as animals age. The bone ossification
occurs in the absence of thyroxine or a biomechanical challenge suggesting instead the
existence of a physiological timer either triggered by sexual maturation or body mass.
However, this ossification can be accelerated in response to exogenous thyroxine, creating in
short time a bone mass that would normally take longer to form. Our findings add to a growing
body of literature indicating that vertebrate skeletal tissues are structurally diverse, and break
many of the established mammalian-centric rules of development and maturation.

Experimental procedures

Animal husbandry, transgenesis, metamorphosis and sample collection

Axolotls (*Ambystoma mexicanum*) husbandry was performed in the CRTD axolotl

facility, following the guidelines previously published (Khattak et al. 2014). Husbandry and

experimental procedures were performed according to the Animal Ethics Committee of the

State of Saxony, Germany. Animals used were selected by its size and it is indicated in each

experiment individually (total length = TL; snout to vent = SV). Sex determination in juvenile

larvae was performed by PCR based on (Keinath et al., 2018).

White axolotls (*d/d*) were used for most of the experiments. Transgenic lines used

included the previously published TgSceI*(Mmus.Col1a2:TFPnls-T2a-ERT2-Cre-ERT2)*[emt]

(referred as *Col1a2-Tg*) and TgSceI*(Mmus.Col1a2:TFPnls-T2a-ERT2-Cre-*

ERT2;Mmus.CAGGS:lp-GFP-3pA-lp-Cherry)[emt] (referred as *Col1a2xLPCherry-Tg*) (Gerber et al.

2016). We generated the transgenic lines C-Ti[t/+]*(Sox9:Sox9-T2a-mCherry)*[emt] (referred as *Sox9-*

Tg) and TgTol2*(Mmus.Tie2:EGFP)*[tsg] (referred as *Tie2-Tg*).

For generation of the *Tie2-Tg* transgenic line, we used the plasmid pSPTg.T2FXK (#54),

a gift from Thomas Sato (Addgene plasmid # 35963). The *Egfp* coding region was cloned 3´

from the promoter together with *Tol2* sequences. Fertilized embryos from *d/d* axolotls were

injected with the *Tie2:EGFP* vector and *Tol2* mRNA as previously described (Khattak et al.

2014). The *Sox9-Tg* reporter line is a targeted Knock-in line created by CRISPR/Cas9 technology

according to the published protocol (Fei et al., 2018). Briefly, a portion of the Sox9 gene

including a part of the second (last) intron and the remaining downstream part of the CDS was

PCRed from the axolotl genomic DNA and inserted into the vector pGEM-T along with a DNA

fragment encoding T2a-Cherry-3xnls. The resulting vector was injected into fertilized eggs

along with the CAS9 protein and the gRNA targeting the sequence

GGACTGCTGGCGAATGCACC, which is found in the intron sequence within the vector and in

the genome. As a result, mCherry fused to the C-terminus of the SOX9 and separated by a T2a

self-cleaving peptide is expressed from the endogenous SOX9 genomic locus.

Metamorphosis was induced in *d/d* animals as previously described (Khattak et al.,

2014). Briefly, axolotls TL 14 cm were anesthetized prior to intraperitoneal injections with 1.5

µL of L-thyroxine (Sigma, T2376) per gram of bodyweight (stock 1 µg/µL in DMSO). Animals

were cleaned regularly and water level was reduced as gills were lost. Only 1 out of 15 animals died during metamorphosis. Tissue was collected at 35 days post injection.

Recombination of *Col1a2xLPCherry-Tg* was induced by keeping animals in separated tanks with water containing (Z)-4-hydroxytamoxifen (4-OHT, Sigma T5648) 2 µM overnight. Animals were then washed and screened two weeks later to ensure conversion.

For tissue collection, animals were anesthetized with 0.007% benzocaine solution. After collection, animals were euthanized by exposing them to. a lethal dosage of anesthesia (0.1% benzocaine) for at least 20 min.

Alcian blue/alizarin red staining

Limbs collected were fixed with formaldehyde 10% at 4°C for 1 week and then washed three times with PBS 1x and dehydrated with serial EtOH washes (25, 50 and 70%). Limbs were either stored at -20°C in EtOH 70% or processed immediately at RT. Epidermis from animals > 12 cm TL was removed. Limbs were stained with alcian blue solution (Sigma A3157; alcian blue 0.0001% in EtOH 60% / glacial acetic acid 40%) for 3 days. Then, samples were rehydrated with EtOH dilution series in H_2O (80, 70, 50, 25%), before treatment with trypsin 1% in borax 30% for 30 min. Limbs were rinsed with KOH 1% and stained with alizarin red solution (Sigma A5533; alizarin red 0.0001% in KOH 1%) for 3 days. Next, limbs were washed with KOH 1% and cleared in KOH 1% / glycerol 20% overnight. Finally, limbs were dehydrated with EtOH dilution series (25, 50, 70, 90 and 3x 100%) and later transferred into serial washes of glycerol/EtOH (1:3, 1:1 and 3:1). Limbs were stored in 100% glycerol at 4°C.

µCT scan

Limbs collected were fixed with formaldehyde 10% at 4°C for 1 week and then washed three times with PBS 1x and dehydrated with serial EtOH washes (25, 50 and 70%) and stored in EtOH 70% at -20°C until measurement. Bone volume and microarchitecture were determined by micro-computed tomography (µCT, vivaCT40, ScancoMedical) of zeugopodial bones using pre-defined script 6. The isotropic voxel size was 10.5 µm (70 kVp, 114 µA, 200 ms integration time). Diaphysis scans were done for the whole zeugopod (both elements at the same time) and included 600 – 1000 slices. ScancoMedical protocols were used for all analyses and 3D reconstructions. Threshold used for analyses in Fig. 1 was 220 mg HA/cm^3, and for Fig. 6 was 150 mg HA/cm^3.

Paraffin sectioning and H&E staining

Skeletal elements were isolated and fixed with MEMFa 1x (MOPS 0.1M pH7.4 / EGTA 2mM / MgSO4x7H2O 1mM / 3.7% formaldehyde) overnight at 4°C, washed with PBS 1x and decalcified with EDTA 0.5M pH7.4 at 4°C for 1 to 4 weeks. After decalcification, samples were washed with PBS 1x and dehydrated with serial EtOH washes (25, 50, 70 and x3 100%). Samples were then incubated 3x with Roti®Histol (Carl Roth, 6640) at RT and 4x with paraffin at 65°C in glass containers. After last incubation, samples were embedded in paraffin (Roti®-Plast, Carl Roth, 6642) using plastic containers and stored at RT. Longitudinal sections of 6 µM thickness were obtained.

H&E staining on paraffin sections was performed following the producer recommendations (Sigma, Procedure No. HHS)

Cryosectioning and immunofluorescence

Skeletal elements were isolated and fixed with MEMFa 1x overnight at 4°C, washed with PBS 1x and decalcified with EDTA at 4°C for 48 hours. Next, the skeletal elements were washed with PBS 1x and dehydrated overnight with sucrose 30% at 4°C. Samples were embedded in O.C.T. compound (Tissue-Tek, 4583) using plastic molds and frozen with dry ice for 1 hour prior to storage at -20°C. Longitudinal sections of 12 µm thickness were cut and mounted on superfrost slides. Slides were kept at -20°C until processed.

For immunofluorescence, slides were dried at RT for at least 1 hour. Sections were washed 3x with PBS 1x + 0.3% Tx-100 prior to blocking with PBS 1x + 0.3% Tx-100 + 10% normal horse serum for 1 hour. Primary antibody incubation was done in blocking solution overnight at 4°C. Sections were then washed 3x with PBS 1x + 0.3% Tx-100 and incubated with secondary antibody 1:300 and nuclear staining 1:10.000 for 2 hours. Finally, sections were washed 3x with PBS 1x + 0.3% Tx-100 and mounted using Mowiol mounting medium (Carl Roth, 0713). The antibodies and nuclear stainings used in this work are listed in table I.

Whole mount immunofluorescence and tissue clearing

Limbs or isolated skeletal elements were fixed with MEMFa 1x overnight at 4°C, washed with PBS 1x and decalcified with EDTA at 4°C for 48 hours. Whole mount immunofluorescence protocol was adapted from (Masselink et al., 2019). Briefly, skeletal

elements were washed overnight with PBS 1x + 0.3% Tx-100 at RT and then blocked in PBS + 0.3% Tx-100 + 10% DMSO + 5% goat serum for 24 hours at 37°C. Samples were incubated with primary antibody anti-GFP in blocking solution for 3 days at 37°C. Next, samples were briefly rinsed with PBS + 0.3% Tx-100 and then washed 4x for 2 hours at 37°C. Samples were blocked again at 37°C overnight and then incubated with secondary antibody for 2 days at 37°C. After incubation, samples were rinsed with PBS + 0.3% Tx-100 and then washed 2x for 2 hours at 37°C. Nuclear staining was done for 1 hour in washing solution at RT and samples were then washed 4x in PBS 1x for 15 min at RT. Isolated skeletal elements were embedded in agarose 1% before clearing.

For clearing, samples were dehydrated with serial washes of EtOH (25, 50, 70, 100%) for 2 hours each at 4°C. Samples were then incubated overnight in EtOH 100% at 4°C prior to clearing with ethyl cinnamate (Sigma, 112372-100G) at RT for at least 2 hours. Samples were imaged the same day. The antibodies and nuclear stainings used in this work are listed in table I.

Table I: Antibodies and nuclear staining.

Antibody / Nuclear Staining	Dilution	Cat. Number
Anti-GFP (chicken)	1:500	Abcam (ab13970)
Anti-OCN (rabbit)	1:250	Abcam (ab198228)
Goat anti-Rabbit, Alexa Fluor 647	1:300	Invitrogen (A-21245)
Goat anti-Chicken, Alexa Fluor 488	1:1000	Invitrogen (A-11039)
TO-PRO™-3 iodide (642/661)	1:10.000	Invitrogen (T3605)
SYTOX™ Green nucleic acid stain	1:10.000	Invitrogen (S7020)

Microscopy

Alcian blue/alizarin red staining imaging was performed on a Zeiss Discovery.V20 stereomicroscope (Plan S 1.0x). H&E staining imaging was performed on a Zeiss Axio Observer.Z1 inverted microscope (Plan-apochromat 20x/0.8). Immunofluorescence imaging was performed on a Zeiss Axio Observer.Z1 inverted microscope with an ApoTome1 system (Plan-apochromat 10x/0.45) and on a Zeiss confocal laser scanning microscope LSM 780 (Plan-apochromat 10x/0.45 or Plan-apochromat 20x/0.8). Whole mount immunofluorescence

imaging was performed on a Zeiss confocal laser scanning microscope LSM780 (Plan
apochromat 10x/0.45).

Statistical analysis

Statistical analysis was performed using the software Prism9 (GraphPad Software, LLC)
for macOS. To assess the differences in TL or SV between sexes (Fig 1C), a paired t test was
performed between samples. To assess the differences in volume between radii and ulnas of
different TL animals (Fig 1G) and between radii and ulnas of paedomorph and metamorphic
animals (Fig 6C), a Two-way ANOVA test was performed, using a post hoc Tukey's multiple
comparisons test for assessing statistical significance between each sample pair. P-values <
0.05 were considered statistically significant.

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Authors contribution

CRG and TSG conceived experiments, analysed data and wrote the manuscript. CRG
conducted experiments. MR, MS and AB contributed experimental data and support. DK
created *Sox9-Tg* line. DK and MR contributed with manuscript revision and discussions. TSG
secured funding for the project.

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Figure legends

Figure 1: Appendicular skeleton is ossified with growth

(A) Quantification of axolotl total length (TL) in cm as a function of its age in months. Animal sex was not considered (n = 213). Y-axis scale was set to logarithmic scale to better represent the growth phases.

(B) Quantification of axolotl snout to vent length (SV) in cm as a function of its age in months. Animal sex was not considered (n = 213). Y-axis scale was set to logarithmic scale to better represent the growth phases.

(C) Quantification of TL and SV in cm in 1-year-old siblings as a function of its sex. (n = 18. 9 males, 9 females. *** $p < 0.001$, student's t test).

(D) Alcian blue/alizarin red staining of limbs from different TL animals. R: radius, U: ulna. Scale bar for 4 and 10 cm: 500 µm. Scale bar for 12 to 20 cm: 1 mm.

(E) *In vivo* alizarin red staining of axolotl TL 5 cm. Scale bar: 500 µm

(F) 3D reconstructions from µCT scan for zeugopodial bones from different TL animals. All bones are scaled. Scale bar: 250 µm.

(G) Quantification of zeugopodial bones volume (cm^3) from different TL animals. (n = 4 per size. a: radius TL 20 cm compared to radius from TL 10, 12, 14 and 16 cm. b: ulna TL 20 cm compared to ulna from TL 10, 12, 14 and 16 cm. c: radius TL 20 cm compared to ulna TL 20 cm. a, b: $p < 0.001$, c: $p < 0.05$, Tukey's multiple comparisons test).

(H) 3D reconstruction from µCT scan for zeugopodial bones from axolotl TL 25 cm. Bones scaled to (E). Scale bar: 500 µm.

Figure 2: Bone is vascularized and marrow cavity is filled with adipocytes

(A) Representative sections of radii stained with H&E from axolotls TL 6 and 12 cm. RZ: resting

zone, PZ: proliferative zone, HZ: hypertrophic zone, MC: marrow cavity, D: distal, P: proximal.

Scale bar: 500 µm.

(A') Inset from TL 6 cm in (A). Arrowhead: calcified ring around HZ. Scale bar: 250 µm

(A'') Inset from TL 12 cm in (A). Arrowhead: discontinuous ossification around diaphysis. Scale

bar: 250 µm

(B) H&E staining in the distal half of the radius from an axolotl TL 16 cm. Scale bar: 500 µm.

(C) Confocal image of anti-GFP whole-mount IF from forelimb or radius from *Tie2-Tg* (red) TL

6 or 12 cm respectively. TO-PRO™3 was used for nuclear staining (white). Image represents a

maximum intensity projection of 3 images (10 µm interval). Arrowheads: bone vascularization.

Scale bar: 250 µm.

(D) Inset from *Tie2-Tg* TL 6 cm in (C) with overlapping brightfield image. Arrowhead: blood

vessel in contact with calcified cartilage. Scale bar: 100 µm.

Figure 3: Sox9 labels chondrogenic cells while Col1a2 labels all skeletal cells

(A) Confocal image of ulnas sections from *Sox9-Tg* (red) axolotls TL 6 and 12 cm. Calcified tissue stained with calcein (green). TO-PRO™3 was used for nuclear staining (white). D: distal, P: proximal. Scale bar: 250 µm.

(B) Confocal image of ulna sections from *Col1a2-Tg* (red) axolotls TL 6 and 12 cm. TO-PRO™3 was used for nuclear staining (white). Scale bar: 250 µm.

(C) Confocal image of anti-Sox9 IF in ulna sections from *Sox9-Tg* in axolotl TL 16 cm. SYTOX™ Green was used for nuclear staining. Scale bar: 250 µm

(D) Inset from (A), *Sox9-Tg* TL 12 cm. Dotted line shows boundary between cartilage and bone (chondro-osseous junction), hc: hypertrophic chondrocytes. Scale bar: 250 µm.

(E) Inset from (B) *Col1a2-Tg* TL 12 cm, os: osteocyte, eo: endosteal cell, hc: hypertrophic chondrocytes. Scale bar: 250 µm.

Figure 4: Col1a2$^+$ cells contribute to OCN$^+$ osteoblasts

(A) Confocal image of anti-OCN IF (red) from representative radii sections from axolotls TL 6 and 12 cm. SYTOX™ Green was used for nuclear staining (white). D: distal, P: proximal. Scale bar: 250 µm.

(B) Inset from (A) anti-OCN IF (red). Dotted line shows boundary between cartilage and bone. Solid arrowheads: hypertrophic chondrocytes. Arrowheads: periosteal cells.

(C) Confocal image of anti-OCN IF from radii sections from *Sox9-Tg* TL 12 cm animals. SYTOX™ Green was used for nuclear staining . Scale bar: 250 µm.

(D) Confocal image of *Col1a2xLPCherry-Tg* (red) ulna sections from tamoxifen-converted axolotls at TL 5 cm and grown to TL 14 cm. SYTOX™ Green was used for nuclear staining (white). Dotted line shows boundary between cartilage and bone, c: cartilage, b: bone, mc: marrow cavity. Scale bar: 250 µm.

(E) Confocal image of *Col1a2xLPCherry-Tg* (red) ulna sections from axolotls TL 14 cm. SYTOX™ Green was used for nuclear staining (white), po: periosteum, eo: endosteum, ad: adipocytes. Scale bar: 100 µm.

(F, F', F'') Confocal image of anti-OCN IF (green) (F) from ulna sections from *Col1a2xLPCherry-Tg* (red) (F') axolotls TL 14 cm. SYTOX™ Green was used for nuclear staining (white). (F'') Merge. Scale bar: 100 µm.

 Figure 5: Bone ossification and marrow cavity expand with age

(A) H&E staining of radii sections from different age axolotls. D: distal, P: proximal, ep:

epiphysis, mc: marrow cavity, b: bone. Scale bar: 500 μm.

(B) Alcian blue/alizarin red staining of forelimb from a 5-year-old axolotl. Scale bar: 2 mm.

(C) Alcian blue/alizarin red staining of foot from a 20-year-old axolotl. Scale bar: 2 mm.

(D) H&E staining of proximal phalanx from 2-year-old axolotl. Scale bar: 1 mm.

(E) H&E staining of humerus from 5-year-old axolotl. Scale bar: 1 mm.

Figure 6: Axolotl metamorphosis accelerates bone ossification

(A) H&E staining from radii sections from paedomorphic and metamorphic axolotls (35 dpi).

D: distal, P: proximal. Scale bar: 500 μm.

(B) 3D reconstructions from μCT scans for zeugopodial bones from aquatic and metamorphic

axolotl. All bones are scaled. Scale bar: 500 μm.

(C) Quantification of zeugopodial bones volume (cm^3) from aquatic and metamorphic axolotl.

(n = 5 for aquatic zeugopods and n = 6 for metamorphic zeugopods. ** $p < 0.01$, * $p < 0.05$,

student's t test).

A — Total length (TL) in Axolotl
B — Snout to vent length (SV) in Axolotl
C — Sex and size correlation
○ Female
● Male

D — 4 cm   10 cm   12 cm   14 cm   16 cm   20 cm
R   U

E — 5 cm
Brightfield / Alizarin red

F — 12 cm   14 cm   16 cm   20 cm
R   U

H — 25 cm
R   U

G — Zeugopodial bones volume
● Radius
○ Ulna
a,c   b

A 6 cm 12 cm

RZ

PZ

HZ

MC

D
P

A'

A'

A"

B 16 cm

HZ

Periosteum

Osteocytes

Adipocytes

Endosteum

C

6 cm

12 cm

D
P

D

Nuclei / anti-GFP (*Tie2-Tg*)

D

Brightfield /
anti-GFP (*Tie2-Tg*)

A   6 cm   D⤑P   12 cm   C

Nuclei / mCherry (*Sox9-Tg*) / Calcein

B   6 cm   12 cm   D

Nuclei / TFP (*Col1a2-Tg*)

C   16 cm

Nuclei   anti-Sox9   mCherry (*Sox9-Tg*)   mCherry (*Sox9-Tg*) / anti-Sox9

D   Cartilage   hc   Bone

Brightfield / mCherry (*Sox9-Tg*)

E   Cartilage   hc   os   Bone   eo

Brightfield / TFP (Col1a2-*Tg*)

A

6 cm

12 cm

P ⤑ D

B

Nuclei / anti-OCN

B

Cartilage

Bone

Nuclei / anti-OCN

C

Cartilage

Bone

mCherry (*Sox9-Tg*)　　　anti-OCN　　　Nuclei

D

14 cm

c

D ⤑ P

b　mc

b

Nuclei / mCherry
(*Col1a2xLPCherry-Tg*)

E

po▷

eo▷　◁ad

◁ad

Nuclei / mCherry
(*Col1a2xLPCherry-Tg*)

F''

Nuclei / anti-OCN /
mCherry(*Col1a2xLPCherry-Tg*)

F

Nuclei / anti-OCN

F'

eo▷　　　　eo▷

◁eo

◁eo　　▷po

Nuclei / mCherry
(*Col1a2xLPCherry-Tg*)

A

11-months   5-years   7-years   20-years

B   5-years

C   20-years

D   Proximal Phalanx (2-years)

E   Humerus (5-years)

A Paedomorph   Metamorphic

D
↑
P

B Paedomorph   Metamorphic

R   U   R   U

C

Zeugopodial bones volume

Volume (cm^3)

● Radius
○ Ulna

Paedomorph Metamorphic Paedomorph Metamorphic

** *